

Suppl 3. Hierarchical clustering heatmaps of DEGs screened in blood based on $|\log_2 \text{fold change}| > 1$ and a corrected P-value < 0.05 were shown, respectively. Notes: (A) CD samples, (B) UC

samples; Red indicates that the expression of genes is upregulated, blue means that the expression of genes is downregulated, and * indicates gene expression value > 5.0 .

