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Microglia have a protective role in viral encephalitis-induced seizure development and hippocampal damage

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1. Introduction

Viral encephalitis is the most common form of brain infection and can cause severe and life-threatening consequences, including epilepsy (Vezzani et al., 2016). Monocyte infiltration and activation of microglia, the resident macrophage population of the brain, are hallmarks of CNS inflammation, including viral infection (Prinz and Priller, 2017). However, the role of these cells in viral clearance and immunopathology is not well defined. At least in part this is due to the problems of differentiating invading monocytes from activated microglia in the brain and the lack of selective tools to manipulate these two types of myeloid cells. In several viral brain infections, activated microglia appears to be involved in inhibition of viral replication and in neurotoxicity, indicating the dual nature of microglia: they contribute to the defense of the CNS but may also bear responsibility for CNS damage (Rock et al., 2004). Microglia expresses various Pattern Recognition Receptors (PRRs) to identify viral signatures called Pathogen Associated Molecular Patterns (PAMPs). Upon stimulation by PAMPs microglia respond by releasing several pro- and anti-inflammatory cytokines such as monocyte chemoattractant protein 1 (MCP1/CCL2), interleukin (IL)-1β, type I interferon (IFN), IFNγ, and tumor necrosis factor (TNF)α (O’Shea et al., 2013).

Recently, a new compound, PLX5622, which acts as specific dietary inhibitor of colony stimulating factor-1 receptor (CSF1R), a tyrosine kinase transmembrane receptor essential for the survival and activation of monocytes and macrophages in the periphery and microglia in the CNS (Erblich et al., 2011; Hamilton and Achuthan, 2013; Elmore et al., 2015), was shown to efficiently eliminate microglia in mice (Dagher et al., 2015). Similar to the less specific CSF1R inhibitor PLX3397 (Elmore et al., 2015), PLX5622 has been used to study the role of microglia in several conditions of brain injury (Acharya et al., 2016; Feng et al., 2016; Beckers et al., 2017; Hilla et al., 2017). Two recent studies...
reported that microglia depletion by PLX5622 increases virus replication and subsequent mortality in response to intracerebral infection with mouse hepatitis virus (MHV; Wheeler et al., 2018) or flaviviruses (West Nile virus and Japanese encephalitis virus; Seitz et al., 2018).

We used PLX5622 in a model of viral encephalitis in mice, in which Thelier’s virus (also termed Thelier’s murine encephalomyelitis virus (TMEV)), a naturally occurring enteric pathogen of the mouse, is used for intracerebral infection (Vezzani et al., 2016; DePaula-Silva et al., 2017). Disease development following TMEV infection depends on the mouse strain. The SJL/J mouse strain develops a multiple sclerosis-like progressive T cell-mediated autoimmune demyelinating disease, characterized by weakness of the hind limbs, which advances to a severe spastic paralysis, and inflammatory demyelination in the spinal cord, while the C57BL/6J (B6) mouse strain develops hippocampal damage and acute seizures, which progress to epilepsy (DePaula-Silva et al., 2017). The striking difference between the mouse strains seems to be partially due to the strong antiviral cytotoxic CD8+ T lymphocyte response observed in B6 mice, which is suppressed by the elevated induction of regulatory CD4+ T cells (Tregs) in SJL/J mice, resulting in viral persistence and demyelination (DePaula-Silva et al., 2017). We now report that elimination of microglia strikingly changes the disease phenotype of TMEV-infected B6 mice, indicating that microglia is much more important in this model of viral encephalitis than previously thought.

2. Materials and methods

2.1. Animals

Four-week-old female JAX® C57BL/6J (B6) mice were purchased from Charles River (Sulzfeld, Germany) and housed in groups under standardized conditions, 12 h/12 h day-night cycle, 50–60% humidity, 22–24 °C temperature and ad libitum tap water. Environmental enrichment was ensured by provision of housing and nesting material. Mice were randomly assigned into four experimental groups (vehicle vs. substance and subsequently into infected vs. mock-infected groups). According to a cooperative agreement between Charles River and The Jackson Laboratory, the JAX® C57BL/6J mouse strain bred by Charles River in Europe is genetically equivalent to that bred by The Jackson Laboratory in the U.S.

All animal experiments were conducted in accordance with the German Animal Welfare Law and were authorized by the local government (LAVES Oldenburg, Germany, permission number 33.9-42502-04-11/0516).

2.2. Treatment

Mice randomly assigned to the vehicle group received vehicle food chow, while mice randomly assigned to the treatment group had access to chow containing 1200 ppm PLX5622 per kilogram food chow (kindly provided by Plexxikon Inc., Berkeley, CA, USA). The regular food uptake of a 20–25 g mouse is 4 g per day, so the predicted overall intake of PLX5622 is 4.8 ppm per mouse per day. Treatment started 21 days prior to infection and continued until animals were sacrificed six to seven days post infection, resulting in a total treatment period of 28–29 days. This treatment protocol was established by Plexxikon and was previously used by other research groups (Dagher et al., 2015).

2.3. Infection

Intracerebral infection with the Daniels (DA) strain of TMEV was performed as previously described by Bröer et al. (2016, 2017) and Waltl et al. (2018). Therefore, a deep isoflurane inhalation anesthesia was induced and either 20 μl TMEV solution (2.44 × 10^7 PFU) or a mock solution were injected into the left parietal cortex of the mice using a free-hand method. The dose of TMEV was based on previous experiments of our group in B6 mice, to ensure a high incidence (> 50%) of early seizures in this mouse strain (Bröer et al., 2016).

2.4. Surveillance of body weight, seizures, and general behavior

Development of acute seizures, weight, general appearance and behavior in the first week after infection was assessed by animal monitoring twice daily for one hour, one h in the morning (between 9 and 12 a.m.) and one h in the afternoon (between 1 and 4 p.m.), with at least 3 h interval between the two recording sessions. Care was taken to avoid any group differences in recording periods by randomly assigning animals to the recording sessions. Recording was performed by experienced researchers, and the choice of recording periods was based on previous experiments with video-EEG monitoring of acute seizures in this model (for more details see Bröer et al., 2016, 2017).

2.5. Perfusion

On days 6 or 7 post infection animals were transcardially perfused with PBS followed by 4% paraformaldehyde. Brains and spleens were removed and left in formaldehyde overnight before being trimmed and embedded in paraffin. Additionally, segments of the spinal cord including the spinal column were removed, left in formaldehyde overnight, decalcified in EDTA for 48 h and afterwards embedded in paraffin. Animals used for flow cytometric analysis or RT-qPCR analysis were perfused with 4°C PBS only. Brains for flow cytometry were harvested and stored in 4°C PBS until being further processed, whereas brains for RT-qPCR were embedded in tissue freezing medium (Leica), frozen in liquid nitrogen and stored at −80°C until further processing. For RT-qPCR analysis of the spinal cord, the spinal column was harvested, the vertebrae removed and the segments of the cord equally processed as brains for RT-qPCR analysis. EDTA-blood for flow cytometry was sampled prior to the start of perfusion analysis and stored at 4°C.

