

# **Supplemental figures**



#### Figure S1. Targeted deletion of SYK in tamoxifen-treated Syk<sup>fl/fl</sup> Cx3cr1<sup>Ert2Cre</sup> mice, related to Figure 1

(A-B) Cre-negative 5xFAD *Syk*<sup>fl/fl</sup> littermate controls (5xFAD mice) and Cre-negative *Syk*<sup>fl/fl</sup> littermate controls (*Syk*<sup>con</sup> mice) received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Spleens and brains were harvested at 5 months of age to evaluate SYK expression in T cells, monocytes, and microglia. T cells, monocytes, and microglia were sorted from single-cell suspensions using respective anti-CD90.2<sup>+</sup>cells), anti-CD11b<sup>+</sup>, and anti-CD11b<sup>+</sup>(microglia) -coated magnetic beads and magnetic column sorting. (A) MACS-sorted splenic T cells and monocytes and MACS-sorted brain microglia were harvested from 5-month-old *Syk*<sup>con</sup> and 5xFAD mice, and levels of SYK protein (top panel) and Actin or Ponceau-stained protein (bottom panel) were measured by Western blotting. (B) Quantification of intensity of SYK protein bands normalized to total Actin in sorted splenic T cells and monocytes, and intensity of SYK protein bands normalized to Ponceau staining in brain microglia from *Syk*<sup>con</sup> and 5xFAD mice. (C-F)





Syk<sup>fl/fl</sup> *Cx3cr1*<sup>ERT2Cre</sup> (Syk<sup>ΔMG</sup> mice) and Cre-negative Syk<sup>fl/fl</sup> littermate controls (Syk<sup>Con</sup> mice) received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Brains and spinal cords were later harvested to evaluate SYK deletion. Microglia were sorted from single-cell suspensions using anti-CD11b<sup>+</sup>-coated magnetic beads and magnetic column sorting. (C) Representative flow cytometry gating strategy used to validate purity of MACS-sorted microglia from naive *Syk*<sup>Con</sup> and *Syk*Δ<sup>MG</sup> combined brain and spinal cord samples. (D) Expression levels of *Sykb* mRNA from MACS-sorted microglia quantified by qPCR. (E) Levels of SYK protein (top panel) and total protein loaded (bottom panel) from MACS-sorted microglia determined by Western blotting and SDS-PAGE with a stain-free gel, respectively. (F) Quantification of intensity of SYK protein bands normalized to intensity of bands from total protein loaded. (G-H) *Syk<sup>Con</sup>* and *Syk*<sup>ΔMG</sup> mice received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Spleens were harvested at 5 months of age to evaluate SYK expression in T cells and monocytes. (G) Levels of SYK protein bands normalized to total Actin in sorted splenic T cells and monocytes from *Syk<sup>Con</sup>* and *Syk<sup>ΔMG</sup>* mice. Statistical significance between experimental groups was calculated by unpaired Student's *t* test (B, D, F, H). ns = not significant, \*\*\*p < 0.0001, \*\*\*\*p < 0.0001. Error bars represent mean ± SEM and each data point represents an individual mouse.







## Figure S2. Microglial SYK deletion in the hippocampus and after disease onset significantly alters AD pathology and microgliosis, however, the loss of SYK does not affect microgliosis in the absence of $A\beta$ , related to Figures 1, 2, and 3

