

Description of Additional Supplementary Files

Supplementary Data 1:

TRAP sequencing of astrocytes during remyelination. Fragments per Kilobase Million (FPKM) and Log2 fold change of genes identified from TRAPseq of Aldh1l1-Rpl10- eGFP LPC-demyelinated lesions at 3, 7, and 10 days post injection (DPI) and no lesion controls. DE-Seq2 and Benjamini-Hochberg test, adjusted P values indicated.