2.6. Histology

Histological analyses of the brain were mainly focused on the dorsal hippocampus, as this brain structure is mainly damaged due to a hippocampal tropism of TMEV in B6 mice and is also associated with development of seizures and epilepsy (DePaula-Silva et al., 2017). Additionally, thalamic, hypothalamic and cortical regions were screened for morphological alterations and staining positivity. Furthermore, paraffin embedded sections of the cervical, thoracic and lumbar region of the spinal cord were examined to assess alterations in these regions.

Inflammatory processes were assessed by histological staining with H&E and additional immunohistochemistry staining with Mac-3 (Bio-Rad ABD Serotec GmbH, Puchheim, Germany), Iba1 (Wako Chemicals GmbH, Neuss, Germany and Abcam, Cambridge, GB), CD3 (DakoCytomation GmbH, Hamburg, Germany), Foxp3 (eBioscience; antibody-clone-FJK-16s-monoclonal/14-5773-80), TMEM119 (Abcam) and Iba1/Mac-3 fluorescent double staining and were analyzed as described recently (Bennett et al., 2016; Bröer et al., 2016; Ciurkiewicz et al., 2017; Waltl et al., 2018). H&E staining of spinal cord was used to examine neuronal necrosis, hypercellularity and perivascular infiltration and scored semi-quantitatively (Gerhauser et al., 2007). To assess the inflammatory signs in more detail, an additional Mac-3 staining was performed and scored semi-quantitatively (Bröer et al. 2016; Waltl et al., 2018) as follows: score: 0, absent (no positive, activated cells within the region); 1, mild (single positive cells found in the region); 2, moderate (up to 30% of region populated with positive cells); 3, severe (> 30% of region populated with positive cells). Data from both hemispheres were analyzed separately except for thalamus/hypothalamus and cerebral cortex, where Mac-3-positive cells were often in medial portions, so that these portions in both hemispheres were analyzed. CD3-positive, Foxp3-positive and TMEM119-positive cells within
the hippocampus and spinal cord were counted.

Neurodegeneration within the hippocampus was assessed using NeuN immunohistochemistry (Merek Millipore, Darmstadt, Germany) and scored semi-quantitatively (for details see Polaschek et al., 2010; Waltl et al., 2018). Also, Fluoro-Jade C (FJC) staining was performed for visualization of degenerating neurons (Groticke et al., 2008; Waltl et al., 2018). Glial fibrillary acid protein (GFAP) was immunostained to evaluate gliosis and astrocytosis in a semi-quantitative manner (Score: 0 = absent, 1 = mild, 2 = moderate, 3 = severe).

Virus antigen was stained using an anti-TMEV (VP1) antibody, and virus-positive cells within the brain were scored in a semi-quantitative manner (score 0, no stained cells; 1, few positively stained cells (< 10%); 2, 10–25% of cells stained; 3, > 25–45% of cells stained; 4, > 45% of cells stained), whereas virus-positive cells within the spinal cord were counted (Kummerfeld et al., 2009; Waltl et al., 2018).

To assess demyelination within the spinal cord, Luxol-Fast-Blue staining was performed and slides were screened for signs of demyelination (Kummerfeld et al., 2009). To assess alterations of the myeloid system in the periphery, spleens were stained for Iba1 and TMEV and analyzed using the positive pixel count of the ImageJ software (Waltl et al., 2018).

All analyses were performed by two experienced researchers blinded to the experimental groups. Data of both researchers was averaged.

2.7. Immune cell isolation and flow cytometry

Immune cells were isolated from whole brain homogenates as recently described (Waltl et al., 2018). Briefly, in accordance to the manufacturer’s specifications brain tissue was processed using the neural tissue dissociation kit (Miltenyi Biotec, Bergisch Gladbach, Germany). Afterwards, a percoll density gradient (30–70% percoll) was used for cell separation. Following density centrifugation, immune cells were washed and stained for 30 min at 4 °C using the following antibodies: CD11b APC-Cy7 (BD Biosciences; Heidelberg, Germany), and CD45.2 Pacific blue (Biolegend, San Diego, CA, USA). Respective isotype control antibodies or unstained samples were used to determine gates. A slightly adjusted protocol was used to isolate T-, and B-lymphocytes from whole brain homogenates. Therefore, the dissociation mix contained collagenase and DNase and the following antibodies were used: CD45.2 Pacific blue, CD8β PerCP-Cy5.5, Ly6C AF700, B220 PE-Cy5, CD4 FITC, CD44 brilliant violet 605, FoxP3 PE, Ly6G PE-Cy7,CCR2 AF647, MHCII PE (Biolegend, San Diego, CA, USA), and CD11c BV605 and CD11b APC-Cy7 (BD Biosciences; Heidelberg, Germany).

For flow cytometric analyses of blood samples, EDTA-blood was stained at 4 °C for 15 min with the following antibodies: B220 Pacific Blue, CD3 AF700 (Biolegend), CD11b APC-Cy7 (BD Biosciences), and CSF1R BV605 (Biolegend) (for more details see Waltl et al., 2018).

Flow cytometry was performed using a LSRII flow cytometer (BD Biosciences) and data was analyzed using FlowJo software (Tree Star, Ashland, OR, USA).

2.8. RT-qPCR

Ribonucleic acid (RNA) was isolated from frozen brain and spinal cord tissue using the RNeasy Lipid Tissue Mini Kit (Qiagen, Hilden, Germany) and subsequently transcribed into cDNA with the Omniscript RT Kit (Qiagen), Random Primers (Promega, Mannheim, Germany), and RNase Out (Invitrogen, Darmstadt, Germany). RT-qPCR was performed for TMEV, IL-1β, IL-6, IL-17, IL-34, Csfr1, TNfa, Ifnγ, Tgfb1, and three housekeeping genes (GAPDH, β-Actin, HPRT) using standard protocols, the Mx3005P qPCR System (Agilent Technologies Deutschland GmbH, Böblingen, Germany), and Brilliant III Ultra-Fast SYBR® QPCR Master Mixes (Gerhauer et al., 2005, 2012). Forward (F) and reverse (R) primers designed for TMEV and murine genes were as follows: TMEV-F 5′-GACCTAATCGAGGCAAAGC-3′; TMEV-R 5′-GTGAGGAGGCCGGAATGAGA-3′; IL-1β-F 5′-ACTCTCCACCGCAATGGAAG-3′; IL-1β-R 5′-AGTCGATTGTCAATGGGAAAC-3′; IL-6-F 5′-GTCTCGGAAATCATGTGGA-3′; IL-6-R 5′-CCAGAAGAATTCTTTAACATAGGC-3′; IL-17-F 5′-ACTCTCCACCGCAATGGAAG-3′; IL-17-R 5′-CTCTCA GGCTCCTCCTTCAG-3′; IL-34-F 5′-GGCCACCTTGGTCACTAAG-3′; IL-34-R 5′-TTTCCCAAGGCCAGCTGAG-3′; Csfr1-F 5′-AGGAAGTTGACAGTGTTGAG-3′; Csfr1-R 5′-CATGGTCTGTCACAGTAGTG-3′; TNFa-F 5′-GCCCTCTCTTCATTTCCCTT-3′; TNFa-R 5′-CACCCTGGTGGTGTGCTA CG-3′; Ifnγ-F 5′-CAGCAGCAGCTATGGAAAG-3′; Ifnγ-R 5′-AATCTG CGTCTGGAGATTT-3′; Tgfb1-F 5′-TTGCTTCAAGCTCCACAGAAG-3′; Tgfb1-R 5′-TGGTTGTGAGAGGCAAGAC-3′. Tenfold serial dilution standards ranging from 10^{8} to 10^{2} copies/μL were used to quantify the results. Experimental variations were corrected by using a normalization factor calculated from the housekeeping genes (Vandemopele et al., 2002). Melting curve analysis was performed to control the specificity of each reaction.

