

Description of Additional Supplementary Files

Supplementary Data 1. Sequencing statistics of datasets

Supplementary Data 2. Differentially expressed genes across developmental stages

Supplementary Data 3. Quality control for HiCAR datasets

Supplementary Data 4. Differentially expressed genes of astrocytes depend on NR3C1 cKO

Supplementary Data 5. List of nuclei obtained from snRNA-seq datasets

Supplementary Data 6. List of DEGs of NR3C1 cKO astrocytes at EAE peak

Supplementary Data 7. Motif enrichment of cCREs for EAE upregulated gene

Supplementary Data 8. Motif enrichment of cCREs for EAE downregulated gene