

# Prostaglandin D<sub>2</sub>-Mediated Microglia/Astrocyte Interaction Enhances Astroglia and Demyelination in *twitcher*

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Prostaglandin (PG) D<sub>2</sub> is well known as a mediator of inflammation. Hematopoietic PGD synthase (HPGDS) is responsible for the production of PGD<sub>2</sub> involved in inflammatory responses. Microglial activation and astroglia are commonly observed during neuroinflammation, including that which occurs during demyelination. Using the genetic demyelination mouse *twitcher*, a model of human Krabbe's disease, we discovered that activated microglia expressed HPGDS and activated astrocytes expressed the DP<sub>1</sub> receptor for PGD<sub>2</sub> in the brain of these mice. Cultured microglia actively produced PGD<sub>2</sub> by the action of HPGDS. Cultured astrocytes expressed two types of PGD<sub>2</sub> receptor, DP<sub>1</sub> and DP<sub>2</sub>, and showed enhanced GFAP production after stimulation of either receptor with its respective agonist. These results suggest that PGD<sub>2</sub> plays an important role in microglia/astrocyte interaction. We demonstrated that the blockade of the HPGDS/PGD<sub>2</sub>/DP signaling pathway using HPGDS- or DP<sub>1</sub>-null *twitcher* mice, and *twitcher* mice treated with an HPGDS inhibitor, HQL-79 (4-benzhydryloxy-1-[3-(1H-tetrazol-5-yl)-propyl]piperidine), resulted in remarkable suppression of astroglia and demyelination, as well as a reduction in twitching and spasticity. Furthermore, we found that the degree of oligodendroglial apoptosis was also reduced in HPGDS-null and HQL-79-treated *twitcher* mice. These results suggest that PGD<sub>2</sub> is the key neuroinflammatory molecule that heightens the pathological response to demyelination in *twitcher* mice.

## Introduction

Prostaglandin (PG) D<sub>2</sub> is a well known mediator of inflammation. It augments vascular permeability (Flower et al., 1976), chemotaxis (Hirai et al., 2001), and antigen presentation (Herve et al., 2003), inhibits platelet aggregation (Whittle et al., 1978), and induces vasodilatation and bronchoconstriction (Wasserman et al., 1980). Hematopoietic PGD synthase (HPGDS) is responsible for the production of PGD<sub>2</sub>, including that linked to immune and

inflammatory responses (Kanaoka and Urade, 2003). HPGDS is expressed in human Th2 lymphocytes (Tanaka et al., 2000), antigen-presenting cells (Urade et al., 1989), mast cells (Urade et al., 1990), and microglia (Mohri et al., 2003).

Microglial activation and astroglia are commonly observed during the neuroinflammation associated with brain injury, infection, and neurodegenerative diseases. However, the molecular mechanism underlying microglial activation and astroglia is still poorly understood. The *twitcher* mouse (C57BL/6J-*GALC*<sup>twi</sup>:*GALC*<sup>twi/twi</sup>) is an authentic animal model of human globoid cell leukodystrophy (Krabbe's disease) resulting from the deficiency of galactosylceramidase (Duchen et al., 1980; Kobayashi et al., 1980). In this model, initial myelination progresses normally (Nagara et al., 1982), but then demyelination begins as a result of the apoptosis of oligodendrocytes (OLs) after postnatal day 30 (P30) (Taniike et al., 1999). Demyelination in *twitcher* progresses in an orderly manner (Taniike and Suzuki, 1994) and is always accompanied by both microglial activation and astroglia throughout the brain (Ohno et al., 1993b). Interestingly, the pathological features of *GALC*<sup>twi/twi</sup> have many characteristics in common with multiple sclerosis (MS). For example, in both conditions, major histocompatibility complex (MHC) molecules contribute to disease progression (Ohno et al., 1993a; Matsushima et al., 1994; Taniike et al., 1997). The absence of

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MHC class II molecules reduces CNS demyelination and microglial/macrophage infiltration in the *twitcher* brain (Matsushima et al., 1994). It has also been reported that there is expression of proinflammatory molecules, such as interleukin-6 (IL-6), tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) (LeVine and Brown, 1997), monocyte chemoattractant protein-1, interferon-induced protein-10, macrophage inflammatory protein-1 $\alpha$  (MIP-1 $\alpha$ ), MIP-1 $\beta$ , and RANTES (regulated on activation normal T-cell expressed and secreted) (Wu et al., 2000) in *twitcher*. Furthermore, we recently reported that TNF $\alpha$  expression increased with the progression of demyelination in *twitcher* and that a TNF $\alpha$  inhibitor suppresses such demyelination (Kagitani-Shimono et al., 2005). These correlations make the *twitcher* mice a model system for investigating the molecular events of neuroinflammation in demyelinating diseases.

In the present study, we found that HPGDS expression was progressively upregulated in activated microglia in the *GALC*<sup>twi/twi</sup> brain. This upregulation was accompanied by the expression of the PGD<sub>2</sub> receptor DP<sub>1</sub> in hypertrophic astrocytes located close to HPGDS-positive (HPGDS<sup>+</sup>) microglia. Using glial cells in primary culture, we demonstrated that activated microglia produced a large amount of PGD<sub>2</sub> synthesized by HPGDS and that astrocytes expressed both DP<sub>1</sub> and DP<sub>2</sub> receptors and were activated by PGD<sub>2</sub>. Using HPGDS- or DP<sub>1</sub>-null *GALC*<sup>twi/twi</sup> mice and *GALC*<sup>twi/twi</sup> mice treated with an HPGDS-inhibitor, we also found that blockade of the HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> signaling pathway resulted in the suppression of astrogliosis and apoptosis of OLs as well as in that of demyelination. This is the first example of a PGD<sub>2</sub>-mediated microglia/astrocyte interaction that enhances neuroinflammation and demyelination, and our findings may cast new light on strategies for the treatment of Krabbe's disease.

## Materials and Methods

**Mice.** All animal experiments were performed in accordance with the Japanese Law for the Protection of Experimental Animals and conformed to the regulations issued by the National Institutes of Health and the Society for Neuroscience. *GALC*<sup>twi/+</sup> mice were originally purchased from The Jackson Laboratory (Bar Harbor, ME), and the mutation was maintained by interbreeding of known heterozygous mice. DNA was extracted from clipped tails for the detection of wild-type or mutant alleles of *GALC*, which was done with primers gtwf2 (5'-CAGTCATTCAGAAAGTCTTCC-3') and gtwr3 (5'-GCCACTGTCTCAGGTGATA-3'; mismatch primer). The PCR fragment of 237 bp of the normal allele was not digested by *EcoRV*, whereas that of the mutant allele was, resulting in a 217 bp fragment.

*HPGDS*<sup>-/-</sup> and *DP*<sub>1</sub><sup>-/-</sup> mice were generated by standard gene-targeting technology with mouse embryonic stem cells derived from the 129 strain and backcrossed to the inbred C57BL/6J strain 15 times. The former strain was created by the Osaka Bioscience Institute and Japan Tobacco Inc. Pharmaceutical Frontier Research Laboratories, and the latter was created at Kyoto University (Matsuoka et al., 2000). Neither strain exhibits any adverse neurological symptomatology. For genotyping of *HPGDS*, primer FW1 (5'-GAGTTGCTGCATCTGACCTTTC-3') and RV1 (5'-TAGCGAATAATTCGGCTCTTCC-3') were used to detect the wild-type allele, and H-FW1 (AAGATCTGTCTTGTGTCGTACGCT-3') and *NeoR8* (5'-GTCCAGATCATCTGATCGAC-5') were used for that of the targeted allele. For the detection of wild-type or mutant alleles of *DP*<sub>1</sub>, primer DP<sub>1</sub> (5'-TCGGTCTTTTATGTGCTCGTG-3') or *Neo1* (5'-CCCGTGATATTGCTGAAGAGC-3') was used together with DP<sub>2</sub> (5'-GAATCATCTGGATGAAACACC-3').

Double-mutant mice (*HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> or *DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>twi/twi</sup>) were generated by mating *HPGDS*<sup>-/-</sup> C57BL/6J (*HPGDS*<sup>-/-</sup>*GALC*<sup>+/+</sup>) or *DP*<sub>1</sub><sup>-/-</sup> C57BL/6J (*DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>+/+</sup>) offspring with *HPGDS*<sup>+/+</sup>*DP*<sub>1</sub><sup>+/+</sup>*GALC*<sup>twi/+</sup>. The F1 litters (*HPGDS*<sup>+/+</sup>*GALC*<sup>twi/+</sup> and *DP*<sub>1</sub><sup>+/+</sup>*GALC*<sup>twi/+</sup>) were mated with each other to produce *HPGDS*<sup>-/-</sup>

*GALC*<sup>twi/twi</sup>, *DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>twi/twi</sup>, and *HPGDS*<sup>+/+</sup>*DP*<sub>1</sub><sup>+/+</sup>*GALC*<sup>twi/twi</sup>. These three types of *GALC*<sup>twi/twi</sup> male mice were used for analysis. The frequency of each genotype in all generations conformed to Mendelian inheritance patterns.

HQL-79 (4-benzhydroxy-1-[3-(1*H*-tetrazol-5-yl)-propyl]piperidine) (Aritake et al., 2006), an inhibitor of HPGDS, was purchased from Cayman Chemical (Ann Arbor, MI). *GALC*<sup>twi/twi</sup> received a daily subcutaneous injection of HQL-79 (suspended in 0.5% methylcellulose; 50 mg · kg<sup>-1</sup> · d<sup>-1</sup>) from P25 to P45. As a control, the same volume of vehicle was injected into *GALC*<sup>twi/twi</sup> during the same period.