2.9. Statistics

GraphPad Prism Version 6 (La Jolla, CA, USA) was used for all statistical analyses. Depending on the distribution of the data, either parametric or nonparametric statistical tests were applied. For comparisons of two groups, either Student’s t-test or Mann-Whitney U test were used; for more than two groups either one-way or two-way ANOVA for parametric or nonparametric data were used, followed by a post-hoc Dunnett’s, Tukey’s or Bonferroni’s multiple comparisons test. The log-rank Mantel-Cox test was used for survival analysis. Seizure frequencies were compared using Fisher’s exact test. A P ≤ 0.05 was considered significant.

3. Results

Supplemental Table S1 gives an overview of the various experimental results of this study.

3.1. CSF1R inhibition depletes microglia in control and infected mice

Mice were treated with PLX5622, an orally bioavailable selective CSF1R inhibitor that crosses the blood–brain barrier (Dagher et al., 2015), for a period of 21 days prior to infection and treatment was continued until animals were killed 6–7 days after infection (Fig. 1). For this purpose PLX5622 was mixed into standard rodent chow at 1200 ppm.

To verify the microglia depletion, we used a microglia-selective antibody (TMEM119; Bennett et al., 2016). As shown in Fig. 2A for the hippocampus and para-hippocampal tissue, treatment with PLX5622 completely depleted microglia in mock-infected control mice and mice infected with TMEV, which was verified by counting TMEM119+ cells in hippocampus (Fig. 2B). A similar lack of TMEM119-positive microglia was observed throughout the brain of PLX5622-treated infected mice.

In infected B6 controls, significant microglia accumulation was seen in hippocampus and para-hippocampal regions (Fig. 2A and B).

We also quantitated CSF1R mRNA in the brain and spinal cord of mice with and without PLX5622, showing a significant increase in CSF1R upon infection in mice not treated with PLX5622 (Fig. 2C and D). In non-infected mice, expression of CSF1R was almost completely abolished by treatment with PLX5622, while a small but significant increase was seen in PLX5622-treated infected mice (Fig. 2D).

As described previously (Howe et al., 2012a; Cusick et al., 2013; Waltl et al., 2018), flow cytometry was used to differentiate CD45^{high} CD11b^{+} cells, which are considered to be mainly brain-infiltrating blood-derived monocytes, from CD45^{low} CD11b^{+} cells, which are considered to be mainly resident microglia, in the brain (Fig. 3A–D).

Therefore, we will name these two cell populations monocytes and microglia in the following. We found that treatment with PLX5622
Fig. 1. Experimental protocol of the study. For details see the Methods section. Overall five experiments were performed with this protocol in groups of mice. First experiments were performed over 7 days following infection. However, the poor health status of the infected PLX5622 treated mice at day 7, resulting in mortality or euthanasia (see Results), led us shorten subsequent experiments to 6 days.

Fig. 2. Depletion of microglia via CSF1R inhibition by treatment with PLX5622 in Theiler’s virus-infected mice. In the representative photomicrographs shown in A, microglia were immunostained with TMEM119. Infection increased TMEM119+ cells in controls, which was completely inhibited by treatment with PLX5622. Quantification of data is shown in B (microglia in hippocampus). Data are shown as boxplots with whiskers from minimum to maximal values; the horizontal line in the boxes represents the median value. In addition, individual data are shown. Sample size: 4–5 mock-infected controls; 3–5 mock-infected mice with PLX5622; 6–8 infected controls; 6–8 infected mice with PLX5622. Significant differences to mock-infected mice are indicated by asterisks (*P < 0.05; **P < 0.01) while significant differences between infected mice treated with PLX5622 and infected controls are indicated by the hash sign (#P < 0.05; ##P < 0.01; ###P < 0.001). (B) TMEM119+ cells in the hippocampus. Note the complete depletion of microglia in PLX5622-treated mice. (C) CSF1R mRNA quantitated by qPCR in the brains of mice. (D) CSF1R mRNA quantitated by RT-qPCR in the spinal cord of mice. All other details in C and D are as described above for B.
almost completely depleted microglia in mock-infected control mice and mice infected with TMEV (Fig. 3B, D, E).

3.2. CSF1R inhibition does not affect monocyte invasion after infection

As shown in Fig. 3D and F, treatment with PLX5622 did not prevent the massive brain invasion of blood-derived monocytes after infection with TMEV. In the blood, infected mice exhibited a significant increase in CSF1R⁺ monocytes, which was significantly reduced by PLX5622, but was still significantly higher than in mock-infected mice treated with PLX5622 (Fig. 3G). The number of Iba1⁺ macrophages in spleen was not affected by treatment with PLX5622 (Fig. 3H).

3.3. CSF1R inhibition leads to an unfavorable ratio between regulatory T cells and effector T cells after infection

Next we examined whether treatment with PLX5622 affects the adaptive immune response to infection with TMEV. Infiltration of CD3⁺ T-lymphocytes in the hippocampus was comparable in both groups (Fig. S1A). In addition, significant invasion of CD4⁺ T lymphocytes (CD4⁺ T effector cells) was observed in the brain of both infected groups, but CD4⁺ cell invasion was significantly lower in the PLX5622-treated group (Fig. 4A). Significant brain invasion of CD8⁺ cytotoxic T lymphocytes (CD8⁺ T effector cells), which are thought to play a significant role in viral clearance in the TMEV model in mice (Libbey and Fujinami, 2011; DePaula-Silva et al., 2017), was observed both in infected controls and infected mice treated with PLX5622, without significant inter-group difference (Fig. 4B). The brain CD4:CD8 ratio was 0.34 in infected controls, demonstrating that brain invasion of CD8⁺ T cells outnumbers invasion of CD4⁺ T cells in this model. As a result of reduced CD4⁺ T cell invasion in the PLX5622-treated group, the CD4:CD8 ratio was further reduced to 0.28. We also determined CD4⁺ and CD8⁺ cells in the spleen of infected mice (data not shown), resulting in a CD4:CD8 ratio of 1.3 in infected controls vs. 1.23 in PLX5622-treated animals.

Interestingly, in addition to the reduced invasion of CD4⁺ T cells in the brain of microglia-depleted infected mice (Fig. 4A), the number of invading CD4⁺ cells expressing the adhesion receptor CD44, which is upregulated after activation of naive T lymphocytes during their responses against viruses (Baaten et al., 2010), was significantly reduced in infected mice with depletion of microglia (Fig. 4C). The brain invasion of Foxp3⁺ regulatory T cells (Tregs) was increased in both infected controls and PLX5622-treated infected mice to a similar extent without significant inter-group difference (Fig. 4D).

