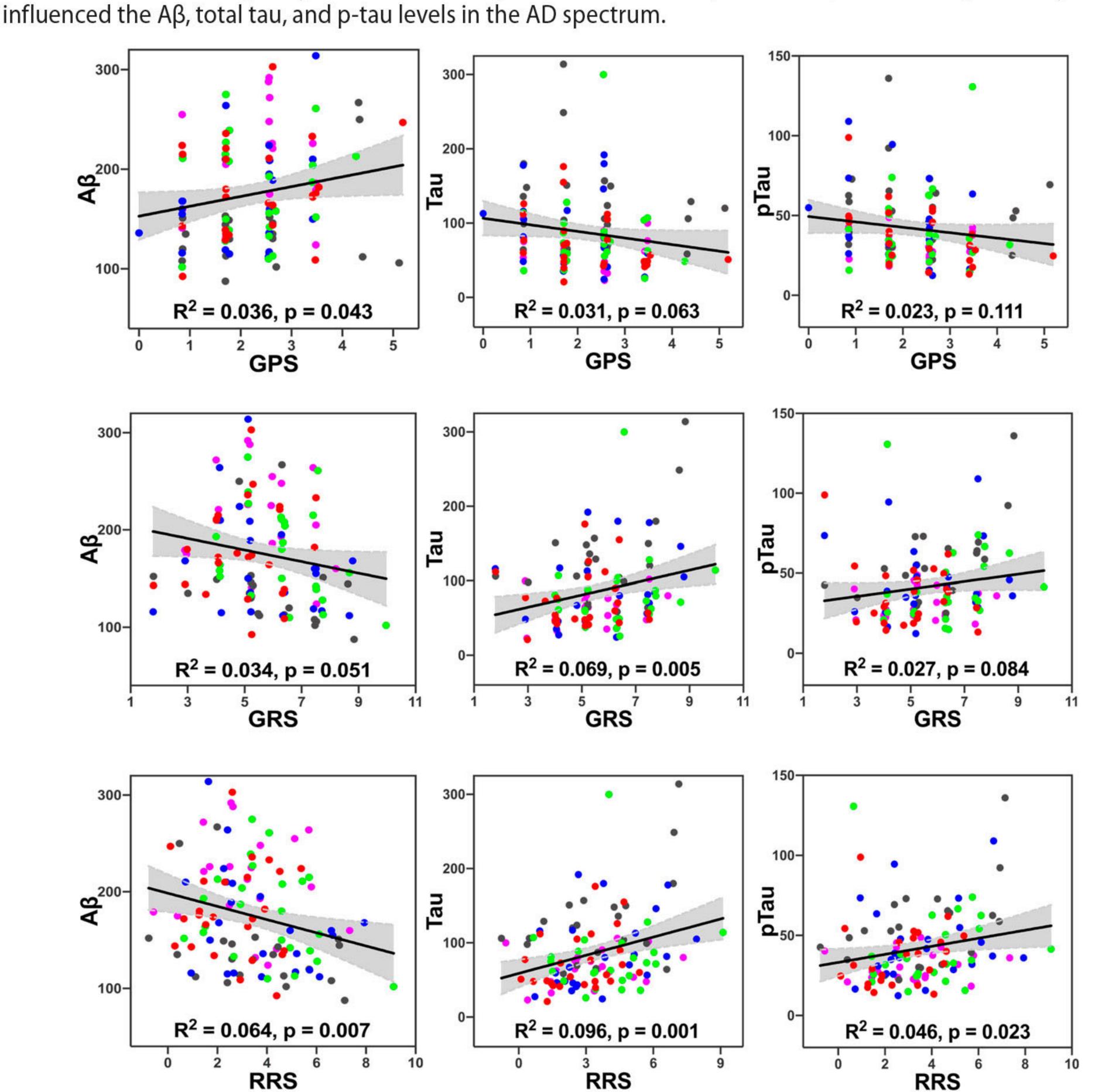
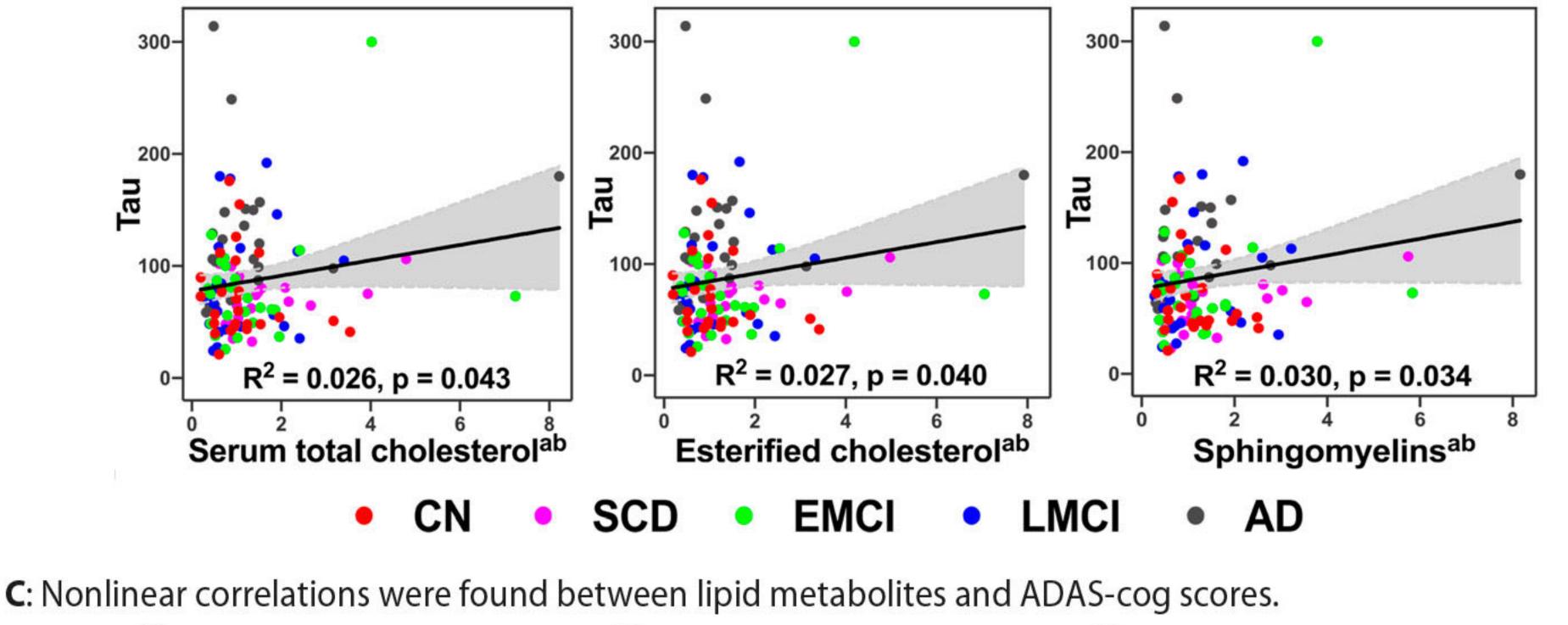
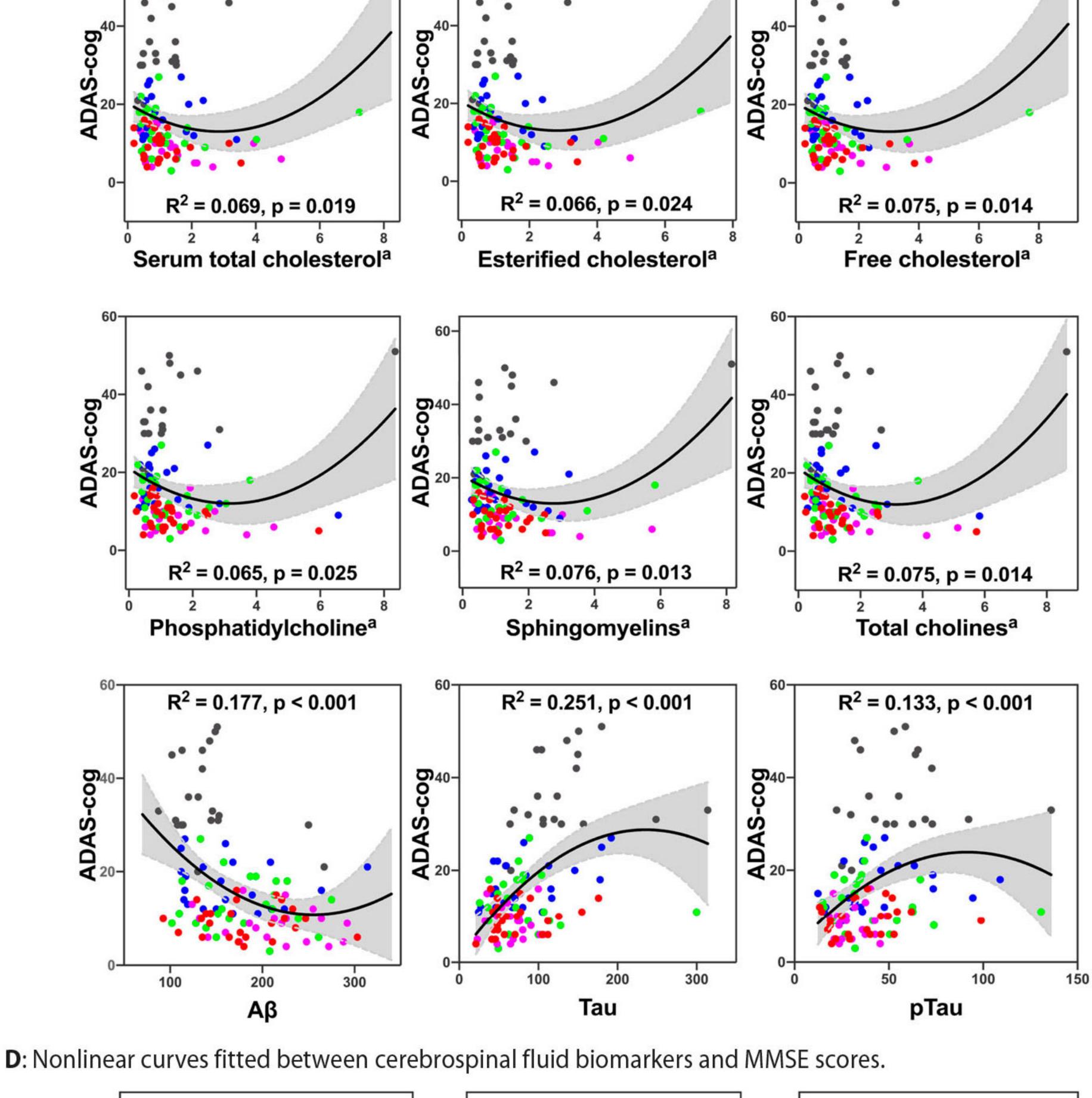
Figure 1. Regression Analyses Among Polygenic Scores, Serum Lipid Metabolites, a Cerebrospinal Fluid Core Biomarkers, and Cognitive Performance in the AD Spectrum*