**Materials.** Rabbit polyclonal (0.1  $\mu$ g/ml) and rat monoclonal (0.2  $\mu$ g/ml) anti-mouse HPGDS antibodies were raised and purified at Osaka Bioscience Institute (Mohri et al., 2003). The preabsorbed antibody was prepared by incubation of the polyclonal antibody with an excess amount of purified mouse HPGDS recombinantly expressed in *Escherichia coli* (Kanaoka et al., 2000). Guinea pig anti-DP<sub>1</sub> antiserum (1:30) was also raised at Osaka Bioscience Institute (Mizoguchi et al., 2001). The other antibodies used were as follows: rabbit polyclonal antibodies against  $\pi$ -class glutathione *S*-transferase ( $\pi$ -GST) (1:1000; Biotrin International, Dublin, Ireland), NG2 (1:200; Chemicon, Temecula, CA), myelin basic protein (MBP) (1:5000; DakoCytomation, Glostrup, Denmark), cow glial fibrillary acidic protein (GFAP) (1:5000; DakoCytomation), and rat monoclonal anti-mouse F4/80 antibody (1:500; Serotec, Oxford, UK). Biotinylated *Ricinus communis*-agglutinin-1 (RCA-1) was purchased from Vector Laboratories (Burlingame, CA) (50  $\mu$ g/ml).

**Measurement of HPGDS activity and PGD<sub>2</sub> content in the brain.** HPGDS activity in the brains of *GALC*<sup>+/+</sup> and *GALC*<sup>twi/twi</sup> mice at P39 ( $n = 4$  each) was determined by using 40  $\mu$ M [1-<sup>14</sup>C]PGH<sub>2</sub> in the presence of 1 mM glutathione, as described previously (Urade et al., 1985). Protein concentrations were determined with bicinchoninic acid reagent (Pierce, Rockford, IL), using bovine serum albumin as a standard as per the protocol of the manufacturer. Quantities of PGD<sub>2</sub>, PGE<sub>2</sub>, and PGF<sub>2 $\alpha$</sub>  in fresh-frozen brains of *GALC*<sup>+/+</sup> and *GALC*<sup>twi/twi</sup> mice decapitated at P39 ( $n = 6$  each) were determined by enzyme immunoassays as described previously (Ram et al., 1997).

**Northern blotting and quantitative PCR.** Total RNA was extracted from mouse cerebrum and cerebellum by the guanidinium thiocyanate-phenol-chloroform method using ISOGEN (Nippon Gene, Tokyo, Japan). Total RNA (10  $\mu$ g) was electrophoresed in an agarose gel, transferred to Zeta Probe nylon membranes (Bio-Rad, Hercules, CA), and hybridized with <sup>32</sup>P-labeled cDNA probes specific for mouse GFAP and glyceraldehyde-3-phosphate dehydrogenase (G3PDH). The blots were visualized by autoradiography with Eastman Kodak (Rochester, NY) XAR-5 film and an intensifying screen. The relative amount of each transcript was estimated by quantifying the associated radioactivity with a BAS-2000 (Fuji Film, Tokyo, Japan).

Quantitative PCR analysis of the amounts of mRNAs for HPGDS, DP<sub>1</sub>, DP<sub>2</sub>, peroxisome proliferator activated receptor  $\gamma$  (PPAR $\gamma$ ), GFAP, and G3PDH was performed by using a LightCycler amplification and detection system (Roche Diagnostics, Indianapolis, IN) as described below. Each sample of total RNA (2  $\mu$ g) was reverse transcribed at 50°C for 30 min into single-stranded cDNA in a reaction mixture containing 20 U of RNase inhibitor, 1  $\times$  RNA PCR buffer, 1 mM dNTP mixture, 2.5  $\mu$ M random primer, and 5 U of avian myeloblastosis virus reverse transcriptase (Takara Biomedicals, Kyoto, Japan). The sequence-specific primers used were as follows: HPGDS forward primer, 5'-GAATAGAACAAGCT-GACTGGC-3'; HPGDS reverse primer, 5'-AGCCAAATCTGTGTTTT-TGG-3'; DP<sub>1</sub> forward primer, 5'-TTTGGGAAGTTCGTGCACTACT-3'; DP<sub>1</sub> reverse primer, 5'-GCCATGAGGCTGGAGTAGA-3'; DP<sub>2</sub> forward primer, 5'-TGGCCTTCTTCAACAGCGT-3'; DP<sub>2</sub> reverse primer, 5'-ACGCAGTTGGGGAATTCG-3'; PPAR $\gamma$  forward primer, 5'-GGAGATCTCCAGTGATATCGACCA-3'; PPAR $\gamma$  reverse primer, 5'-ACGGCTTCTACGGATCGAACT-3'; GFAP forward primer, 5'-TTCCTGGAACAGCAAAACAAGCGCT-3'; GFAP reverse primer, 5'-CTGTCTATACGCAGCCAGGTTGTTCTC-3'; inducible nitric oxide synthase (iNOS) forward primer, 5'-CAGCAATATAGGCTCATCCA-GAG-3'; iNOS reverse primer, 5'-GTGGTGTAGGACAATCCA-CAACT-5'; G3PDH forward primer, 5'-TGAACGGGAAGCTCAC-TGG-3'; and G3PDH reverse primer, 5'-TCCACCACCCTGTTGCT-3'.

The constructs used to create a standard curve were made by cloning each amplified fragment into the *Hind*III site of a pGEM vector (Promega, Madison, WI). The number of copies was calculated by plotting a dilution series on this standard curve in each PCR experiment. The PCR mixture contained *Taq* polymerase, X LightCycle DNA master SYBR Green (Roche Diagnostics) reaction buffer, 3 mM MgCl<sub>2</sub>, and 12.5 pmol of each primer. After PCR had been performed for 40 cycles of denaturation (95°C for 1 s), annealing (57°C for 5 s), and enzymatic chain extension (72°C for 10 s), the products for HPGDS, DP<sub>1</sub>, DP<sub>2</sub>, PPARγ, GFAP, and G3PDH were detected at 87, 83, 83, 91, 87, and 84°C, respectively. All PCR products were visualized under UV light after electrophoresis in an agarose gel containing ethidium bromide and were subsequently sequenced to verify that only the specific polymerization from the intended mRNA had occurred.

**Western blotting analysis.** The brains were homogenized in 3 vol by weight of PBS. After centrifugation at 16,000 × *g* for 20 min, the supernatant and the precipitate were used in Western blotting for HPGDS and MBP, respectively. Protein was electrophoresed in 15/25% SDS-polyacrylamide gradient gel (Daiichi, Tokyo, Japan), and, subsequently, the separated proteins were electrophoretically transferred to an Immobilon polyvinylidene difluoride membrane (Millipore, Bedford, MA). The membrane was reacted with the rabbit polyclonal anti-HPGDS antibody (0.5 μg/ml) or anti-β-actin polyclonal antibody (1:1000; Abcam, Cambridge, UK) at 4°C overnight. After alkaline phosphatase-conjugated anti-rabbit IgG (1:3000) had been applied, HPGDS and MBP were detected by using a 5-bromo-4-chloro-indolyl-phosphate/nitroblue-tetrazolium-chloride developing kit (Amersham Bioscience, Uppsala, Sweden).

**Immunocytochemistry.** *GALC*<sup>twi/twi</sup> and *GALC*<sup>+/+</sup> mice at P20, P30, P40, and P45 were used for immunocytochemical analysis. Under deep ether anesthesia, the mice were perfused via the heart with physiological saline, followed by 4% paraformaldehyde in 0.1 M sodium phosphate, pH 7.4, for 10 min. The brains were removed and postfixed in the same fixative overnight. After a coronal slice had been taken, they were routinely embedded in paraffin. Additionally, two each of the *GALC*<sup>twi/twi</sup> and *GALC*<sup>+/+</sup> mice at P45 were perfused with physiological saline only and processed for the preparation of fresh-frozen sections. Both paraffin and frozen sections (5 μm thick) were mounted on 3-aminopropyl-triethoxysilane-coated slides.

HPGDS immunostaining was performed as described previously (Mohri et al., 2003). Rabbit anti-mouse HPGDS antibody was applied followed by biotinylated anti-rabbit IgG and avidin-biotin complex (ABC) from an ABC Elite kit (Vector Laboratories). Immunoreactivity was visualized by the addition of diaminobenzidine hydrochloride (DAB) (Dotite, Kumamoto, Japan).

The procedures for double labeling were followed as described previously (Mohri et al., 2003). Briefly, in the case of double immunocytochemistry for HPGDS and F4/80, after the primary antibodies had been applied, the sections were sequentially incubated with Texas Red-conjugated anti-rat IgG antibody (20 μg/ml; ICN, Aurora, OH) and biotinylated anti-rabbit IgG antibody (20 μg/ml; Vector Laboratories), followed by FITC-conjugated avidin D (20 μg/ml; Vector Laboratories). In the case of double labeling for HPGDS and biotinylated RCA-1, after incubation with the primary antibody or lectin, the sections were sequentially incubated with the Texas Red-conjugated anti-rabbit IgG antibody (ICN), followed by FITC-conjugated avidin D. The fluorescence was examined with an Axiovert 100M microscope connected to a Zeiss (Oberkochen, Germany) laser-scanning microscope. The frozen sections, optimal for immunostaining of DP<sub>1</sub>, were fixed with 100% ethanol at -20°C for 30 min and then with acetone at room temperature for 5 min. After the sections had been incubated for 5 min with 0.1 M sodium phosphate, pH 7.4, containing 0.005% saponin (Nakarai Tesque, Kyoto, Japan) and 20 μM (4-amidino-phenyl)-methane-sulfonyl fluoride (Wako, Osaka, Japan), they were incubated with guinea pig anti-mouse DP<sub>1</sub> antibody. They were then serially reacted with biotinylated anti-guinea pig antibody (dilution 1:400; Southern Biotechnology, Birmingham, AL) and ABC. The reaction using DAB as the chromogen was enhanced by adding 5 mg/ml nickel (II) ammonium sulfate hexahydrate (Nakarai Tesque) to the reaction solution. The double immunostaining

for DP<sub>1</sub> and F4/80 was performed on unfixed frozen sections as explained in the paragraph describing the double immunostaining of paraffin sections.