(A-G) 5xFAD *Syk*<sup>AMG</sup> mice and 5xFAD littermate controls received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Brains were then harvested at 5 months of age to evaluate plaque sphericity and microgliosis in the hippocampus. (A-B) Microglial response to plaques measured by (A) the sphericity of ThioS-labeled and Imaris-rendered A<sub>β</sub> plaques in the hippocampus of matched sagittal sections. (B) Quantification of sphericity with 1.00 being the most spherical, combined data from a total of 50–100 plaques from 3 matching brain sections per mouse. (C-E) Microglia were imaged by labeling with lba1 (green) surrounding A<sub>β</sub> plaques labeled with ThioS (pink) to assess microglial coverage and proximity to plaques. (C) Representative images of lba1 and ThioS staining in the hippocampus of matching sagittal brain sections. (D) Quantification of microglia numbers in the field of view (FOV) in 40  $\mu$  m images. (E) Quantification of the number of microglia within a 15 and 30  $\mu$  m radius surrounding ThioS-labeled A<sub>β</sub> plaques. (F) Representative Imaris rendering of lba1<sup>+</sup> microglia in the hippocampus of 5xFAD and 5xFAD Syk<sup>AMG</sup> mice. (G) Sholl analysis quantification from a total of 12 microglia from 3 matching brain sections per mouse (5xFAD n = 7, 5xFAD Syk<sup>AMG</sup> n = 7). (H-N) 5xFAD Syk<sup>AMG</sup> mice and 5xFAD mice received tamoxifen food for 2 weeks starting at 4 months of age and then mice were returned to regular food for the remainder of the experiment. Brains were then harvested at 8 months of age in this delayed deletion model to evaluate amyloid beta (A<sub>β</sub>) load in the brain. (H-I) Immunofluorescence staining of A<sub>β</sub> (D54D2, red; DAPI, blue) was performed on sagittal sections. (K) Quantification of spherical, combined data from a total of 50–100 plaques from 3 matching brain sections per mouse. (L-N) Microglia were imaged by labeling with 1.00 being the most spherical, combined data from a total of 50–100 plaques from 3 matching bra





Quantification of microglial numbers in the FOV in 40  $\mu$  m images. (N) Quantification of the number of microglia within a 15 and 30  $\mu$  m radius surrounding ThioSlabeled A $\beta$  plaques. (O-X) 5xFAD *Syk*<sup>ΔMG</sup> mice, 5xFAD mice, *Syk*<sup>ΔMG</sup> mice, and *Syk*<sup>con</sup> mice received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. (O-Q) Mouse body weight was measured at 3 and 6 months of age, while memory and anxiety-related behaviors were evaluated in the Morris water maze (MWM) and elevated plus maze (EPM) at 4 months of age. (O) Mouse body weight recorded in grams (g) across experimental groups. (P) Average velocity in the MWM on day 4 of the acquisition phase of the test. (Q) Distance traveled in the EPM. (R-V) Brains from *Syk*<sup>ΔMG</sup> and *Syk*<sup>con</sup> mice were harvested at 5 months of age to evaluate microgliosis in the absence of A $\beta$ . (R) Iba1 (green) and Ki67 (blue) staining was performed on sagittal sections to evaluate *Syk*<sup>ΔMG</sup> and *Syk*<sup>con</sup> microglial numbers and proliferation. (S) Quantification of microglial numbers in the FOV averaged from 3 matching cortical sections per mouse. (T) Quantification of Ki67<sup>+</sup> microglia in the FOV of the cortex and hippocampus of mice averaged from 3 matching brain sections per mouse. (U) Representative Imaris rendering of Iba1+ microglia in the cortex of *Syk*<sup>Con</sup> and *Syk*<sup>ΔMG</sup> mide *Syk*<sup>ΔMG</sup> and *Syk*<sup>ΔMG</sup> n = 5). (W-X) Brains were harvested from 5xFAD *Syk*<sup>ΔMG</sup> and 5xFAD microglia apoptosis by TUNEL staining. (W) 5xFAD *Syk*<sup>ΔMG</sup> and 5xFAD microglia labeled with Iba1 (green) and TUNEL (pink) surrounding A $\beta$  plaques. (X) Quantification of TUNEL volume within Iba1<sup>+</sup> microglia as a measure of apoptosis. Statistical significance between experimental groups was calculated by an unpaired Student's t test (B, D-E, I, K, M–N, P-Q, S-T, X), two-way ANOVA with a Bonferroni post-hoc test (G, V), and one-way ANOVA with multiple comparisons (Q). ns = not significant, \*p < 0.05, \*\*p < 0.01, \*\*





### Figure S3. Effects of SYK deficiency on microglial gene expression under steady-state conditions and in response to A $\beta$ pathology, related to Figure 4