However, in the hippocampus, which is the main target of TMEV infection in B6 mice because of tropism of the virus to this region (Libbey and Fujinami, 2011), a significant invasion of Foxp3⁺ Tregs was only observed in infected mice treated with PLX5622 (Fig. 4E). A
similar finding was obtained for the spinal cord (Fig. 4F). Since CD8+ T cell effector function is suppressed by elevated induction of Tregs (Richards et al., 2011), the significant increase in Tregs following CSF1R inhibition may lead to an unfavorable hippocampal and spinal cord ratio between Tregs and effector T cells, thus reducing antiviral immunity in these regions. This possibility is substantiated by the high increase in brain mRNA expression of the immunosuppressive cytokine IL-10 in the brain of infected PLX5622-treated mice (see below), which is released by Tregs and suppresses the activation of cytotoxic T cells (O'Shea et al., 2013).

Compared to mock-infected controls, a modest but statistically significant brain infiltration of neutrophils was observed in infected controls. In the PLX5622-treated group only a trend (P = 0.0826) for neutrophil infiltration was observed, but both infected groups did not differ from each other (Fig. S1B).

3.4. Microglia depletion by CSF1R inhibition increases distribution and persistence of Theiler's virus in the brain

B6 mice typically survive the acute viral encephalitis and clear TMEV within 2 weeks after infection (Libbey and Fujinami, 2011). Furthermore, in this mouse strain the virus exhibits tropism to the CA1 and CA2 regions of the hippocampus following intracerebral infection (Libbey and Fujinami, 2011). As shown in Fig. 5A, C and D, in most B6 control mice Theiler’s virus was already eliminated from the hippocampus by day 6/7, except for some staining in the CA2 in a few mice. In contrast, in mice treated with PLX5622, virus antigen was present throughout the hippocampus (Fig. 5B), resulting in significant increases in virus antigen-positive cells in the CA1/CA2 region of the hippocampus and other hippocampal regions (CA3, dentate gyrus) in which normally the virus is not present (Fig. 5C and D). At higher magnification, localization of the virus in neuronal cell bodies was visible in the hippocampus. In addition, virus antigen was infrequently observed outside neurons, associated with cells (most likely macrophages) around small vessels in the hippocampus. Furthermore, virus antigen was detectable in cerebral cortex, thalamus, hypothalamus (Fig. 5E), striatum and several other brain regions of PLX5622-treated mice, demonstrating that treatment with this compound increased the brain distribution of the virus. This is also illustrated by the presence of virus antigen in the spinal cord, in which the virus was not present in infected B6 vehicle controls (Fig. 5F). In addition to TMEV protein, we also determined TMEV mRNA, substantiating the significantly reduced elimination of the virus by treatment with PLX5622 (Fig. 5G and H).

3.5. CSF1R inhibition increases brain inflammation in the TMEV model

Immunohistochemistry with an antibody against Mac-3 was used to determine neuroinflammation in the hippocampus of infected mice. Mac-3 (also known as CD107b and LAMP-2) is a glycoprotein that is expressed at the plasma membrane of both activated microglia and
C. TMEV CA1/2 ipsilateral

D. TMEV CA3c ipsilateral

E. TMEV Thalamus & Hypothalamus

F. TMEV Spinal cord

G. TMEV mRNA Brain

H. TMEV mRNA Spinal cord

(caption on next page)
invading monocytes (Prinz et al., 2008). As shown in Fig. 6, at one week after infection, only some infected B6 control mice exhibited Mac-3+ cells in the hippocampus, whereas significant increase in Mac-3+ cells was observed in the cerebral cortex, thalamus and hypothalamus. Neuroinflammation was increased by treatment with PLX5622 in that significant increase in Mac-3+ cells was also observed in hippocampal CA1, CA2, CA3, and dentate gyrus (Fig. 6). Mac-3 labeling in the contralateral hemisphere was similar to that in the ipsilateral hemisphere shown in Fig. 6.

For further evaluating the activation status of myeloid cells in the brain, the amount of Iba1- and Mac-3-immunolabeled cells was compared in the hippocampus. Ionized calcium binding adaptor molecule 1 (Iba1) is a calcium-binding protein that labels both resting and activated microglia and infiltrating monocytes, whereas Mac-3 labels only activated microglia and invading monocytes (Greter et al., 2015). Regarding Iba1-labeling, almost no such labeling was observed in mock-infected controls (Fig. 7A), while the number of Iba1+ cells significantly increased in infected controls (Fig. 7B). In PLX5622-treated infected mice, the number of Iba1+ cells was lower in the CA1/CA2 region compared to infected controls (Fig. 7D) but not in other hippocampal regions (CA3c shown as example in Fig. 7E). However, a difference in Iba1+ cell shape was observed in that both round and ramified cells were observed in infected controls (Fig. 7B), representing monocytes and resting and activated microglia, whereas only round cells were observed in PLX5622-treated mice (Fig. 7C), substantiating the depletion of microglia as indicated by flow cytometry.

Depending on the hippocampal sub-region and treatment, on average 20–70% of the Iba1-labeled cells in infected mice were also positive for Mac-3. The number of Iba1/Mac-3 double labeled cells in the hippocampus did not significantly differ between infected controls and PLX5622-treated mice (Fig. 7F and G). However, when the activation percentage was calculated (i.e., the percentage of Iba1/Mac-3 double-labeled cells in percent of all Iba1-labeled cells), it was significantly higher in PLX5622-treated mice than infected controls in the CA3c (Fig. 7I) and perivascular area (data not shown).

For further characterizing the neuroinflammation following infection, we determined perivascular cuffing and hypercellularity in HE stained sections of the hippocampus. Perivascular cuffing, the accumulation of lymphocytes and monocytes in a dense mass around vessels, is a typical sign of viral encephalitis (Libbey and Fujinami, 2011). Similarly, hypercellularity, which results from disease-associated infiltration of lymphocytes and monocytes but also from astrocytosis and microgliosis, is an indicator of neuroinflammation. As shown in Suppl. Fig. S1C and D, infected B6 vehicle control mice only rarely exhibited perivascular infiltrates (PVI) at 1 week after infection, whereas a significant number of such infiltrates were observed in the hippocampus of PLX5622-treated mice. The same was true for hypercellularity (Supplemental Fig. S1E).

In addition to microglia, astrocytes are activated in response to TMEV infection in B6 mice, which can be labeled by glial fibrillary acidic protein (GFAP); GFAP+ astrocytes are particularly prominent in the hippocampus of infected mice (Libbey and Fujinami, 2011). In the present experiments, the number of GFAP+ astrocytes was only rarely different between mock-infected controls and TMEV-infected B6 control mice, whereas a marked astrogliosis was observed in PLX5622-treated mice (Supplemental Figs. S1F and S2). Within the hippocampus, GFAP+ astrocytes were primarily located in the tissue adjacent to the damaged cell layers (Supplemental Fig. S2). Gliosis was also observed in the corpus striatum of all infected mice (not illustrated).

### 3.6. CSFIR inhibition increases neurodegeneration in the TMEV model

When neurons were stained by NeuN, the most marked neuronal damage in infected B6 vehicle controls was observed in the CA2 region of the hippocampus (Fig. 8A, E–H). Treatment with PLX5622 significantly increased neuronal damage in the hippocampus (Fig. 8B, E–H). Significant neuron loss was also observed in parahippocampal cortex (not illustrated).