**Terminal deoxynucleotidyl transferase-mediated biotinylated UTP nick end-labeling (TUNEL) histochemistry** combined with immunocytochemistry for π-GST was performed as described previously (Taniike et al., 1999). π-GST-positive TUNEL-positive cells were counted in the cerebral parenchyma of paraffin sections prepared from mice at P45 (*n* = 4 each).

**Primary cultures of mouse microglia and astrocytes.** We prepared primary cultures of microglia and astrocytes from wild-type mouse brains at P1 and P7, respectively. Cerebral cortices were dissected, and the leptomeninges were completely removed. The tissues were minced, suspended in PBS containing 0.05% trypsin (Invitrogen, Grand Island, NY) and 0.01% DNase I (Sigma, St. Louis, MO), and then incubated for 10 min at 37°C. After incubation and centrifugation, the cell pellets were washed three times with PBS. The cells were then filtered through a 75 μm nylon mesh, centrifuged, suspended in DMEM (Nakalai Tesque) containing 10% fetal bovine serum (FBS) (JRH Bioscience, Lenexa, KS), 100 IU/ml penicillin, and 100 μg/ml streptomycin (Invitrogen), and transferred to culture dishes. For microglial cultures, the medium was changed to DMEM containing 10% FBS after a 24 h culture period. This medium was exchanged for fresh medium twice weekly thereafter. The supernatant including microglial cells was collected and subcultured at 1 × 10<sup>5</sup> cells per well (six-well plate). After incubation in DMEM without FBS for 6 h, the microglia were treated with A23187 (5 μM; Sigma) for 30 min. When the microglial cells were treated with HQL-79, the inhibitor was added 15 min before A23187 treatment. The medium after treatment of the cells with these compounds was measured for its content of PGs.

For astrocyte cultures, the medium was changed to DMEM containing 20% FBS after the initial 24 h culture period. Then, 2 d later, the medium was changed back to that containing 10% FBS. Once the astrocytes in the primary culture reached confluence, the cells were rinsed with PBS, suspended in trypsin-containing PBS, and subcultured at 5 × 10<sup>5</sup> cells per well (six-well plate). After a 24 h incubation in DMEM containing 1% FBS, the astrocytes were incubated with BW245C (0.1–100 nM; Cayman Chemical) or 13, 14-dihydro-15-keto-PGD<sub>2</sub> (DK-PGD<sub>2</sub>) (0.1–100 nM; Cayman Chemical) for 6 h at 37°C. After the cells had been washed with PBS, RNA was extracted from them and subjected to quantitative reverse transcription (RT)-PCR. We confirmed these cells to be either microglia or astrocytes by their positive immunoreaction with anti-RCA-1 and F4/80 or GFAP, respectively.

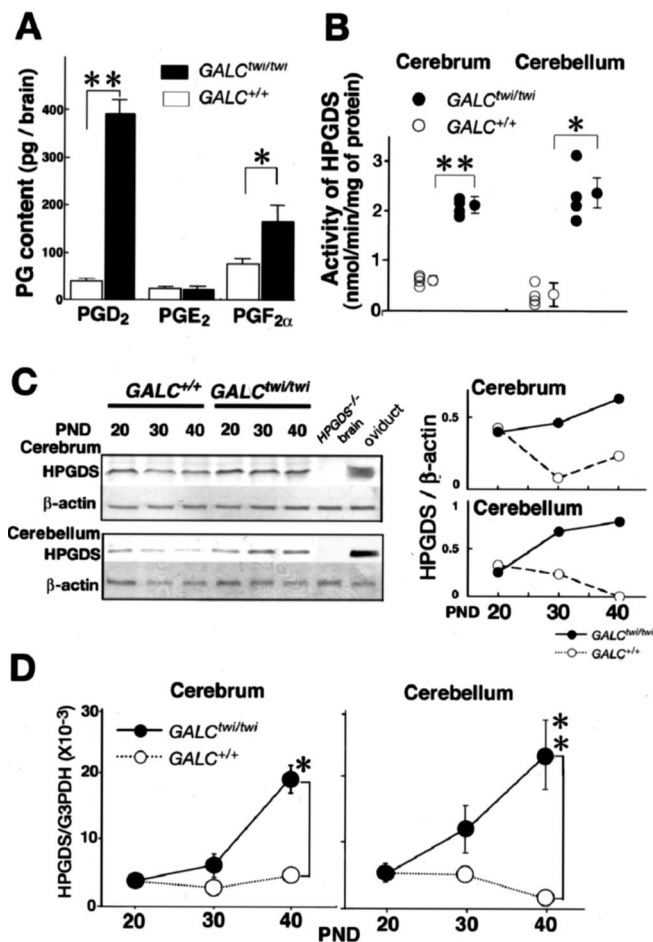
**Statistical analysis.** Values were expressed as the mean ± SE. Data were analyzed by using the two-tailed *t* test; values of *p* < 0.05 were considered to be significant.

## Results

### HPGDS expression is increased with progression of demyelination in *GALC*<sup>twi/twi</sup>

First, we measured the levels of PGD<sub>2</sub>, PGE<sub>2</sub>, and PGF<sub>2α</sub> in the brain of 39-d-old *GALC*<sup>twi/twi</sup> mice (Fig. 1A), in which demyelination was in progress. The PGD<sub>2</sub> content was increased 10-fold in the *GALC*<sup>twi/twi</sup> brain (35.0 ± 9.0 in *GALC*<sup>+/+</sup> vs 390.0 ± 32.0 pg/brain in *GALC*<sup>twi/twi</sup>, mean ± SE; *n* = 6), whereas the other PGs did not show such a drastic increase. That is, the PGE<sub>2</sub> content remained the same (20.0 ± 6.0 vs 19.0 ± 12.0), and that of PGF<sub>2α</sub> increased only twofold (71.0 ± 13.0 vs 160.0 ± 35.0). At P39, the HPGDS enzymatic activity was threefold higher in the *GALC*<sup>twi/twi</sup> cerebrum (0.62 ± 0.05 vs 2.10 ± 0.07 nmol · min<sup>-1</sup> · mg<sup>-1</sup> protein; *n* = 4) and fivefold higher in the *GALC*<sup>twi/twi</sup> cerebellum (0.32 ± 0.10 vs 2.30 ± 0.28; *n* = 4) (Fig. 1B). Western blot analysis also revealed that the level of HPGDS protein in the cerebrum and cerebellum of the *GALC*<sup>twi/twi</sup> brain at P40 was 2- and 10-fold higher, respectively, than that in the same brain regions of the *GALC*<sup>+/+</sup> mice (Fig. 1C).

The level of HPGDS mRNA in the *GALC*<sup>+/+</sup> brains remained constant in the cerebrum and gradually decreased in the cerebel-



**Figure 1.** Increase in PGD<sub>2</sub> in *GALC<sup>twi/twi</sup>* brains together with upregulation of HPGDS. **A**, Quantities of PGD<sub>2</sub>, PGE<sub>2</sub>, and PGF<sub>2α</sub> in fresh-frozen brains of decapitated *GALC<sup>+/+</sup>* (white columns) and *GALC<sup>twi/twi</sup>* (black columns) mice at P39 were determined by enzyme immunoassays. *n* = 6. \**p* < 0.05, \*\**p* < 0.01. **B**, HPGDS activity in the cerebrum and cerebellum of *GALC<sup>+/+</sup>* (white circles) and *GALC<sup>twi/twi</sup>* (black circles) mice at P39 was determined by using 40 μM [<sup>14</sup>C]PGH<sub>2</sub> in the presence of 1 mM glutathione. *n* = 4. \**p* < 0.05, \*\**p* < 0.01. **C**, Western blot analysis of HPGDS and β-actin in the brain from *GALC<sup>+/+</sup>* and *GALC<sup>twi/twi</sup>*. The top and bottom lanes contained the homogenate (5 μg of protein per lane) from the cerebrum and the cerebellum, respectively. The homogenates of the brain of *HPGDS<sup>-/-</sup>* mice and of the oviduct of the wild-type mice were used as the negative and positive control, respectively, of the HPGDS detection. The graphs on the right show the profiles of HPGDS/β-actin amount by morphometrical analysis of Western blots of *GALC<sup>+/+</sup>* and *GALC<sup>twi/twi</sup>* at P20, P30, and P40. **D**, Quantitative PCR analysis of the contents of mRNAs for HPGDS and G3PDH in *GALC<sup>+/+</sup>* and *GALC<sup>twi/twi</sup>* brains was performed by using a LightCycler amplification and detection system. *n* = 4. Data are mean ± SE. \**p* < 0.05, \*\**p* < 0.01. PND, Postnatal day.

lum, whereas it gradually increased in *GALC<sup>twi/twi</sup>* brains after P30. By P40, the mRNA level in *GALC<sup>twi/twi</sup>* was 4.5-fold higher in the cerebrum and 16.5-fold higher in the cerebellum than in the corresponding *GALC<sup>+/+</sup>* brain regions (Fig. 1D).

