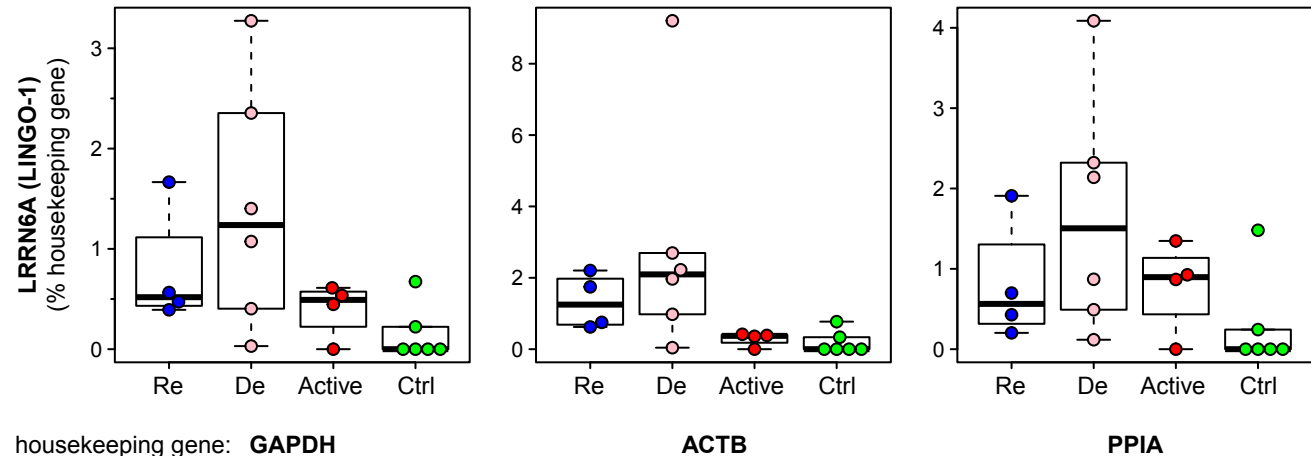




Suppl. Figure 2: LRRN6A (LINGO-1) gene expression in different MS lesion types and control white matter. LINGO-1 was quantified by qPCR TLDA and is shown normalized to the 3 housekeeping genes GAPDH, ACTB, and PPIA. Each symbol represents one dissected lesion area or control tissue. Re: remyelinated, De: demyelinated inactive, Active: actively demyelinating lesions; Ctrl: control white matter. Boxplots include medians, 1st/3rd quartiles, and whiskers extending up to 1.5x of the interquartile range.