(A-K) SxFAD Syk<sup>AMG</sup>, 5xFAD, Syk<sup>AMG</sup>, and Syk<sup>con</sup> mice received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Brains were later harvested at 5 months of age to evaluate microglial activation. (A-D) Microglia from 5-month-old 5xFAD Syk<sup>AMG</sup>, 5xFAD, Syk<sup>AMG</sup>, and Syk<sup>con</sup> mice were sorted from single-cell brain suspensions using anti-CD11b<sup>+</sup>-coated magnetic beads and magnetic column sorting, finally, RNA-Seq was performed. (A) Representative flow cytometry gating strategy used to validate purity of MACS-sorted microglia from 5xFAD Syk<sup>AMG</sup>, 5xFAD, Syk<sup>AMG</sup>, and Syk<sup>con</sup> brain samples. (B) Volcano plots depicting differentially expressed genes (FDR<0.1) where 37 genes are downregulated and 7 genes are upregulated in Syk<sup>AMG</sup> microglia compared to Syk<sup>con</sup> microglia. (C) Heatmap representation of the 20 most significantly upregulated and downregulated (FDR<0.1) genes between Syk<sup>AMG</sup> and Syk<sup>con</sup> mice. (E-I) Immunohistochemistry validation of RNA-Seq findings in 5-month-old 5xFAD Syk<sup>AMG</sup> and 5xFAD mice. (E-I) Immunohistochemistry validation of RNA-Seq findings in 5-month-old 5xFAD Syk<sup>AMG</sup> and 5xFAD mice. (E) Jamen 119 (pink) surrounding Aβ plaques labeled with ThioS (blue). (F) Quantification of Tmem119 vol colocalized with Iba1. (G) 5xFAD Syk<sup>AMG</sup> and 5xFAD microglia labeled with Iba1 (green) and CLEC7A area found surrounding  $\sim$ 30 plaques from 3 matching brain sections per mouse.





<sup>(</sup>J-K) Comparison of the transcriptional changes that arise in microglia with the loss of either SYK or TREM2 in 5xFAD mice.  $5xFAD Syk^{\Delta MG}$  and 5xFAD brains were harvested at 5 months of age and their microglia were sorted from single-cell brain suspensions using anti-CD11b<sup>+</sup>-coated magnetic beads and magnetic column sorting to perform RNA-Seq. 8-month-old 5xFAD  $Trem2^{-/-}$  and littermate 5xFAD microglia were FACS-sorted and analyzed by RNA-Seq in (Griciuc et al., 2019). (J) Venn diagram of significantly upregulated and downregulated genes in 5xFAD  $Trem2^{-/-}$  and 5xFAD  $Syk^{\Delta MG}$  microglia compared with their respective littermate 5xFAD controls (FD < 0.05). (K) Molecular function (MF) term enrichment scatterplot highlighting major functions that are repressed in 5xFAD  $Trem2^{-/-}$  and 5xFAD  $Syk^{\Delta MG}$  microglia in comparison to littermate 5xFAD microglia. Statistical significance between experimental groups was calculated by unpaired Student's *t* test (F, H-I). \*p < 0.05, \*\*\*p < 0.001. Error bars represent mean  $\pm$  SEM and each data point represents an individual mouse.



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Figure S4. Genetic ablation of SYK in microglia leads to increased levels of microglial lipid droplet accumulation and ROS production in 5xFAD mice, related to Figure 5

(A-D) 5xFAD  $Syk^{\Delta MG}$  mice and 5xFAD littermate controls received tamoxifen food for 2 weeks starting at 4 months of age and then mice were returned to regular food for the remainder of the experiment. Microglia from 8-month-old 5xFAD  $Syk^{\Delta MG}$  and 5xFAD mice were sorted from single-cell brain suspensions using flow cytometry. (A) Representative histograms of BODIPY-labeled lipid-droplet accumulation in CD11b<sup>+</sup>CD45<sup>int</sup> microglia. (B) Mean Fluorescence Intensity (MFI) quantification of BODIPY in 5xFAD and 5xFAD  $Syk^{\Delta MG}$  microglia. (C) Representative histograms of CellROX-labeled reactive oxygen species (ROS) in CD11b<sup>+</sup>CD45<sup>int</sup> microglia. (D) MFI quantification of CellROX in 5xFAD and 5xFAD and 5xFAD and 5xFAD syk^{\Delta MG} microglia. (E-F) 5xFAD  $Syk^{\Delta MG}$  mice and Cre-negative 5xFAD mice received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Brains were later harvested and Iba1 staining was performed at 5 months of age to evaluate microglial activation. (E) 5xFAD  $Syk^{\Delta MG}$  and 5xFAD microglia labeled with Iba1 (green) and CD68 (blue) surrounding A $\beta$  plaques labeled with ThioS (pink). (F) Volumetric quantification of CD68 normalized to Iba1 volume. Statistical significance between experimental groups was calculated by unpaired Student's *t* test (B, D, F). ns = not significant, \*p < 0.05. Error bars represent mean  $\pm$  SEM and each data point represents an individual mouse.