#### Activated microglia/macrophages in *GALC<sup>twi/twi</sup>* strongly express HPGDS

Immunocytochemical analysis at P40 showed that HPGDS was localized in resting (ramified) microglia in the *GALC<sup>+/+</sup>* cerebellum (Fig. 2A, arrows and inset). In the *GALC<sup>twi/twi</sup>*, the level of HPGDS protein was remarkably increased in activated microglia (Fig. 2B), which possessed irregular thick processes in the region of demyelination. Confocal double immunostaining revealed that ~100% of the HPGDS<sup>+</sup> cells were positive for RCA-1 (Fig. 2C), a marker for microglia and endothelial cells (Szumanska et

al., 1987). RCA<sup>+</sup> endothelial cells were negative for HPGDS (Fig. 2C, arrowhead). All cells positive for F4/80, a marker for activated microglia (Austyn and Gordon, 1981), were also positive for HPGDS (Fig. 2D, arrows). HPGDS<sup>+</sup> cells were positive for neither GFAP, a marker for astrocytes, π-GST, a marker for OLs, nor NG2, a marker for OL precursor (data not shown).

#### DP<sub>1</sub> receptor is induced in hypertrophic astrocytes of *GALC<sup>twi/twi</sup>*

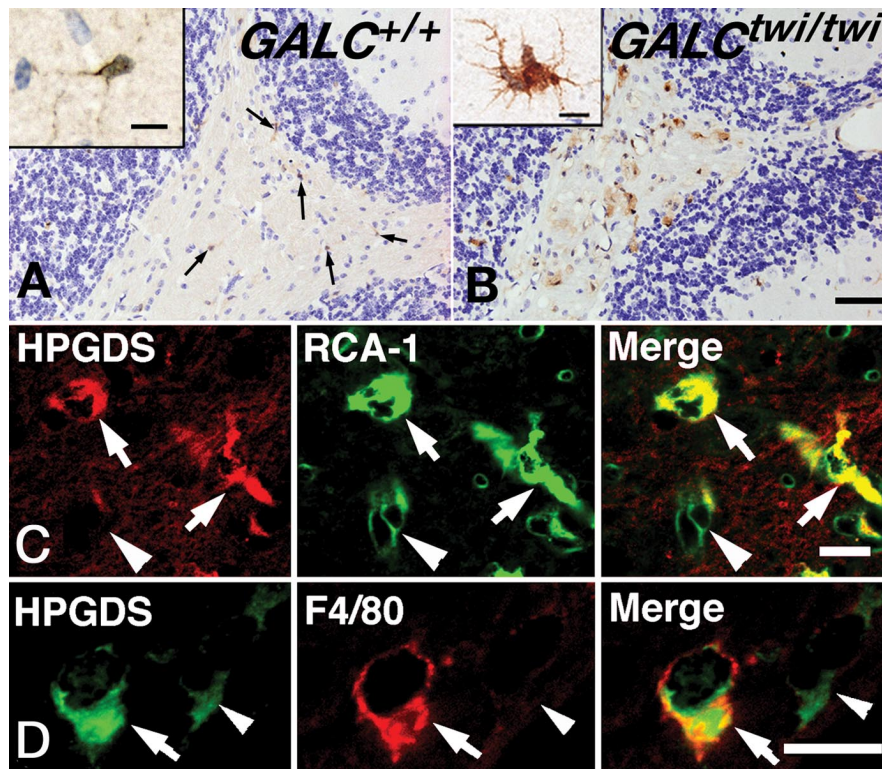
We next quantified the mRNAs of two types of membrane-bound receptors for PGD<sub>2</sub> (i.e., DP<sub>1</sub> and DP<sub>2</sub> receptors), as well as the mRNA of PPARγ, a nuclear receptor for PGD<sub>2</sub> metabolites (Forman et al., 1995; Kliewer et al., 1995). The DP<sub>1</sub> mRNA level remained constant in the cerebrum and the cerebellum of the *GALC<sup>+/+</sup>* mice from P20 through P40 but increased twofold in the cerebrum of the *GALC<sup>twi/twi</sup>* brains and fourfold in their cerebellum at P40 (Fig. 3A). The expression profile of DP<sub>1</sub> mRNA in the cerebrum and cerebellum in the *GALC<sup>twi/twi</sup>* brain during development was thus similar to that of HPGDS (Fig. 1C,D). The contents of DP<sub>2</sub> and PPARγ mRNAs in the brain were ~1/10 and 1/100, respectively, of the content of DP<sub>1</sub> mRNA and did not differ significantly between *GALC<sup>+/+</sup>* and *GALC<sup>twi/twi</sup>* mice at any of the ages examined (Fig. 3A).

Previous studies have revealed that, in the normal mouse, DP<sub>1</sub> receptor immunoreactivity was not detected in the brain except in the arachnoid trabecular cells of the basal forebrain (Mizoguchi et al., 2001). In *GALC<sup>twi/twi</sup>* mice, ~100% of DP<sub>1</sub>-positive cells in brain parenchyma were GFAP-positive astrocytes, and numerous DP<sub>1</sub><sup>+</sup> cells were recognized in the HPGDS-rich areas such as the corpus callosum and cerebellar white matter (CWM) (Fig. 3B) at P40–P45. The regional distribution profile was almost the same as that for HPGDS<sup>+</sup> microglia (Fig. 2B). These DP<sub>1</sub> receptor-positive cells were identified as hypertrophied astrocytes because they possessed large soma with thick-branched processes (Fig. 3B, inset) and expressed GFAP as judged by double immunostaining (Fig. 3C). DP<sub>1</sub> receptor-positive cells were not identical to, but were intermingled with, F4/80<sup>+</sup> microglia in the CWM (Fig. 3D), thus suggesting that DP<sub>1</sub> receptor-positive astrocytes were located close to HPGDS<sup>+</sup> activated microglia. These results suggest that activated microglia interact with hypertrophied astrocytes via PGD<sub>2</sub>/DP<sub>1</sub> signaling in *GALC<sup>twi/twi</sup>* mice.

#### Cultured microglia produce PGD<sub>2</sub> by HPGDS action and cultured astrocytes upregulate GFAP expression when stimulated by DP

Using primary cultures of microglia, we investigated whether these cells produced PGD<sub>2</sub> by HPGDS catalysis. As shown in Figure 4A, after treatment with 5 μM A23187, a Ca<sup>2+</sup> ionophore, PGD<sub>2</sub> release was increased ~23-fold compared with that of controls, whereas PGE<sub>2</sub> release was increased only 3.8-fold, and that of PGF<sub>2α</sub> remained unchanged. PGD<sub>2</sub> production was suppressed dose dependently by treatment with HQL-79, a specific inhibitor of HPGDS (Matsushita et al., 1998).

To investigate the effect of PGD<sub>2</sub> on astrocyte activation, we determined the contents of DP<sub>1</sub> and DP<sub>2</sub> receptor mRNAs in cultured mouse astrocytes before and after stimulation with selective agonists for DP<sub>1</sub> [BW245C (Sharif et al., 2000)] and DP<sub>2</sub> [DK-PGD<sub>2</sub> (Gervais et al., 2001)] (Fig. 4B). Astrocytes in primary culture contained mRNAs of both DP<sub>1</sub> and DP<sub>2</sub> receptors in almost equal amounts. The DP<sub>1</sub> receptor mRNA content was increased after stimulation with either agonist in a dose-dependent manner from 1 to 100 nM and reached a level ~1.5-fold higher than that without stimulation, whereas the DP<sub>2</sub> re-



**Figure 2.** Upregulation of HPGDS in microglia of *GALC*<sup>twi/twi</sup> brains at P45. **A, B**, HPGDS immunostaining visualized by using DAB in the cerebellum of *GALC*<sup>+/+</sup> (**A**) and *GALC*<sup>twi/twi</sup> (**B**). The inset and arrows in **A** indicate resting HPGDS<sup>+</sup> microglia. Scale bars: **B**, 50  $\mu$ m; insets, 10  $\mu$ m. **C, D**, Confocal images for HPGDS (red in **C** and green in **D**) and microglial markers, i.e., RCA-1 (green in **C**) and F4/80 (red in **D**). Arrows in **C** and **D** point to HPGDS<sup>+</sup> microglia, and arrowheads in **C** point to RCA<sup>+</sup> HPGDS<sup>-</sup> endothelial cells, judged from the morphology. Arrowheads in **D** indicate HPGDS-positive/F4/80-negative resting microglia. Scale bars: **C, D**, 10  $\mu$ m.

ceptor mRNA content remained almost unchanged after stimulation with either agonist. The GFAP mRNA content was also increased in a dose-dependent manner from 1 to 100 nM after stimulation with either agonist, becoming twofold higher than that in the untreated control cultures.

