

eFigure 9 Serum neurofilament light chain (sNfL) levels in young and old MS patients. sNfL levels were evaluated in the serum of young (≤ 50 years) and old (≥ 50 years) patients with multiple sclerosis (MS) (young: n=35, old: n=34; relapsing-remitting MS (RRMS): young: n=16, old: n=14; primary progressive (PPMS): young: n=19, old: n=20). Demographic data of study subjects are depicted in eTable 1. (A) sNfL levels of MS patients. (B) sNfL levels of RRMS and PPMS patients. (C) Correlation analysis of sNFL levels with age of MS patients (n=69). (D) Correlation analysis of sNFL levels with the Expanded Disability Status Scale (EDSS) score of MS patients (n=69). (E) Correlation analysis of sNFL levels with disease duration of MS patients (n=69). (F) Correlation analysis of sNFL with frequencies of CD28⁻ CD4 T cell of young (*left*) and old (*right*) MS patients. (G) Correlation analysis of sNFL with frequencies of CD28⁻ CD8 T cells of young (left) and old (right) MS patients. (H) Correlation analysis of sNFL with frequencies of CD57⁺ CD4 T cells of young (*left*) and old (*right*) MS patients. (I) Correlation analysis of sNFL with the expression of KLRG1 on memory CD8 T cells of young (left) and old (right) MS patients. (J) Correlation analysis of sNFL with the expression of KLRG1 on central memory (CM) CD8 T cells of young (left) and old (right) MS patients. Data are displayed as boxplots of the median and the 25th and 75th percentile ± IQR. Statistical analysis was conducted by two-tailed Mann-Whitney test. For correlation analysis, the Pearson product-moment correlation coefficients (Pearson's R) were computed. Differences were considered statistically significant with the following P-values: *P < 0.05, **P < 0.01, ***P < 0.010.001 and ****P < 0.0001