

Expanded View Figures

Figure EV1. Immune cell analysis in EAE and TNP-ATP-treated EAE mice.

- A Correlation between the neurological score and the number of Iba1⁺ cells in gray matter and white matter of the spinal cord.
- B Flow cytometric quantification of immune cell infiltrates (gating as in Fig 2E) in the brain of TNP-ATP-treated or non-treated mice analyzed at the peak of symptoms after EAE induction (n = 8). Data are presented as mean \pm s.e.m. and were analyzed by Student's *t*-test. *P < 0.05.



Figure EV2. Heatmap of pro-inflammatory and anti-inflammatory genes at EAE peak.

Expression of most pro-inflammatory genes and some anti-inflammatory genes was increased in the spinal cord of EAE mice (n = 3) at the peak phase, but TNP-ATP did not induce significant changes (n = 3). Tables indicate statistical significance between control and EAE (*) and between EAE and EAE + TNP-ATP ([#]). Statistics were performed with Student's *t*-test.



Figure EV3. Pro-inflammatory and anti-inflammatory polarization in wild-type and P2X4 $^{-\prime-}$ microglia.

Staining for iNOS (red) and mannose receptor (MRC1, green) in control, proinflammatory, and anti-inflammatory microglia from wild-type and P2X4^{-/-} mice. Scale bar = 50 μ m (n = 4 experiments performed in triplicate). Data are presented as mean \pm s.e.m. and were analyzed by Student's *t*-test. *^{*H*}P < 0.05, **P < 0.01. Symbols indicate significance versus control (*) or versus pro-/antiinflammatory microglia ([#]).



Figure EV4. Functional role of P2X4R in oligodendrocytes in vitro.

- A Effect of ATP γ S (10 μ M) \pm ivermectin or TNP-ATP (10 μ M) on oligodendrocyte differentiation quantified on the basis of the number of MBP⁺ cells/total cell number and the area of the cells. Area was calculated with ImageJ software (n = 3).
- B Oligodendrocyte cell viability after 24-h incubation with BzATP, a broad-spectrum agonist of purinergic receptors, in the absence or presence of TNP-ATP (10 μ M) (left) or ATP γ S (right) in the absence or presence of IVM (3 μ M) (n = 3).
- C Western blot analysis of BDNF in control and IVM-treated wild-type and P2X4^{-/-} microglia (n = 3).
- D Bdnf mRNA levels in the spinal cord of WT and $P2X4^{-/-}$ mice in control and after EAE induction (n = 4 mice/group).

Data information: Data are presented as mean \pm s.e.m. and were analyzed by one-way ANOVA (B) and Student's *t*-test (C, D). **P* < 0.05, ****P* < 0.001. Source data are available online for this figure.



Figure EV5. Role of P2X4R in myelin phagocytosis and lysosome function.

- A Spinal cord sections of wild-type (n = 6) and P2X4^{-/-} mice (n = 7) 35 days after EAE induction stained for MBP. Scale bar = 50 and 10 (inset) μ m. Inset images come from different fields at higher magnification. Right, demyelination score analyzed in MBP-stained sections.
- B Effect of IVM (3 μ M) on myelin endocytosis (1 h) and degradation (3 days) in microglia from wild-type and P2X4^{-/-} mice (n = 3).
- C Treatment with IVM (3 μ M; 16 h) increased late endosome–lysosome (LEL) size in microglia from wild-type and P2X4^{-/-} mice. Representative images and histogram for LEL size distributions under the conditions indicated (n = 20 cells from two independent experiments). Scale bar = 10 μ m.
- D IVM-treated microglia (3 μ M) and anti-inflammatory microglia showed an acidic shift in lysosomal pH, as analyzed on the basis of fluorescein/rhodamine–dextran ratio (n = 20 cells from two independent experiments). Scale bar = 10 μ m.

Data information: Data are presented as mean \pm s.e.m. and were analyzed by one-way ANOVA. "P < 0.05, **P < 0.01, ***P < 0.001. Symbols indicate significance versus control (*) or versus pro-inflammatory microglia ([#]).