#### Inhibition of HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> receptor signaling alleviates astrogliosis of *GALC*<sup>twi/twi</sup>

To assess the contribution of HPGDS and the DP<sub>1</sub> receptor to astrogliosis in *GALC*<sup>twi/twi</sup> mice, we generated *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (supplemental Fig. 1A, available at www.jneurosci.org as supplemental material) and *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (supplemental Fig. 1B, available at www.jneurosci.org as supplemental material) mice and then compared the mRNA content and the immunocytochemical profile of GFAP in the brain of these double-mutant mice at P45 with those of *GALC*<sup>twi/twi</sup> mice. The content of psychosine, a potent neurotoxin and an accumulated substrate of the missing enzyme galactosylceramidase in *GALC*<sup>twi/twi</sup>, was not significantly different among *GALC*<sup>twi/twi</sup>, *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, and *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> in the cerebrum (8.86 ± 0.45, 8.65 ± 2.47, and 9.07 ± 3.56 ng/mg protein, respectively; *n* = 4) or cerebellum (25.12 ± 2.21, 29.59 ± 4.94, and 35.78 ± 8.98 ng/mg protein, respectively; *n* = 4). Therefore, we consider the gliotic factors, with respect to the primary metabolic defect, to be uniform in all strains.

GFAP immunostaining at P45 revealed that only weakly GFAP<sup>+</sup> astrocytes with thin and delicate processes (Fig. 5A, inset) were present in both the cerebrum and cerebellum (Fig. 5A, E) of *GALC*<sup>+/+</sup> mice. In *GALC*<sup>twi/twi</sup> mice, intensely GFAP<sup>+</sup>

astrocytes with enlarged soma and many thick processes (Fig. 5B, inset) were numerous and diffusely distributed in both cerebrum and cerebellum (Fig. 5B, F). In *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (Fig. 5C, G) and *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (Fig. 5D, H) mice, the GFAP immunoreactivity was slightly more intense than that in *GALC*<sup>+/+</sup> mice but remarkably weaker than that in *GALC*<sup>twi/twi</sup> mice, and the processes of GFAP<sup>+</sup> astrocytes were clearly thinner than those in the *GALC*<sup>twi/twi</sup> mice. The suppression of astrogliosis was more remarkable in *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> than in *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> mice. Quantitative RT-PCR and Northern blot analyses revealed that the GFAP mRNA content in the cerebellum at P40 was decreased to 15 and 43% of that of *GALC*<sup>twi/twi</sup> mice in *HPGD*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> and *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> mice, respectively (Fig. 5I).

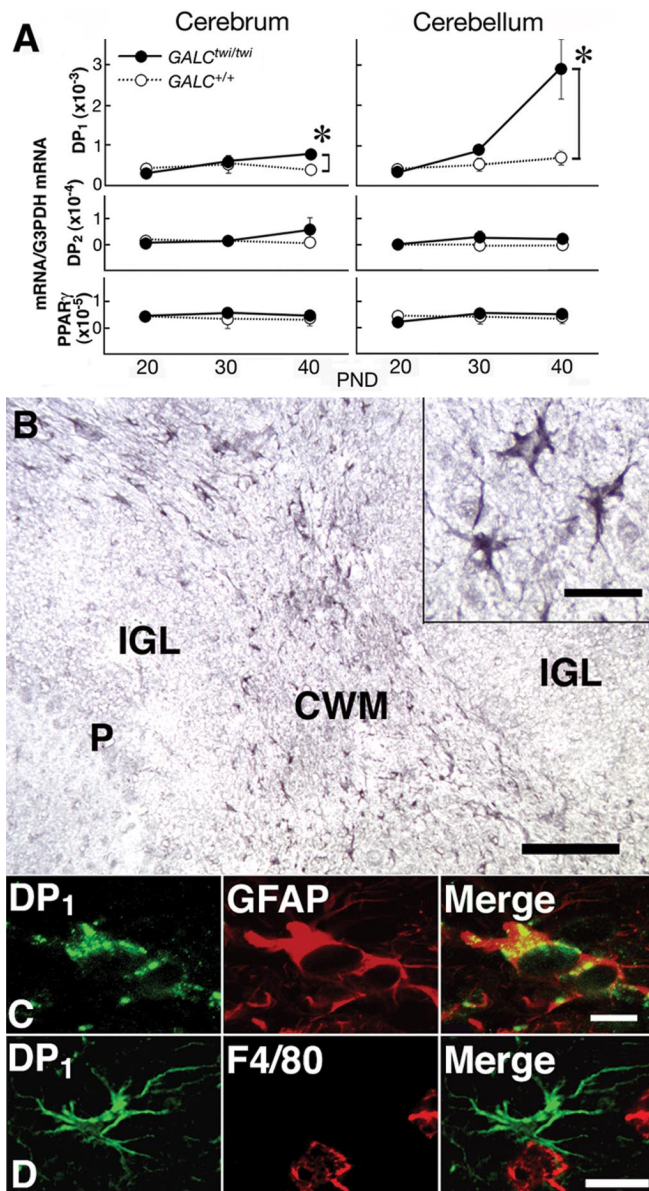
Furthermore, when we treated *GALC*<sup>twi/twi</sup> mice with HQL-79, the astrogliosis of *GALC*<sup>twi/twi</sup> mice (Fig. 6A, B) was markedly suppressed in both the cerebrum (Fig. 6D) and cerebellum (Fig. 6E). DP<sub>1</sub> receptor-positive astrocytes in the *GALC*<sup>twi/twi</sup> mice (Fig. 6C) were also decreased in number after the HQL-79 treatment (Fig. 6F). Quantitative RT-PCR and Northern blot analyses (Fig. 6G) revealed that the GFAP mRNA content in the *GALC*<sup>twi/twi</sup> cerebellum at P45 was decreased to 29% of the control

by the inhibitor treatment.

#### Inhibition of HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> receptor signaling suppresses apoptosis of OLs and demyelination of *GALC*<sup>twi/twi</sup>

MBP immunostaining revealed that demyelination was severe in the *GALC*<sup>twi/twi</sup> brain (Fig. 7A), yet myelin was well conserved in the brains of *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, and HQL-79-treated *GALC*<sup>twi/twi</sup> (Fig. 7B–D). Western blotting analysis for MBP of these mice confirmed the immunocytochemical observation (Fig. 7E, F). As shown in Figure 7F, relative amounts of MBP were significantly retained in the brains of *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (*n* = 5), and HQL-79-treated *GALC*<sup>twi/twi</sup> (*n* = 7) mice. Especially MBPs of 18.5 and 14 kDa, which are expressed in the early phase of myelination, were better preserved in *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, *DP1*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> and HQL-79-treated *GALC*<sup>twi/twi</sup> than those of 21.5 and 17 kDa MBPs, expressed in the later phase. As shown in Figure 7G, the number of  $\pi$ -GST-positive TUNEL-positive cells, which indicate apoptotic OLs, was significantly decreased in *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> and HQL-79-treated *GALC*<sup>twi/twi</sup> (Fig. 7G). To determine whether proinflammatory molecule levels were changed in those mice, we measured the mRNAs for several proinflammatory molecules, such as iNOS, IL-1 $\beta$ , IL-6, and TNF $\alpha$ . The level of iNOS, but not that of IL-1 $\beta$ , IL-6, or TNF $\alpha$ , was significantly reduced in *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> and HQL-79-treated *GALC*<sup>twi/twi</sup> animals (*n* = 4) (Fig. 7H).

These results indicate that the genetic or pharmacological blockade of the HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> signaling pathway not only

