Figure S1

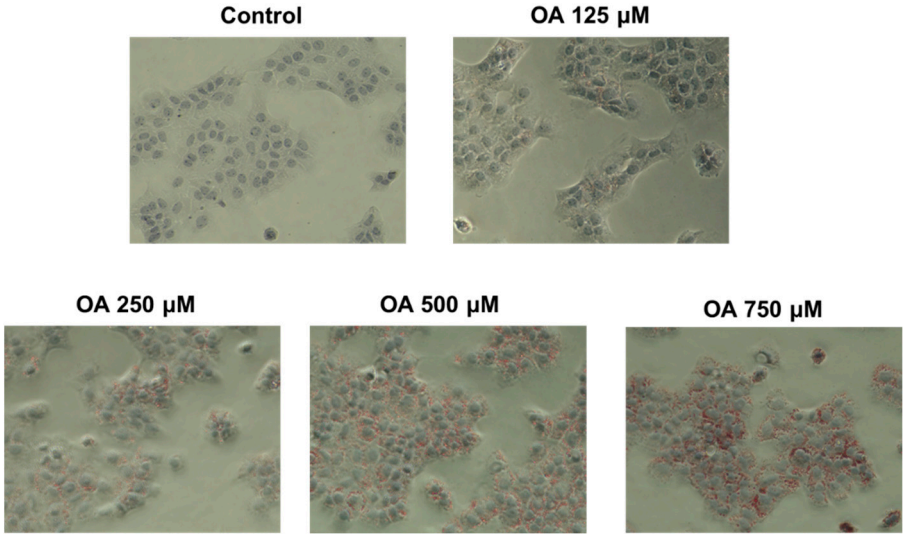


Figure S1. Representative views of oil red O staining of AML-12 cells after a series of dose of OA incubations.

Figure S2

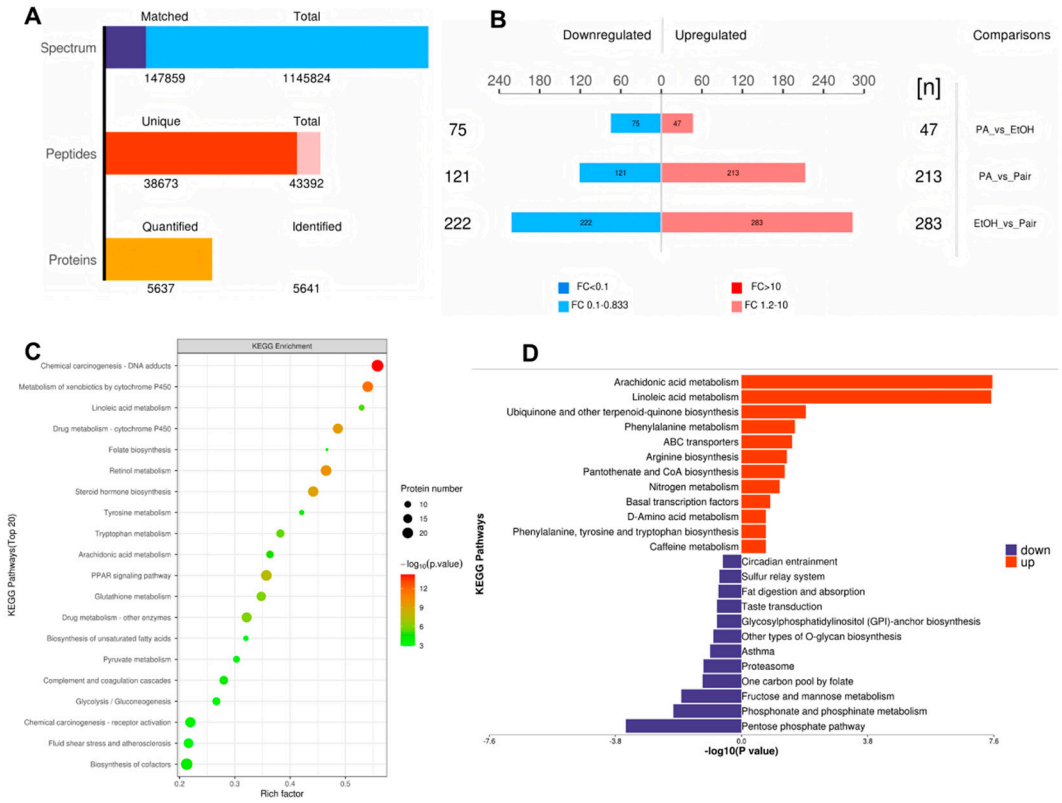


Figure S2. The data of proteomic analysis. (A) The number of chromatograms, peptide segments, proteins and identified proteins in proteomics. (B) The number of DEPs identified in the various comparison groups. (C) KEGG functional enrichment analysis of DEPs in EtOH group versus

Control group. (D) The separate enrichment analyses for up- and down-regulation DEPs in EtOH group versus Control group.

Figure S3

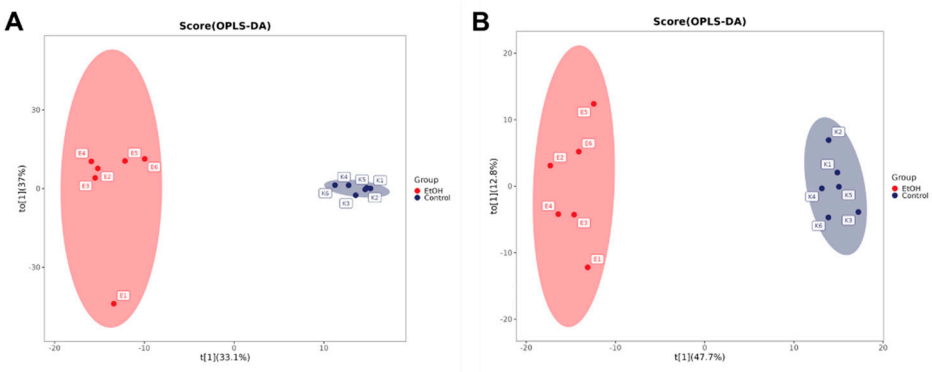


Figure S3. OPLS-DA scores of samples in EtOH and Control group in positive mode (A) and negative mode (B), respectively.