

significant associations (p<0,05 and passing FDR corrected at q=0.2) are displayed. Associations are

colored according to the Spearman's p, with positive associations on red and negative on blue scale.

Proteins are colored according to Gene Ontology (GO) annotations. Additionally, proteins involved in glycolysis, blood coagulation and extracellular matrix (ECM) interaction are indicated. Spearman's ρ

cases = 151; healthy controls = 142. Two splicing variants are reported for KNG1 (P01042 and P01042-2,

correlations between all analyzed CSF proteins and CSF metabolites are presented in Table S3. N for AD

marked with asterisk).