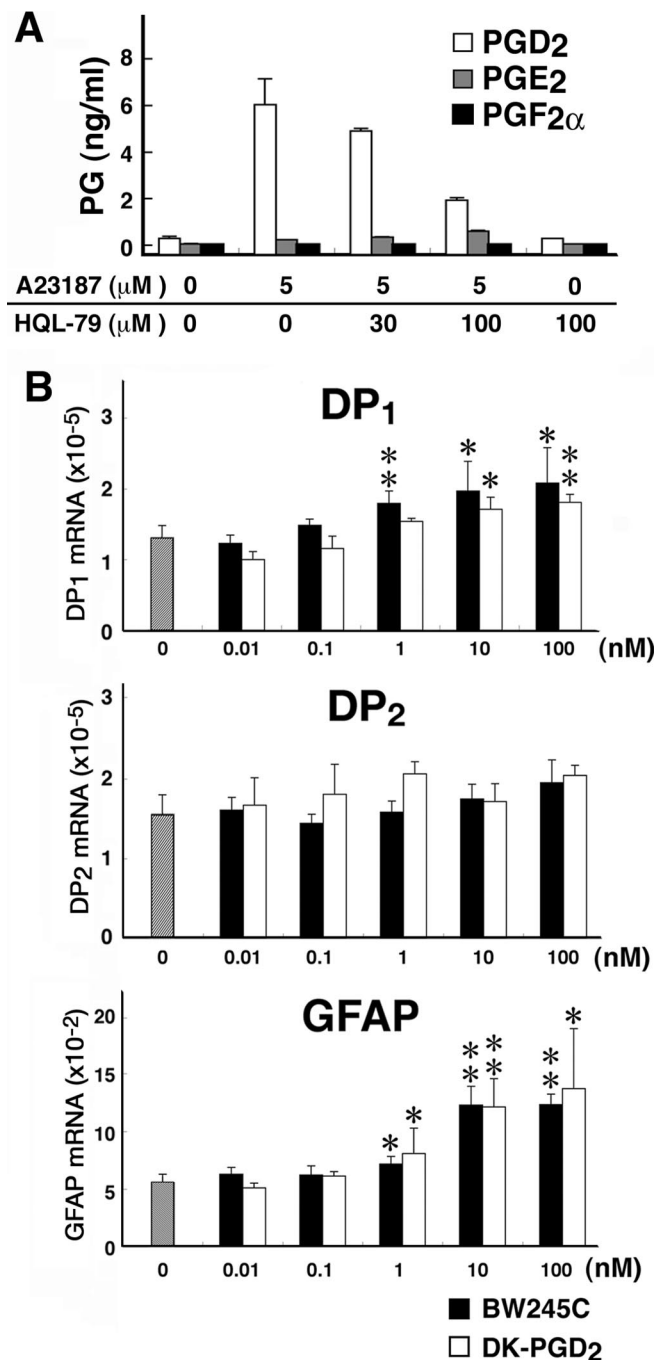


**Figure 3.** Induction of DP<sub>1</sub> in *GALC<sup>twi/twi</sup>* hypertrophic astrocytes. **A**, Quantitative PCR analysis of the contents of mRNAs for DP<sub>1</sub>, DP<sub>2</sub>, PPAR $\gamma$ , and G3PDH mRNA in *GALC<sup>+/+</sup>* (white circles) and *GALC<sup>twi/twi</sup>* (black circles) brains was performed by using a LightCycler amplification and detection system.  $n = 4$ . \* $p < 0.05$ . PND, Postnatal day. **B**, DP<sub>1</sub> immunostaining in the *GALC<sup>twi/twi</sup>* cerebellum at P45. DP<sub>1</sub><sup>+</sup> cells are recognized in the CWM but not in the internal granular layer (IGL) or Purkinje cell layer (P). Scale bar, 100  $\mu$ m. Inset, High-power view of DP<sub>1</sub><sup>+</sup> cells. Scale bar, 10  $\mu$ m. **C**, Double immunostaining for DP<sub>1</sub> (green) and GFAP (red) in hypertrophied astrocytes. Scale bar, 5  $\mu$ m. **D**, DP<sub>1</sub><sup>+</sup> astrocytes (green) and F4/80<sup>+</sup> microglia (red) are intermingled in the CWM, although DP<sub>1</sub> and F4/80 are not detectable in the same cells. Scale bar, 10  $\mu$ m.

suppressed the hypertrophic change in astrocytes or astrogliosis in *GALC<sup>twi/twi</sup>* mice but also suppressed iNOS expression and subsequent enhancement of demyelination.

#### Inhibition of HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> receptor signaling diminishes symptomatology of *GALC<sup>twi/twi</sup>*

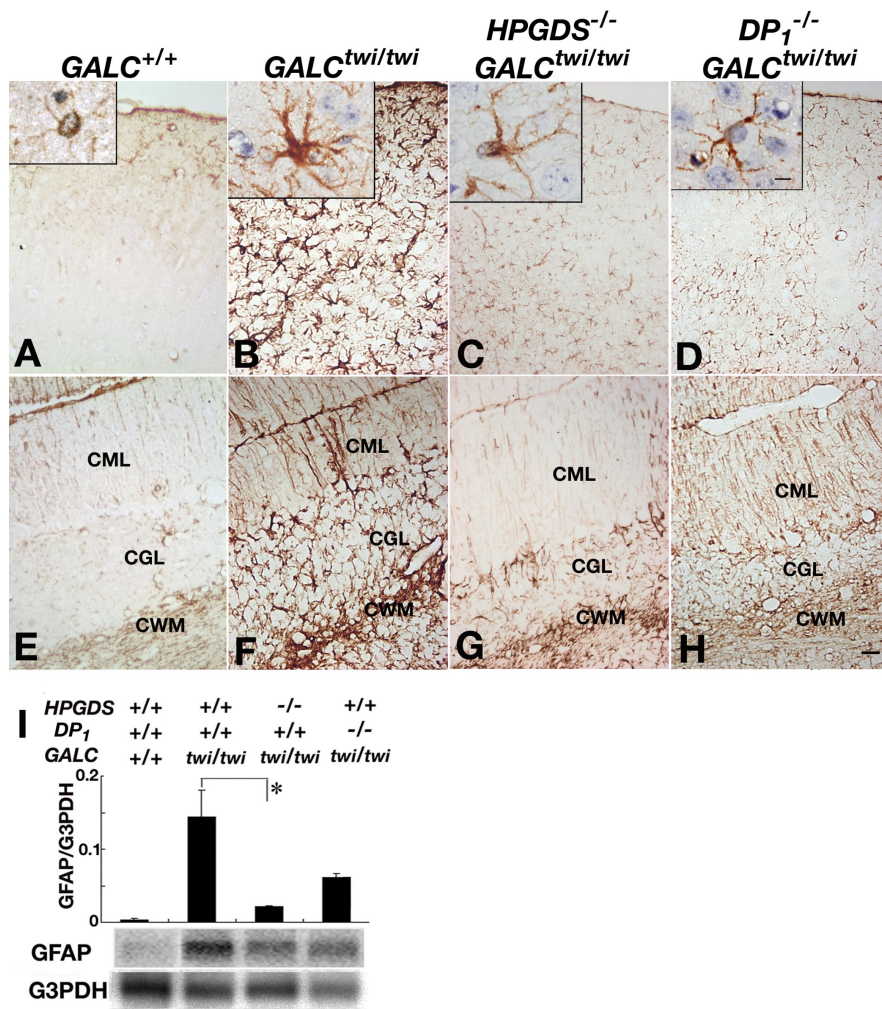
The lifespan of *HPGDS<sup>-/-</sup>GALC<sup>twi/twi</sup>* and *DP<sub>1</sub><sup>-/-</sup>GALC<sup>twi/twi</sup>* mice ( $47.3 \pm 4.5$  and  $46.2 \pm 0.5$  d, respectively) was slightly longer than that of *GALC<sup>twi/twi</sup>* mice ( $45.5 \pm 0.4$  d) despite the fact that the severe peripheral neuropathy, which primarily determines the lifespan of *GALC<sup>twi/twi</sup>* mice, was observed across all



**Figure 4.** **A**, PG production by primary cultures of microglia 30 min after treatment with A23187 in the absence or presence of HQL-79. After incubation in DMEM without FBS for 6 h, the microglia were treated with 5 mM A23187 for 30 min. HQL-79 was added 15 min before A23187 treatment. PG contents in the medium were quantified by enzyme immunoassays. White columns, PGD<sub>2</sub>; gray columns, PGE<sub>2</sub>; black columns, PGF<sub>2 $\alpha$</sub> .  $n = 3$ . **B**, mRNA levels of DP<sub>1</sub>, DP<sub>2</sub>, and GFAP before (gray column) and after stimulation of primary cultures of astrocytes with BW245C (black columns) or DK-PGD<sub>2</sub> (white columns). Once the astrocytes in the primary culture reached confluence, the cells were rinsed with PBS and subcultured at  $5 \times 10^5$  cells per well (6-well plate). After a 24 h incubation in DMEM containing 1% FBS, the astrocytes were incubated with BW245C (0.1–100 nM) or DK-PGD<sub>2</sub> (0.1–100 nM) for 6 h at 37°C.  $n = 3$ . \* $p < 0.05$ , \*\* $p < 0.01$ .

three groups of mice. The lifespan of HQL-79-treated *GALC<sup>twi/twi</sup>* mice was significantly longer than that of the vehicle-treated mice ( $47.8 \pm 1.3$  and  $44.0 \pm 0.7$  d, respectively;  $n = 5$ ;  $p < 0.03$ ).

Spasticity and ataxic symptoms of *HPGDS<sup>-/-</sup>GALC<sup>twi/twi</sup>* and *DP<sub>1</sub><sup>-/-</sup>GALC<sup>twi/twi</sup>* mice at P45 were remarkably milder than



**Figure 5.** Alleviated astrogliosis in the mutant *twitcher* lacking HPGDS or DP<sub>1</sub> gene. **A–H**, GFAP immunostaining in the cerebrum (**A–D**) and the cerebellum (**E–H**) of *GALC*<sup>+/+</sup> (**A, E**), *GALC*<sup>twi/twi</sup> (**B, F**), *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> (**C, G**), and *DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>twi/twi</sup> (**D, H**) mice. Insets show higher-magnification view of GFAP-immunoreactive astrocytes for each genotype. Scale bars: **H**, 50 μm; insets, 5 μm. CGL, Cerebellar granular layer; CML, cerebellar molecular layer. **I**, Quantitative RT-PCR (top) and Northern blot analyses (bottom) for GFAP and G3PDH in *GALC*<sup>+/+</sup>, *GALC*<sup>twi/twi</sup>, *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, and *DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>twi/twi</sup>. The quantitative PCR analysis of the contents of mRNAs for GFAP and G3PDH was performed by using a LightCycler amplification and detection system. For Northern blotting, total RNA (10 μg) was electrophoresed in an agarose gel, transferred to nylon membranes, and hybridized with <sup>32</sup>P-labeled cDNA probes specific for mouse GFAP and G3PDH. *n* = 3–5. \**p* < 0.05.

those of *GALC*<sup>twi/twi</sup> mice. *GALC*<sup>twi/twi</sup> mice barely staggered with remarkable intentional tremor, whereas *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> mice moved about noticeably faster at P45. When suspended by the tail, *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup> animals retained the righting reflex, exhibiting a milder tremor (supplemental movie A, available at www.jneurosci.org as supplemental material). Clinical symptoms including spasticity and ataxic symptoms were also milder in HQL-79-treated *GALC*<sup>twi/twi</sup> than in vehicle-treated *GALC*<sup>twi/twi</sup> mice. HQL-79-treated *GALC*<sup>twi/twi</sup> moved around smoothly at P45 and retained the righting reflex with milder tremor when suspended by the tail (supplemental movie B, available at www.jneurosci.org as supplemental material).

