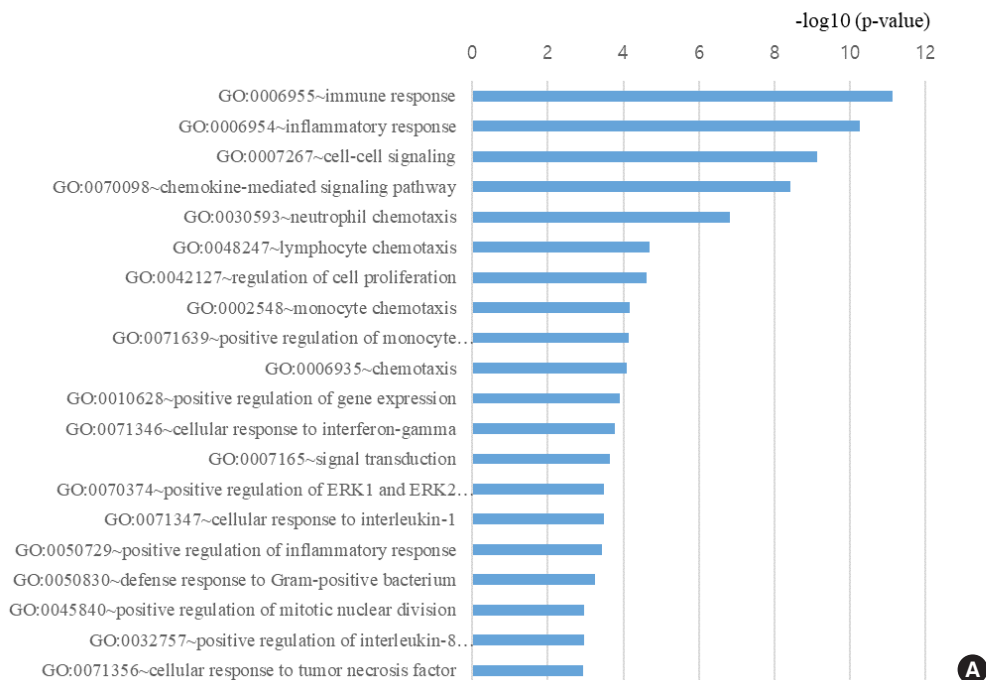
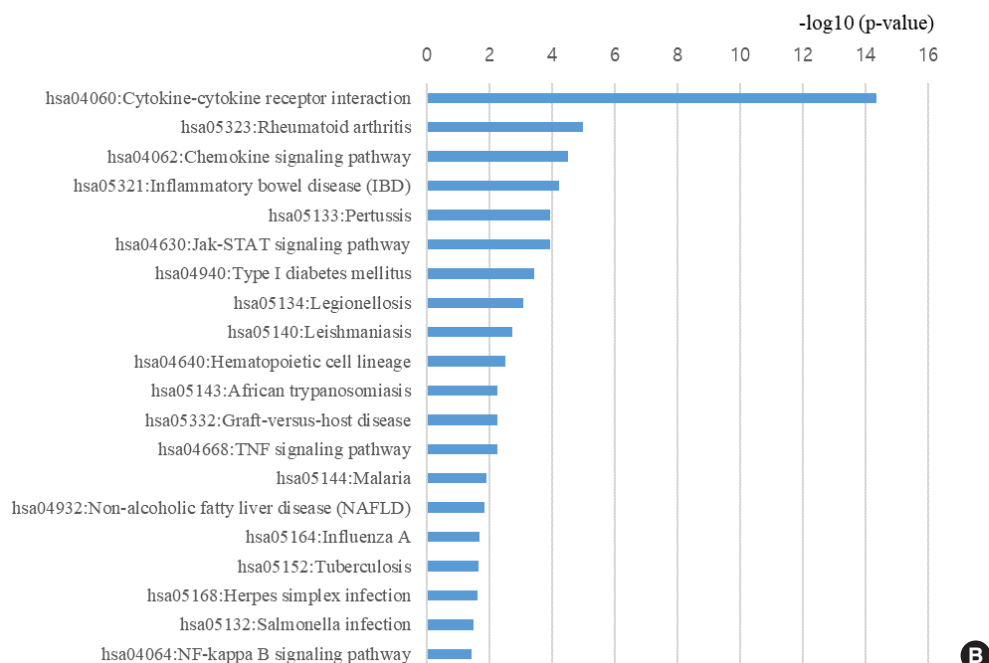


Top 20 enrichment GO Biological process



Top20 enrichment KEGG pathway



Supplemental Fig. S3. Gene Ontology (GO) term enrichment analysis for biological process (A) and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway analysis (B) of the cytokines with 1.5-fold difference between subjects with normoglycemia and patients with prediabetes. Analysis was performed using the Database for Annotation, Visualization and Integrated Discovery (DAVID) tool with the human genome as a reference.