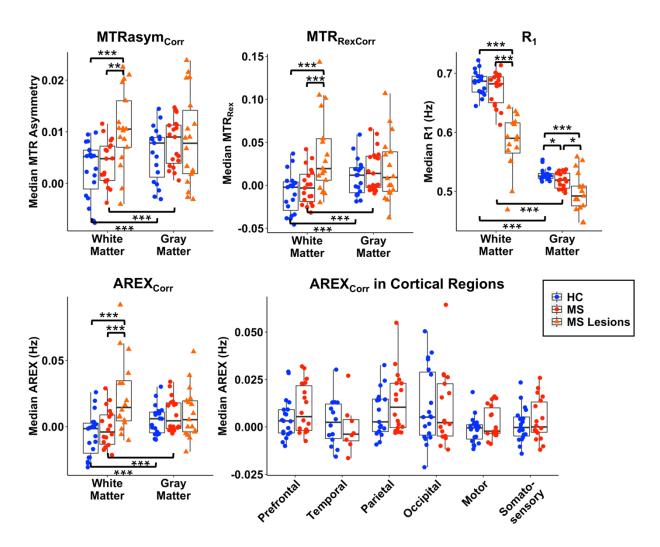
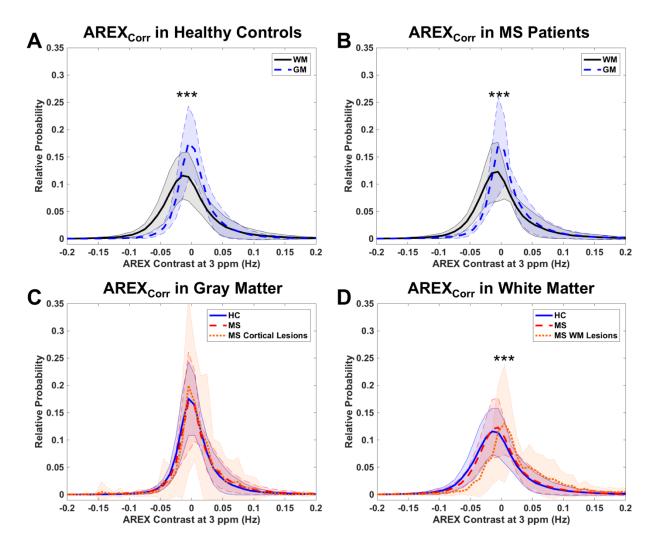


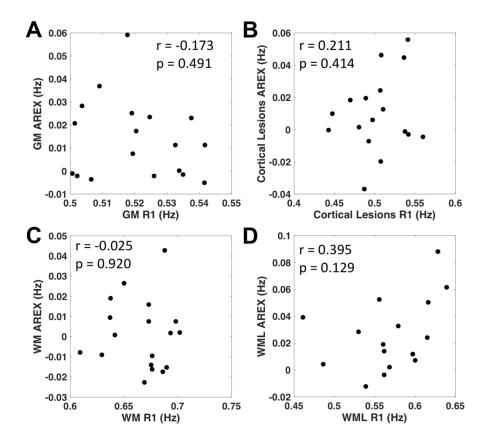
## Supplementary Material



**Supplementary Figure 1.** Median values of MTRasym<sub>Corr</sub>, MTR<sub>RexCorr</sub>, AREX<sub>Corr</sub>, and R<sub>1</sub> are shown for WM, GM, WM lesions, and cortical lesions for group comparisons. AREX<sub>Corr</sub> values are also shown for each cortical GM region. WM lesion values differ significantly from normal-appearing WM in MS patients and from healthy control (HC) WM for all CEST-derived indices and for R<sub>1</sub>. Significant differences in GM and cortical lesions are only present in R<sub>1</sub>. No group differences are significant for AREX<sub>Corr</sub> within cortical regions. All indices differ significantly between GM and WM within each group. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001



**Supplementary Figure 2.** Average histograms of AREX<sub>Corr</sub> values are shown for WM, GM, WM lesions, and cortical lesions for within-group (**A**, **B**) and between-group (**C**, **D**) comparisons (shaded area represents standard deviation). AREX<sub>Corr</sub> values are significantly greater in GM than in WM within the healthy control (HC) and MS patient groups. AREX<sub>Corr</sub> is increased significantly in WM lesions compared to normal-appearing WM in MS patients and compared to WM in healthy volunteers. \*\*\*p<0.001 for mean value comparison.



**Supplementary Figure 3.** Mean  $AREX_{Corr}$  values do not correlate with corresponding mean  $R_1$  values in each tissue region of interest in patients with MS. Plots are shown for normal-appearing GM **(A)**, cortical lesions **(B)**, normal-appearing WM **(C)**, and WM lesions **(D)** with the corresponding Pearson correlation coefficient r and p-value.