The main cause of the death of *GALC*<sup>twi/twi</sup> mice is peripheral and cranial nerve palsy, which leads to difficulty in eating and limits the lifespan. MBP immunostaining (Fig. 7A–D) and electron microscopy (data not shown) revealed that demyelination was attenuated in the CNS, but not in the sciatic nerve, of *HPGDS*<sup>-/-</sup>*GALC*<sup>twi/twi</sup>, *DP*<sub>1</sub><sup>-/-</sup>*GALC*<sup>twi/twi</sup>, and HQL-79-treated *GALC*<sup>twi/twi</sup> mice. A similar discrepancy between CNS and

PNS has been reported previously in the case of *GALC*<sup>twi/twi</sup> mice receiving bone-marrow transplantation (Toyoshima et al., 1986; Ichioka et al., 1987). Furthermore, Escobar et al. (2005) recently reported that transplantation of umbilical-cord blood into babies with infantile Krabbe’s disease resulted in improvement in the CNS but not in the PNS.

Using immunocytochemistry, we found that DP<sub>1</sub> receptor expression was upregulated in the CNS but not detected in the sciatic nerve of *GALC*<sup>twi/twi</sup> (data not shown), suggesting that PGD<sub>2</sub> may not be involved in progression of the pathology of the PNS of *GALC*<sup>twi/twi</sup>. This coincides with a distinction between the CNS and PNS pathologies in *twitcher* in which Schwann cells, unlike OLs, proliferate in response to demyelination. Furthermore, apoptotic death does not appear to occur in the peripheral nerves (Komiya and Suzuki, 1992). The demyelination mechanism, therefore, appears to be distinct between CNS and PNS, particularly regarding the contribution of support cells as well as PGD<sub>2</sub> to the demyelination.

### Discussion

#### PGD<sub>2</sub>-mediated enhancement of microglial activation and astrogliosis during neuroinflammation

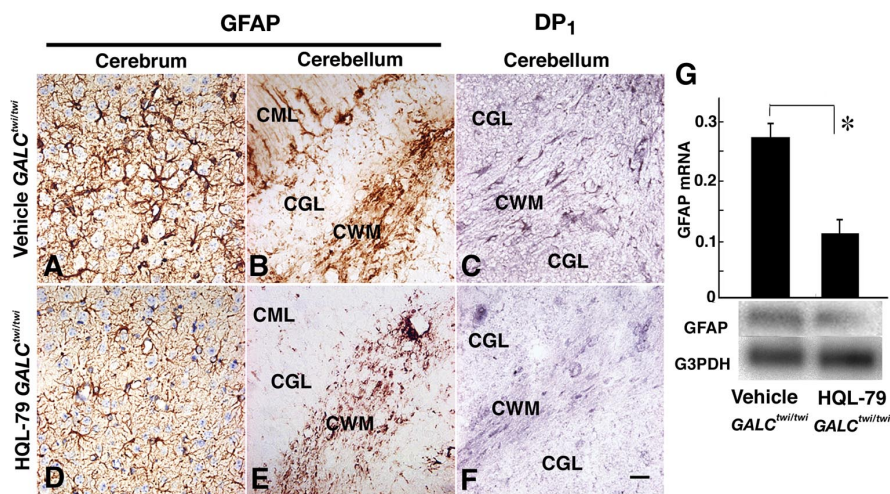
Microglia are believed to act primarily as a line of defense for the brain, because they express MHC molecules, secrete proinflammatory cytokines (Woodroffe et al., 1991), and produce NO (Wood et al., 1994) and PGs (Gebicke-Haerter et al., 1989; Minghetti and Levi, 1995; Bauer et al., 1997). Cyclooxygenases (COXs) are the key enzymes for the production of PGs and are upregulated in the CNS in various neurodegenerative diseases, including Alzheimer’s disease (Yermakova et al., 1999), Parkinson’s disease (Teismann et al., 2003), prion disease (Walsh et al., 2000; Deininger et al., 2003), spinal cord injury (Schwab et al., 2000), and ischemia (Govoni et al., 2001). Several investigators have reported selective COX inhibitors to be neuroprotective (McGeer and McGeer, 1995; Stewart et al., 1997; Pasinetti, 1998; Lim et al., 2000; Govoni et al., 2001), suggesting that some classes of PGs may play important roles in neurodegeneration. Positive immunoreactivities for COX-1 and COX-2 are mostly observed in microglia (Pasinetti and Aisen, 1998; Yermakova et al., 1999; Walsh et al., 2000). Several reports have demonstrated that microglia produce PGD<sub>2</sub> and PGE<sub>2</sub> *in vitro* (Gebicke-Haerter et al., 1989; Minghetti and Levi, 1995; Bauer et al., 1997). Between these two prostanoids, PGE<sub>2</sub> has been proposed to act as a mediator in inflammatory processes (Petrova et al., 1999; Kim et al., 2001) of neurodegenerative diseases such as amyotrophic lateral sclerosis (Ilzecka, 2003), whereas the direct mechanism of PGD<sub>2</sub> involvement in neurodegenerative diseases has not been well studied.

The pathological phenotypes within the CNS of HPGDS- or DP<sub>1</sub>-null *twitcher* mice are quite different from those seen in their peripheral immune responses. We reported previously that DP<sub>1</sub><sup>-/-</sup> mice used in the ovalbumin-induced asthma model exhibited a remarkably decreased accumulation of lymphocytes and proinflammatory cytokines in their lungs compared with wild-type mice (Matsuoka et al., 2000). In contrast, HPGDS<sup>-/-</sup> mice used in this same model did not show any changes in the lymphocyte and cytokine components in their lungs (K. Aritake and Y. Urade, unpublished results). Together, these results suggest that the above functional peripheral defect of DP<sub>1</sub><sup>-/-</sup> mice may not be simply caused by the deficiency of DP<sub>1</sub> signaling but may be attributable to the shunting of PGD<sub>2</sub> signaling to the DP<sub>2</sub> pathway. Based on these results, we believe that the mild gliosis and demyelination in the HPGDS- and DP<sub>1</sub>-null *twitcher* mice are mainly caused by the suppression of neuroinflammation around the HPGDS-expressing microglia and DP<sub>1</sub>/DP<sub>2</sub>-possessing astrocytes, although it is possible that decreased cell infiltration into the CNS in these mice leads to the suppression of secondary gliosis.

We demonstrated here that PGD<sub>2</sub> was actively produced in the GALC<sup>twi/twi</sup> brain (Fig. 1A) by HPGDS upregulated in activated microglia (Figs. 2, 4A). In contrast, the level of PGE<sub>2</sub> was unchanged in the GALC<sup>twi/twi</sup> brain when compared with that in the GALC<sup>+/+</sup> brain (Fig. 1A), and microsomal PGE synthase was not induced in the GALC<sup>twi/twi</sup> brain, as judged from the results of quantitative RT-PCR and immunocytochemistry (data not shown). Furthermore, only a small amount of PGE<sub>2</sub> release was observed in cultured microglia after activation (Fig. 4A). Therefore, PGD<sub>2</sub> is thought to play more important roles than PGE<sub>2</sub> in the processes of neuroinflammation that induce microglia activation and astrogliosis.

In addition, we demonstrated that astrocytes had DP receptors and that PGD<sub>2</sub> stimulation accelerated astrogliosis (Figs. 3, 4B). We also showed that inhibition of HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> receptor signaling by genetic depletion of HPGDS or DP<sub>1</sub>, or administration of HQL-79, reduced astrogliosis and ameliorated the clinical symptoms of GALC<sup>twi/twi</sup>. As shown in Figure 4B, cultured astrocytes expressed both DP<sub>1</sub> and DP<sub>2</sub> receptors and increased GFAP production when stimulated by either a DP<sub>1</sub> or DP<sub>2</sub> agonist. This may explain the finding that the astrogliosis of HPGDS<sup>-/-</sup> GALC<sup>twi/twi</sup> mice was milder than that of DP<sub>1</sub><sup>-/-</sup> GALC<sup>twi/twi</sup> mice, in which PGD<sub>2</sub>-induced gliosis progressed in a DP<sub>2</sub> receptor-mediated manner. Conversely, in HPGDS<sup>-/-</sup> GALC<sup>twi/twi</sup> mice, PGD<sub>2</sub> was not produced in the neuroinflammatory loci, signal transduction through both DP<sub>1</sub> and DP<sub>2</sub> receptors was blocked, and astrocytic activation was minimal.

In this study, we unexpectedly found that blockade of the HPGDS/PGD<sub>2</sub> signaling resulted in suppressed apoptosis of OLs (Fig. 7G). This suppression may have contributed to the amelioration of demyelination in HPGDS<sup>-/-</sup> GALC<sup>twi/twi</sup> and HQL-79-treated GALC<sup>twi/twi</sup> mice. However, DP<sub>1</sub><sup>-/-</sup> GALC<sup>twi/twi</sup> mice exhibited milder symptoms and their demyelination was clearly suppressed, as judged from the results of the immunohistochemical and Western blot analyses (Fig. 7C, E, F) compared with the



**Figure 6.** HPGDS-inhibitor alleviated astrogliosis via inhibition of DP<sub>1</sub>. GALC<sup>twi/twi</sup> mice received a daily subcutaneous injection of 50 mg · kg<sup>-1</sup> · d<sup>-1</sup> HQL-79 from P25 to P45. As a control, the same volume of vehicle was injected into GALC<sup>twi/twi</sup> during the same period. **A–C**, Vehicle-treated GALC<sup>twi/twi</sup>. **D–F**, HQL-79-treated GALC<sup>twi/twi</sup>. **A, B, D, E**, GFAP immunostaining. **C, F**, DP<sub>1</sub> immunostaining. Scale bar, 50 μm. **G**, Quantitative RT-PCR (top) and Northern blot analyses (bottom) for vehicle-treated and HQL-79-treated GALC<sup>twi/twi</sup>. *n* = 3. \**p* < 0.05. CGL, Cerebellar granular layer; CML, cerebellar molecular layer.