For further analysis of neurodegeneration in the hippocampus and other brain regions, sections were stained with Fluoro-Jade C (FJC), a sensitive and specific fluorescent marker of dying neurons (Schmued et al., 2005). In mock-infected controls, no FJC-stained neurons were observed (not illustrated), while intense staining of pyramidal cells was observed in the CA1 and CA2 layers of some infected B6 vehicle controls (Fig. 8C). In infected mice treated with PLX5622, FJC-labeled neurons were not only observed in CA1 and CA2 but all regions of the hippocampal formation, including CA3, the granule cell layer of the dentate gyrus and the dentate hilus (Fig. 8D). Furthermore, FJC-labeled neurons were observed in the cortex. Counting of FJC-positive cells in the hippocampus substantiated the impression of visual inspections of sections in that infected mice treated with PLX5622 exhibited significantly more degenerating neurons in the hippocampus than infected controls (see Fig. 8I and J as examples; significant differences also observed in CA3a, CA3c, and dentate gyrus).

### 3.7. CSFIR inhibition induces inflammation and neurodegeneration in the spinal cord of infected mice

In contrast to SJL/J mice, B6 mice typically do not develop a chronic inflammatory demyelinating disease in the spinal cord, so B6 mice are considered to be resistant to the TMEV-induced demyelinating disease (Libbey and Fujinami, 2011). Consistent with this view, infected B6 control mice did not show any indication of cell death or inflammation in the spinal cord (Fig. 9C) except for a moderate increase in CD3+ T lymphocytes and Mac-3 labeling (Fig. 9K and L). Furthermore, myelin staining did not reveal any demyelination (Fig. 9E). In contrast, PLX5622-treated mice exhibited TMEV antigen (Fig. 9A), necrosis of motor neurons (Fig. 9B and G) and inflammation in the spinal cord (Fig. 9D, H–J), thus explaining the paralysis observed in these mice (see below). Also, a significant increase in FoxP3+ regulatory T cells was only observed in PLX-treated mice (Fig. 4D). Demyelination was not observed in PLX5622-treated mice (Fig. 9F), which may be due to the fact that mice were killed 6–7 days after infection, because demyelination is known to develop slowly in sensitive mouse strains such as SJL/J (DePaula-Silva et al., 2017).
Fig. 6. Neuroinflammation as indicated by Mac-3 labeling of activated microglia and infiltrating monocytes in the brain of Theiler’s virus-infected mice. (A) A representative photomicrograph of an infected control mouse. (B) A representative photomicrograph of an infected mouse with PLX5622 treatment. Scale bar in B = 200 µm. (C–H) Semiquantitative data for expression of Mac-3. Data are shown as boxplots with whiskers from minimum to maximal values; the horizontal line in the boxes represents the median value. In addition, individual data are shown. Sample size: 4 mock-infected controls; 5 mock-infected mice with PLX5622; 8 infected controls; 8 infected mice with PLX5622. Significant differences to mock-infected mice are indicated by asterisks (*P < 0.05; **P < 0.01; ***P < 0.001) while significant differences between infected mice treated with PLX5622 and infected controls are indicated by the hash sign (#P < 0.05). (C) Scores for Mac-3 labeling in the ipsilateral CA1 of the hippocampus. At the time chosen (6–7 days after infection), significant difference to mock controls is only observed for infected mice treated with PLX5622. (D) Scores for Mac-3 labeling in the ipsilateral CA2 of the hippocampus. Significant difference to mock controls is only observed for infected mice treated with PLX5622. (E) Scores for Mac-3 labeling in the ipsilateral CA3c of the hippocampus. Significant difference to mock controls is only observed for infected mice treated with PLX5622. (F) Scores for Mac-3 labeling in the ipsilateral dentate gyrus (DG) of the hippocampus. Significant difference to mock controls is only observed for infected mice treated with PLX5622. (G) Scores for Mac-3 labeling in the cerebral cortex. Significant difference to mock controls is observed for both infected groups. (H) Scores for Mac-3 labeling in the thalamus and hypothalamus. Significant difference to mock controls is observed for both infected groups.
The image presents a study on the effects of infection and treatment with PLX on the expression of markers Iba1 and Mac-3. The study involves both mock control and infected conditions.

**A**
- **Mock, control**
- **Infected, control**
- **Infected, PLX**

**B**
- **DAPI**
- **Iba1**
- **Mac-3**
- **DAPI/Iba1/Mac-3**

**D**
- **Iba1 CA1/CA2**
- **No. of Iba1 cells**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

**E**
- **Iba1 CA3c**
- **No. of Iba1 cells**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

**F**
- **Iba1/Mac-3 CA1/CA2**
- **No. of Iba1/Mac-3 cells**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

**G**
- **Iba1/Mac-3 CA3c**
- **No. of Iba1/Mac-3 cells**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

**H**
- **Activation Percentage CA1/CA2**
- **% of Mac-3 Iba1**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

**I**
- **Activation Percentage CA3c**
- **% of Mac-3 Iba1**
- **Mock control**, **Infected control**, **Mock PLX**, **Infected PLX**

*(caption on next page)*
Fig. 7. Neuroinflammation as indicated by Iba1 and Mac-3 labeling and double labeling. While Iba1 labels both resting and activated microglia and infiltrated monocytes in the brain, Mac-3 only labels activated microglia and infiltrated monocytes. (A) Representative photomicrographs of a mock-infected control mouse. The first graph shows labeling of cell nuclei by DAPI in blue, the second graph Iba1 labeled cells in green, the third graph Mac-3 labeled cells in red, and the fourth graph cells labeled for both Iba1 and Mac-3. Only few Iba1 and no Mac-3 labeled cells are visible, consistent with the lack of neuroinflammation in mock-infected mice. (B) Representative photomicrographs of an infected control mouse. Note the marked increase in both Iba1 and Mac-3 labeling. Both ramified and round cell types are visible. Iba1 and Mac-3 double labeled cells are indicated in yellow. (C) Representative photomicrographs of an infected mouse with PLX5622 treatment. Note the lack of ramified cells, which is due to microglia depletion. Scale bar in A, B, and C = 100 μm. (D-I) Quantitative data for expression of Iba1, Iba1/Mac-3 double labeled cells, and the “activation percentage” indicated by Iba1/Mac-3 double labeling. Data are shown as boxplots with whiskers from minimum to maximal values; the horizontal line in the boxes represents the median value. In addition, individual data are shown. Sample size: 4 mock-infected controls; 4-5 mock-infected mice with PLX5622; 7-8 infected mice with PLX5622. Significant differences to mock-infected mice are indicated by asterisks (\(^*\) \(P < 0.05\); \(^{**}\) \(P < 0.01\); \(^{***}\) \(P < 0.001\)). (D) Number of Iba1+ cells in the CA1/C2 layers of the ipsilateral hippocampus. Note the significant reduction in such cells in infected PLX5622-treated mice. (E) Number of Iba1+ cells in the CA3c layer of the ipsilateral hippocampus. (F) Number of Iba1/Mac-3 double-labeled cells in the CA1/C2 layers of the ipsilateral hippocampus. (G) Number of Iba1/Mac-3 double-labeled cells in the CA3c layer of the ipsilateral hippocampus. (H) Activation percentage in the CA1/C2 layers of the ipsilateral hippocampus. (I) Activation percentage in the CA3c layer of the ipsilateral hippocampus. Note the higher percentage of Iba1/Mac-3 double labeled cells in infected PLX5622-treated mice. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

3.8. CSF1R inhibition increases morbidity and mortality in the TMEV model

In infected B6 vehicle controls, animals did not show any signs of severe impairment due to infection and the following encephalitis as demonstrated by the Total Clinical Score (TCS) and weight development during the first days after infection (Fig. 10A). However, this strikingly changed in the PLX5622 treated group starting from day five post-infection (Fig. 10A). Animals showed severe paralysis of the hind limbs (Fig. 10B), bent-back posture, and a strong decrease in activity and food uptake. This health impairment was also displayed by severe loss of weight (Fig. 10C). Hence, some animals, which strongly showed these symptoms, had to be sacrificed before day seven to avoid suffering. Furthermore, one animal of the PLX5622-treated group died following infection. Survival analysis resulted in a highly significant difference between infected B6 vehicle controls and PLX5622-treated mice (Fig. 10D).