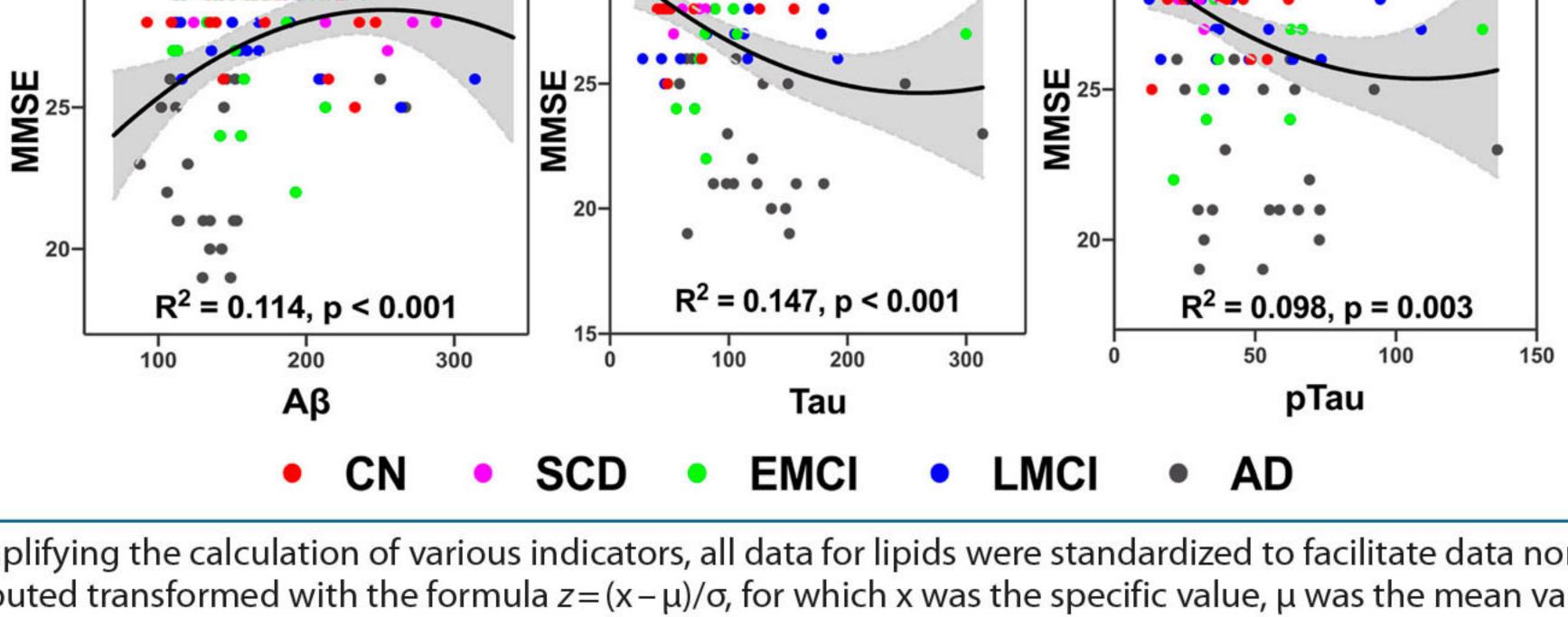
A: Linear regression between polygenic scores and cerebrospinal fluid biomarkers revealed that the GPS was correlated with AB levels but not total tau and p-tau levels; in contrast, GRS was significantly correlated with total tau levels and marginally associated with Aß levels but not with p-tau levels, while RRS significantly