GALC<sup>twi/twi</sup> mice, despite the fact that the number of TUNEL-positive cells was not significantly decreased (Fig. 7G). Therefore, secondary demyelination caused by neuroinflammation might also have been suppressed in the DP<sub>1</sub><sup>-/-</sup> GALC<sup>twi/twi</sup> mice, leading to decreased spasticity and improved ataxic symptoms.

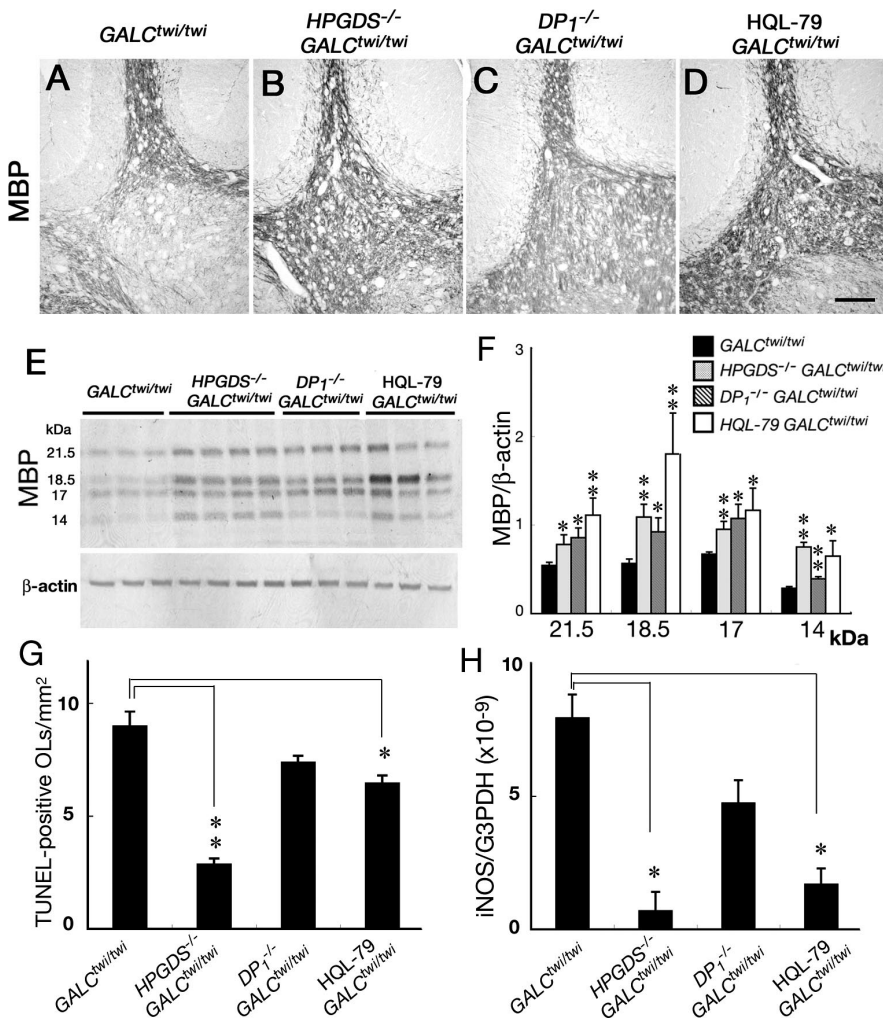
#### Predicted mechanism of the HPGDS/PGD<sub>2</sub>/DP<sub>1</sub> signaling pathway leading to enhanced astrogliosis and demyelination

Based on the findings described in this present study, we hypothesize the molecular events leading to neuroinflammation in the GALC<sup>twi/twi</sup> brains to be the following (supplemental Fig. 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). First, myelin and/or OL debris produced by the apoptotic death of OLs in the GALC<sup>twi/twi</sup> brain (Taniike et al., 1999) activates microglia to synthesize PGD<sub>2</sub> through the action of HPGDS. The PGD<sub>2</sub> produced by these activated microglia then stimulates both DP<sub>1</sub> and DP<sub>2</sub> receptors on neighboring astrocytes in a paracrine manner and upregulates the expression of DP<sub>1</sub>, but not that of DP<sub>2</sub>, receptors on these astrocytes. The activation of DP<sub>1</sub>, a G<sub>s</sub>-coupled receptor (Hirata et al., 1994), increases the intracellular cAMP level in the astrocytes to induce hypertrophy of these cells. DP<sub>2</sub> stimulation also reportedly elevates the intracellular Ca<sup>2+</sup> level. Engagement of DP<sub>2</sub>, a G<sub>αi</sub>-coupled receptor (Hirai et al., 2001), also increases GFAP expression, presumably through elevation of the intracellular Ca<sup>2+</sup> level.

We reported previously that OLs express lipocalin-type PGD synthase (L-PGDS) but not HPGDS (Taniike et al., 2002). Although L-PGDS and HPGDS catalyze the same reaction to produce PGD<sub>2</sub>, these two enzymes have quite different roles within the context of demyelination. In L-PGDS<sup>-/-</sup> GALC<sup>twi/twi</sup>, both OL apoptosis and gliosis were enhanced (Taniike et al., 2002), unlike the case of HPGDS<sup>-/-</sup> GALC<sup>twi/twi</sup>, in which gliosis was ameliorated. Therefore, we hypothesize that L-PGDS localized in OLs plays a protective role in *twitcher* mice and that HPGDS, upregulated in activated microglia, is involved in the secondary pathology in this disease model.

The instigation of the pathological condition in *twitcher* is attributable to the accumulation of psychosine, yet it has been reported that inflammatory cells and many cytokines are also involved in the progression of the disease (Wu et al., 2000). Al-





**Figure 7.** Demyelination suppressed via inhibition of neuroinflammation in *GALC<sup>twi/twi</sup>*. **A–D**, MBP immunostaining. Cerebellum of mice at P45. Scale bar, 100  $\mu$ m. **E**, Western blot analysis of MBP (top) and  $\beta$ -actin (bottom). The top and bottom lanes contained the homogenate (2  $\mu$ g of protein per lane) from the cerebellum. **F**, Morphometrical analysis of MBP protein content in the brain of *GALC<sup>twi/twi</sup>* (black columns;  $n = 4$ ), *HPGDS<sup>-/-</sup> GALC<sup>twi/twi</sup>* (gray columns;  $n = 5$ ), *DP1<sup>-/-</sup> GALC<sup>twi/twi</sup>* (hatched columns;  $n = 5$ ), and HQL-79-treated *GALC<sup>twi/twi</sup>* (white columns;  $n = 7$ ). \* $p < 0.05$ , \*\* $p < 0.01$ . **G**, TUNEL histochemistry combined with immunocytochemistry for  $\pi$ -GST was performed, and  $\pi$ -GST-positive TUNEL-positive cells were counted in the basal ganglia of the brains of paraffin sections prepared from mice at P45 ( $n = 4$  each). \* $p < 0.05$ , \*\* $p < 0.01$ . **H**, Quantitative analysis of iNOS mRNA in the cerebellum at P45.  $n = 4$ . \* $p < 0.05$ .

though the toxicity of psychosine plays a significant role in the apoptotic death of OLs and in preliminary demyelination, the ensuing pathology is intensified by cytokines and chemokines released by activated microglia and possibly by astrocytes as well. In this sense, the demyelinating process closely resembles that of pattern III of MS, in which the lesions contain an inflammatory cell infiltration, activated microglia, and OL apoptosis but without any deposition of Ig and complement (Lucchinetti et al., 2000). Considering this, along with reports that MHC class II-null *twitcher* mice show only mild pathology (Matsushima et al., 1994) and that inhibiting the proinflammatory cytokine TNF $\alpha$  suppresses demyelination and apoptosis of OLs in *GALC<sup>twi/twi</sup>* (Kagitani-Shimono et al., 2005), we believe that secondary neuroinflammation is a critical component in the progression of the degenerative condition found in *twitcher*. Furthermore, because HPGDS inhibition in *GALC<sup>twi/twi</sup>* significantly suppressed the disease process, including apoptosis of OLs, it is reasonable to surmise that neuroinflammation in *twitcher* mice exacerbates demyelination and functional deficits.

Germane to our contention is identifying the physiological role of astrogliosis within the disease model. Although still a subject not yet clarified despite intense discussion, we speculate that hypertrophic astrocytes may be related to neuroinflammation through their production of cytokines. This study, then, represents the first example of a PGD<sub>2</sub>-mediated microglia/astrocyte interaction in neuroinflammation. Here, PGD<sub>2</sub> was shown to be an important mediator of neuroinflammation, and it was demonstrated that the HPGDS inhibitor HQL-79, an orally active compound remarkably effective at attenuating neuroinflammation-associated astrogliosis, was an able attenuating substance. Our present results encourage additional study into both the role of PGD<sub>2</sub> in inflammatory and demyelinating diseases as well as the use of HPGDS inhibitors as novel strategies for anti-neuroinflammatory therapy.

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