Figure 3: Mass Spectrometry Analysis (MS /MS) of APOA2 and A2MG

The MS /MS spectra show the results of mass spectrometry analysis for two proteins, APOA2 (apolipoprotein A- II) and A2MG (Alpha-2-macroglobulin). The MS /MS analysis provides detailed information about the peptide sequences and possible post-translational modifications (PTMs) in these proteins.

Table 3: Presents a summary of identified peptides with their corresponding fold change (FC) values, log2-transformed fold change (log2(FC)), p-values, and the negative logarithm of p-values (|-LOG10(p)|).