B: Significant linear correlations were found between several lipid metabolism-related markers and total tau but not Aβ and p-tau levels of cerebrospinal fluid in the AD spectrum.







30-

^aFor simplifying the calculation of various indicators, all data for lipids were standardized to facilitate data normally distributed transformed with the formula $z = (x - \mu)/\sigma$, for which x was the specific value, μ was the mean value, and σ the standard deviation. Then, the z values of lipids were transformed using an inverse logarithm algorithm with a base of 2. Considering the opposite effects of HDL compared with LDL or VLDL, the values of HDL-related markers

30-

were further converted into their reciprocal form. ^bOne-tailed *P* values < .05 were considered significant only in Figure 1B. *Gray bands indicated 95% confidence intervals.

Abbreviations: $A\beta = amyloid \beta$ peptides 1 to 42, AD = Alzheimer's disease, ADAS - cog = Alzheimer's Disease

Assessment Scale-13-item cognitive subscale, CN = cognitively normal, EMCI = early amnestic mild cognitive impairment, GPS = genetic protective score, GRS = genetic risk score, HDL = high-density lipoprotein, LDL = lowdensity lipoprotein, LMCI = late mild cognitive impairment, MMSE = Mini-Mental State Examination, p-tau = tau phosphorylated at the threonine-181 position, RRS = relative risk score, SCD = subjective cognitive decline, VLDL = very low density lipoprotein.