3.9. CSF1R inhibition facilitates seizure induction in the TMEV model

When determining the daily incidence of seizures across the 7 days after infection, maximum seizure occurrence was reached significantly more rapidly in PLX5622-treated mice than in vehicle controls, indicating that microglia depletion facilitated seizure induction (Fig. 11A). However, overall seizure incidence was not affected by microglia depletion: about 68% of the infected B6 vehicle controls exhibited seizures in the first week after infection compared to 78% of the PLX5622-treated group, which was not significantly different (Fig. 11B). No seizures were observed in the mock-infected groups. Most infected mice exhibited several seizures, but no significant inter-group difference in seizure frequency was observed (Fig. 11C). Similarly, the maximum seizure severity was not different in the two infected groups (Fig. 11D).

3.10. CSF1R inhibition alters cytokine and interferon mRNA expression in brain and spinal cord

In a series of previous studies in TMEV-infected B6 mice, release of the proinflammatory cytokines TNF-α and IL-6 by invading monocytes was described as particularly important for the seizure phenotype of these mice (DePaual-Silva et al., 2017). When these two cytokines were determined by RT-qPCR in the present study, significant increases in IL-6 and TNF-α mRNAs were observed in the brain (Fig. 12A, Supplemental Fig. S3A) and spinal cord (Fig. 12D, Supplemental Fig. S3E) of infected control mice. In infected PLX5622-treated mice, the brain increase in IL-6 was 20-fold more marked than the increase determined in infected controls (Fig. 12A), indicating increased release of this cytokine by invading monocytes. In addition to IL-6, a 6.6-fold more marked increase in expression of IL-10 was determined in the brain of infected PLX5622-treated mice compared to infected controls (Fig. 12B). Furthermore, a 2.3-fold more marked increase in expression of IFNγ was determined in the brain of infected PLX5622-treated mice compared to infected controls (Fig. 12C). In the spinal cord, the increase in IFNγ in infected PLX5622-treated mice was 49-fold higher than the increase determined in infected controls (Fig. 12F).

No marked differences between groups with or without PLX5622 were observed for IL-1β, IL-34, and TGF-β1 (Supplemental Fig. S3). IL-17 was below detection level in both mock and TMEV infected mice, including PLX5622 treated animals. Supplemental Fig. S4 summarizes the selective increases in cytokine and interferon mRNA levels in infected PLX5622-treated mice vs. infected controls, demonstrating the preferential increase of IL-6, IL-10 and IFNγ in the brain and IFNγ in the spinal cord of PLX5622-treated mice. Since IFNγ is preferentially released by cytotoxic CD8+ T cells whereas IL-10 is released by Tregs in the infected brain (Oleszak et al., 2004; DePaula-Silva et al., 2017), we calculated the ratio between the increase in IFNγ and IL-10 in the brain of infected controls vs. infected PLX5622-treated mice. This ratio was 5.57 in infected controls, indicating higher functionality of T effector cells vs. Tregs as regularly observed in TMEV-infected B6 mice (DePaula-Silva et al., 2017), whereas the ratio was 0.84 after infected PLX5622-treated mice, indicating suppression of T effector cells by activated Tregs.

4. Discussion

Intracerebral infection of B6 mice with Theiler’s virus is the first, and currently only, rodent model of viral encephalitis-induced seizures and epilepsy (Vezzani et al., 2016; DePaula-Silva et al., 2017). This model is ideally suited to investigate the mechanisms underlying the generation of seizures and hippocampal pathology developing during and after viral encephalitis. In contrast to B6 mice, which rapidly clear the virus, the virus persists in the CNS of SJL/J mice, which do not develop seizures and hippocampal damage, but a slowly progressing inflammatory demyelination in the spinal cord, making the TMEV infection in SJL/J mice a widely used model of multiple sclerosis (Oleszak et al., 2004; Procaccini et al., 2015; DePaula-Silva et al., 2017). While TMEV persistence in SJL/J mice is important to induce chronic demyelination, the trigger mechanism for demyelination is inflammation and the induction of autoimmunity (DePaula-Silva et al., 2017). The persistence of TMEV in the CNS of SJL/J mice is thought to be related, at least in part, to the high induction of Tregs during the acute stage of the infection, because Tregs can secrete immunosuppressive cytokines (such as IL-10), thereby blocking activation and effector function of cytotoxic CD8+ T cells that are important for viral clearance (Richards et al., 2011). In contrast to SJL/J mice, the lower number of Tregs induced in B6 mice after TMEV infection has been suggested to allow for rapid viral clearance by CD8+ T effector cells (Richards et al., 2011).
Indeed, Tregs, characterized by expression of the transcription factor forkhead box P3 (Foxp3), are known to modulate the immune response to viral infection, and numerous infection models have demonstrated that Treg depletion increases viral clearance and IFNγ production by CD8+ T cells (Groux et al., 1997; Dittmer et al., 2004; He et al., 2004; Eggena et al., 2005). However, in TMEV-infected B6 mice, ablation of Foxp3+ Tregs during the acute disease phase of TMEV infection did not accelerate virus clearance (Pradeeth et al., 2014). In contrast, ablation of cytotoxic CD8+ T cells rendered B6 mice susceptible to develop enhanced chronic demyelinating neuroinfection comparable to the encephalomyelitis observed in SJL/J mice, which might partly be mediated by boosting the suppressive capacity of Tregs on viral elimination (Ciurkiewicz et al., 2017).

In addition to CD8+ T cells, it has been suggested that CD4+ T cells may confer protection against TMEV persistence and demyelination in resistant mouse strains such as B6 (Murray et al., 1998; Kang et al., 2005; Mohindru et al., 2006). Indeed, genetic deletion of either CD4+ or CD8+ cells from B6 mice resulted in viral persistence and demyelination during the chronic stage of disease (Murray et al., 1998). Interestingly, neither CD4+ nor CD8+ cells are required for development of acute seizures following infection with TMEV in B6 mice (DePaula-Silva et al., 2018).

