

Description of Additional Supplementary Files

File: Supplementary Data 1.

Description: Quantification of myelinated axons/100 μm^2 per animal in normal and cuprizone-treated animals.

File: Supplemental Data 2.

Description: All significantly enriched gene ontology terms for genes downregulated with miR-145-5p overexpression in OLs.

File: Supplemental Data 3.

Description: All significantly enriched gene ontology terms for genes upregulated with miR-145-5p overexpression in OLs.