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SUPPLEMENTARY DATA

Spatiotemporal dysregulation of neuron-glia related genes and pro-/anti-inflammatory miRNAs in the *5xFAD* mouse model of Alzheimer's disease

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Supplementary Table S1	Gene expression profile by RT-qPCR analysis revealed no significant differences when simultaneously considering the genotype (WT/5xFAD), age (6/9 months) and brain region (HPC, hippocampus/PFC, prefrontal cortex).	Link
Supplementary Table S2	miRNA-target table for the Analyzed microRNAs and targets selected for this study. Obtained in the online platform https://www.mirnet.ca/	Link
Supplementary Table S3	Node table for the analyzed microRNAs and targets selected for this study. We only considered miRNAs and/or targets with a node degree of at least 2. Obtained in the online platform https://www.mirnet.ca/	Link
Supplementary Table S4	Bivariate Pearson's correlation coefficients and respective *p*-values obtained between all miRNAs and genes.	Link
Supplementary Table S5	List of microRNAs analyzed by RT-qPCR and their primer sequences.	Link
Supplementary Table S6	List of genes and respective primer sequences used for mRNA analysis by RT-qPCR.	Link
Supplementary Table S7	Raw data used for correlational analysis in hippocampus (HPC) and prefrontal cortex (PFC) using the corrplot package in RStudio software.	Link