The present study shows that depletion of microglia is an effective means to retard viral clearance and exacerbate the TMEV-induced disease in B6 mice, thus defining the role of microglia in this model of viral encephalitis. Previously, understanding the roles that microglia play in a neurotropic viral infection and how these roles affect the immune response to viruses has been complicated by differentiating microglia from extensively infiltrating monocytes with potentially overlapping functions during neuroinflammation (Terry et al., 2012; Prinz and Priller, 2014; Prinz et al., 2017). Furthermore, while microglia, the resident macrophages of the CNS, are often touted as the first responders to a CNS viral infection and are known to respond rapidly to injury (Rock et al., 2004; Carrithers, 2014), there is little direct evidence to show that microglia are needed for an optimal host response to pathogens. Selective and complete depletion of microglia by chronic treatment with the CSF1R inhibitor PLX5622 is an ideal means to study the functions that microglia use in responding to viruses. PLX5622 is a potent inhibitor of CSF1R tyrosine kinase activity (K uninsured = 5.9 nM) with at least 50-fold selectivity over 4 related kinases, and over 100-fold selectivity against a panel of 230 kinases (Dagher et al., 2015). As shown here, by the PLX5622 treatment protocol used, CSF1R mRNA expression in the CNS was almost completely lost, which is a consequence of the depletion of microglia that express this mRNA.

Recent studies in the TMEV model in B6 mice have suggested that both seizures and hippocampal damage in this mouse strain are primarily due to infiltration of blood-derived monocytes and cytokines released by these cells (Howe et al., 2012a,b; Cusick et al., 2013), resulting in apoptotic death of neurons in the CA1/CA2 area of the hippocampus (Buenz et al., 2009; Howe et al., 2012b). Indeed, suppression of monocyte infiltration by deleting the chemokine receptor CCR2 prevented hippocampal damage following TMEV infection, but did not prevent encephalitis-associated seizures (Käufner et al., 2018), thus challenging previous assumptions with less selective approaches (Cusick et al., 2013). The present study shows that depletion of microglia by PLX5622 exacerbates hippocampal damage and increases progression of seizures following viral encephalitis, indicating that microglia may exhibit a protective role in this model, at least in the early phase following infection. Furthermore, neuronal damage, associated with severe motor impairment, occurred in the spinal cord, which is typically not affected in B6 mice. The severe clinical impairment and enhanced mortality in infected mice treated with PLX5622 did not allow evaluating whether spinal cord demyelination occurs at later stages of the infection as observed in TMEV-infected SJL/J mice.

The first study that reported the effect of microglia depletion in B6 mice on viral encephalitis used intracranial infection with a neurotropic variant of the JHMV strain of MHV (Wheeler et al., 2018), which induces mild acute encephalitis and acute and chronic demyelinating disease (Bergmann et al., 2006). Treatment with PLX5622 increased CNS replication of MHV and subsequent mortality in B6 mice, which was explained by ineffective innate and virus-specific T cell responses resulting from microglia depletion (Wheeler et al., 2018). Indeed, microglia depletion led to a markedly reduced brain infiltration of CD4+ T cells and Foxp3+ Tregs as well as IFNγ expression at 7 days after infection (Wheeler et al., 2018). The present data show that these consequences of microglia depletion are virus specific, because we did not observe a decrease but a significant increase in infiltration of Foxp3+ Tregs in hippocampus and spinal cord and only a relatively small reduction in brain invasion of CD4+ T cells in TMEV-infected, PLX5622-treated mice compared to infected controls, although the invasion of CD4+ CD44+ T cells was markedly reduced by microglia depletion. Furthermore, in apparent contrast to the study of Wheeler et al. (2018), IFNγ expression was significantly increased in brain and spinal cord of PLX5622-treated mice in response to TMEV infection.

In a recent study using PLX5622 for determining the role of microglia in flaviviral pathogenesis in female Swiss-Webster mice (Seitz et al., 2018), depletion of microglia resulted in increased mortality and viral titer in the brain following infection with either West Nile virus (WNV) or Japanese encephalitis virus (JEV). Interestingly, the expression of several pro-inflammatory genes was increased in virus-infected, microglial-depleted mice compared to virus-infected, untreated controls. T cell responses were not examined in the study of Seitz et al. (2018).

In both previous studies on depleting microglia in viral CNS infection models, Iba1 was used as a marker for microglia (Seitz et al., 2018; Wheeler et al., 2018). However, Iba1 is not selective for microglia but also labels monocytes that invade the brain during viral encephalitis (Greter et al., 2015), which is also shown here. For the present study,
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G Spinal cord, neuronal necrosis

H Spinal cord, PVI grey matter

I Spinal cord, PVI white matter

J Spinal cord, hypercellularity grey matter

K Spinal cord, CD3+ T lymphocytes (average of all parts)

L Spinal cord, Mac-3 grey matter

(caption on next page)
Fig. 9. Neuroinflammation and neurodegeneration in the spinal cord. (A) A representative photomicrograph of virus antigen staining in the spinal cord of an infected PLX5622-treated mouse. No such staining was observed in infected controls (see Fig. 5F for quantitative data). (B) Neuronal necrosis in the spinal cord (arrows) as indicated in H&E-stained sections of an infected PLX5622-treated mouse. (C) Mac-3 labeled cells in the spinal cord of an infected control. (D) Mac-3 labeled cells in the spinal cord of an infected PLX5622-treated mouse; note the increase in Mac-3 labeled cells vs. (C). (E) Luxol fast blue (LFB) staining of myelin in the spinal cord of an infected control. (F) LFB staining in the spinal cord of an infected PLX5622-treated mouse. No indication of demyelination was observed in E or F. (G-L) Quantitative data for neuronal necrosis (G), perivascular infiltrates (PVI; H, I), hypercellularity (J), CD3+ T lymphocyte infiltration (K), and Mac-3 labeled cells (L). Data are shown as boxplots with whiskers from minimum to maximal values; the horizontal line in the boxes represents the median value. In addition, individual data are shown. Sample size: 4 mock-infected controls; 5 mock-infected mice with PLX5622; 8 infected controls; 8 infected mice with PLX5622. Significant differences to mock-infected mice are indicated by asterisks (*P < 0.05; **P < 0.01) while significant differences between infected mice treated with PLX5622 and infected controls are indicated by the hash sign (#P < 0.05; ##P < 0.01). All parameters shown in G-L demonstrated the increased neuroinflammation and neurodegeneration induced by treatment with PLX5622 in the spinal cord of C57BL/6 (B6) mice. All scale bars = 200 µm. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Fig. 10. Clinical data from infected mice. Experiments were performed several times (see Fig. 1) and data summarized; sample size: 19 mock-infected controls; 20 mock-infected mice with PLX5622; 28 infected controls; 27 infected mice with PLX5622. (A) Illustrates the “total clinical score”, describing the overall clinical impairment, ranging from 0 (healthy) to 10 (maximal clinical impairment) (see Methods). Infected controls did not differ from mock-infected controls (except for seizures, which were assessed separately as shown in Fig. 11), while infected PLX5622-treated mice developed severe clinical symptoms, including paralysis of the hind limbs, bent-back posture, and a strong decrease in general activity and food uptake. Data are shown as means ± SEM and were analyzed by two-way ANOVA; significant differences between infected PLX5622-treated mice to all other groups are indicated by the hash sign (#P < 0.0001). (B) A representative picture of a PLX5622-treated infected mouse with paralysis of hind limbs, which was never observed in infected controls. (C) Body weight development. Data are shown as means ± SEM and were analyzed by two-way ANOVA; significance between infected and mock-infected mice is indicated by asterisk (*P < 0.01), while significance between PLX5622-treated and vehicle-treated infected mice is indicated by the hash sign (#P < 0.01). Note the much more severe weight loss in infected PLX5622-treated mice. (D) Survival curves for infected controls and infected PLX5622-treated mice. Mice that had to be killed before day 7 (the planned end of the experiment) because of poor health status were included in survival analysis. Statistical evaluation of data by the log-rank (Mantel-Cox) test indicated a highly significant (P < 0.0001) difference between groups.
we chose the much more selective microglia marker TMEM119 (Bennett et al., 2016), demonstrating that PLX5622 results in complete microglia depletion at the chosen dosing protocol. This was confirmed by flow cytometry.

