



Supplementary Fig. 4. Functional enrichment by Gene Ontology (GO) and Kyoto Encyclopedia of Genes and Genomes (KEGG) analysis. (A, B) GO analysis based on biological process (BP), cellular component (CC), and molecular function (MF). (C, D) KEGG analysis. PI3K, phosphoinositide 3-kinase; ECM, extracellular matrix; AGE-RAGE, advanced glycation end-products-receptor for advanced glycation end-products.