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Figure S5. Syk<sup> $\Delta$ MG</sup> mice have increased numbers of cytokine producing T cells infiltrating the CNS but have modestly reduced peripheral T cell responses during EAE, related to Figure 6

(A-H) *Syk*<sup>ΔMG</sup> mice and *Syk*<sup>con</sup> littermate controls received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Mice were later immunized with MOG + CFA and pertussis toxin at 8–14 weeks of age to induce experimental autoimmune encephalomyelitis (EAE). (A-E) Spinal cords were then harvested from mice during the effector phase of clinical disease (at least 30 days post-immunization) to evaluate immune cell responses by flow cytometry. (A) Representative flow cytometry gating strategy. (B) Quantification of immune cell populations. (C) Representative dot plots of effector cytokine-producing CD4<sup>+</sup> T cells after no stimulation or 5 h *ex vivo* stimulation with PMA and ionomycin in the presence of monensin. (D) Quantification of the frequencies of effector cytokine producing CD4<sup>+</sup> T cells. (E) Quantification of the total numbers of effector cytokine producing CD4<sup>+</sup> T cells. (F-H) Spleens were harvested from mice during the effector phase of clinical disease (at least 30 days post-immunization) to evaluate immune cell responses by flow cytometry. (F) Representative dot plots of EAE effector cytokine producing CD4<sup>+</sup> T cells. (E) Quantification of the total numbers of effector cytokine producing cD4<sup>+</sup> T cells. (F-H) Spleens were harvested from mice during the effector phase of clinical disease (at least 30 days post-immunization) to evaluate immune cell responses by flow cytometry. (F) Representative dot plots of EAE effector-cytokine producing CD4<sup>+</sup> T cells from EAE effector phase spleens after no stimulation or 5 h *ex vivo* stimulation with PMA and ionomycin in the presence of monensin. (G) Quantification of effector cytokine producing by CD4<sup>+</sup> T cells. T cells from EAE effector phase spleens after no stimulation or 5 h *ex vivo* stimulated *ex vivo* with MOG<sub>35-55</sub> peptide for 48 h. Levels of indicated analytes were measured by multiplex cytokine assay. (I) Flow cytometry-based sorting of ZsGreen<sup>+</sup> microglia from Syk<sup>Δ</sup>







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Figure S6. Disruption of SYK signaling in microglia during demyelinating disease leads to impaired microglial response and reduced OPC proliferation, related to Figure 7

(A-B)  $Syk^{\Delta MG}$  mice and  $Syk^{con}$  littermate controls received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food for the remainder of the experiment. Mice were harvest at 8 months of age and total myelin levels were evaluated in the corpus callosum. (A)  $Syk^{\Delta MG}$  mice and  $Syk^{con}$  corpus callosum labeled with myelin basic protein (MBP; green) and DAPI (blue). (B) Quantification of percent area covered by MBP in the corpus callosum at baseline. (C-J)  $Syk^{\Delta MG}$  mice and  $Syk^{con}$  mice received tamoxifen food for 2 weeks starting at 3 weeks of age and then mice were returned to regular food. Adult (8–12 month old) mice were later fed a diet consisting of 0.3% cuprizone for 5 weeks to induce demyelination. Mice were then either harvested after 5 weeks of cuprizone treatment (demyelination group) or returned to normal chow for one additional week before being harvesting (remyelination group). Control mice were not introduced to the cuprizone diet. (C-D) Representative images and quantification of the number of Iba1<sup>+</sup> cells (green) in the corpus callosum. (E-F) Representative images of Iba1<sup>+</sup> microglia (green) and CD68 (pink) expression in the corpus callosum. (H) Volume of CD68 colocalized to the volume of Iba1 in the corpus callosum. (G) Representative images of proliferating Ki67<sup>+</sup> (pick) Pdgfra<sup>+</sup> Olig2<sup>+</sup> (blue; green) oligodendrocyte precursor cells (OPCs) in the corpus callosum. (J) Quantification of Ki67<sup>+</sup> OPCs in the corpus callosum. Statistical significance between experimental groups was calculated by unpaired Student's *t* et (B) and one-way ANOVA with multiple comparisons (D, F, H, J). ns = not significant, \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. Data are mean ± SEM and combined from two independent experiments and each data point represents an individual mouse.