CSF1R, the receptor affected by PLX5622, is also expressed on other myeloid cells, including brain perivascular macrophages and infiltrating hematopoietic monocytes (Hamilton and Achuthan, 2013). As a consequence, we observed a significant decrease in circulating CSF1R⁺ monocytes in the blood of mock infected mice after treatment with PLX5622. After infection, blood CSF1R⁺ monocytes increased in both controls and PLX5622-treated groups, although to a slightly lower degree in the PLX5622-treated mice. This was not associated with any obvious reduction in brain infiltration of blood-derived monocytes as both demonstrated by flow cytometry and immunolabeling by Mac-3 and Iba1. Instead, both Mac-3 and Iba1 indicated an increase of myeloid cells in the hippocampus and spinal cord of infected mice after depletion of microglia, indicating that infiltrating hematopoietic monocytes/macrophages try to compensate for the lack of microglia. In line with this idea, mRNA expression of IL-6, which is released by both invading monocytes and activated microglia (Oleszak et al., 2004), was increased 20-fold in microglia-depleted brains of infected mice.

Interleukin-6 (IL-6) plays an important role in the development and progression of inflammatory responses, in part by facilitating the generation of IL-17-producing Th17 cells (Kothur et al., 2016). Many viral infections, including TMEV, result in the vigorous production of IL-6 (Kothur et al., 2016; DePaula-Silva et al., 2017). B6 mice carrying an IL-6 transgene, resulting in the production of excessive IL-6, develop a TMEV-induced demyelinating disease accompanied by an increase in viral persistence and an elevated Th17 cell response in the CNS (Hou et al., 2014). IL-17 producing Th17 cells have been shown to promote...
persistent viral infection by inhibiting cytotoxic T cell function and apoptotic cell death (Hou et al., 2009). As shown by Hou et al. (2014), excessive IL-6 promotes the generation of Th17 cells, and the resulting IL-6 and IL-17 synergistically promote viral persistence by protecting virus-infected cells from apoptosis and CD8+ T cell-mediated target destruction. These data suggest that the 20-fold increase in IL-6 following microglia depletion by PLX5622 may be involved in the altered responses to TMEV infection observed in the present experiments. However, we did not observe an increase in IL-17 in microglia-depleted mice following TMEV infection, demonstrating low numbers of Th17 cells in this early phase of the disease (6–7 days p.i.).

Instead, we observed a marked increase in brain mRNA expression of the immunosuppressive cytokine IL-10 in the brain of infected PLX5622-treated mice compared to infected B6 controls. Among other cell types, IL-10 is secreted by Tregs, which were significantly increased in the brain of infected PLX5622-treated mice. Thus, similar to TMEV infection in SJL/J mice, microglia depletion in B6 mice seems to lead to an unfavorable ratio between Tregs and T effector cells, resulting in suppression of T cell activation and effector function. This interpretation of the data was substantiated when calculating the ratio between IFN-γ and IL-10 in the brain of infected controls vs. infected mice treated with PLX5622, demonstrating a reversal in this ratio from 5.57 to 0.84. This indicates that IL-10 suppressed functionality of IFN-γ-releasing T effector cells following microglia depletion.

In the spinal cord, however, a massive, 50-fold increase of IFN-γ was observed. This increase in IFN-γ may be critically involved in disease exacerbation. In neuroinflammatory CNS disorders, IFN-γ is elevated mainly in viral encephalitis, suggesting activation of lymphocytes in the CNS which participate in viral clearance (Kothur et al., 2016). IFN-γ, which is produced by activated natural killer (NK) and T cells, is important for virus clearance and induces many immunomodulatory effects, including activation of macrophages, promotion of leukocyte adhesion to allow trafficking of cells to the CNS, direct antiviral and antiproliferative effects, and induction of the release of other cytokines, including TNF-α and IL-1 (Bowen and Olson, 2013). IFN-γ has been implicated in the clearance of virus from CNS following infection with a wide variety of viruses, including TMEV (Rodriguez et al., 2003). However, at high concentrations as observed following microglia depletion here, IFN-γ is neurotoxic through a neuron specific, calcium-permeable complex of IFN-γ-receptor and the AMPA subtype of glutamate receptors (Mizuno et al., 2008). Thus, IFN-γ in concert with other cytokines such as IL-6β may contribute to the increased degeneration of hippocampal and spinal cord motor neurons and clinical signs observed after microglia depletion in TMEV-infected mice.

This, however, does not necessarily explain the facilitation of seizures observed in infected PLX5622-treated mice. Here, a direct effect of microglia could be important. By debris clearance and continuously contacting dendritic spines to regulate structural synaptic changes, microglia plays a fundamental role in facilitating the reorganization of neuronal circuits and triggering repair in CNS injury (Neumann et al., 2009; Kim et al., 2013). Proper synaptic pruning (or engulfment) by microglia has been shown to modulate seizure threshold (Kim et al., 2013), which could explain the facilitated seizure induction observed after microglia depletion in Thiel’s virus-infected mice.

Overall, based on the present data, the following scenario can be postulated. Depletion of microglia reduces the local immune response to infection, resulting in increased virus proliferation. Enhanced induction of Tregs in hippocampus and spinal cord and resulting IL-10 release seem to add to the reduced immune response to viral infection observed after microglia depletion. On the other hand, markedly

![Fig. 12.](image-url)
increased IL-6, and, in the spinal cord, IFNγ levels contribute to neurodegeneration and elevated seizure susceptibility. Also, the increased astrogliosis observed in infected, microglia-depleted mice may contribute to neuroinflammation and related processes, although a neurotoxic subtype (A1) of astrocytes, which is induced by neuroinflammatory microglia (Liddelow et al., 2017), should be reduced or absent in microglia-depleted mice. Our results demonstrate a critical role for microglia in the context of viral encephalitis, affecting virus replication and innate and adaptive immune responses as well as disease progression and severity. However, as indicated by the significant differences in response to microglia depletion between the study of Wheeler et al. (2018) with a coronavirus (MHV) and the present study with a picornavirus (TMEV), both performed in B6 mice, the role of microglia in viral encephalitis markedly depends on the specific virus inducing the disease. Furthermore, the severity of viral encephalitis certainly has an impact on specific microglia response. The present findings enhance our understanding of microglia-T cell crosstalk that seems to play a vital role in neuroinflammation, whether through soluble mediators or via contact-dependent interactions (Schettes et al., 2017).

Author contributions

IW and WL conceived the study; IW, CK, IG, CC and LG acquired and validated the data and executed the experiments; IW, CK, IG, CC, LG, and WL analyzed the data; WL wrote the manuscript and all authors revised the manuscript.

Conflict of interest

The authors declare no competing financial interests.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.jibb.2018.09.006.

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