



Supplementary Figure 2. CD8⁺ T cells co-cultured with WT or GPR84^{-/-} MDSCs were used for RNA-seq.

(A) Principal component analysis (PCA) of the gene expression profiles between groups. (B) Volcano map of differential genes. (C) Heat map of the hierarchical clustering of the gene expression matrix. (D) Gene ontology (GO) enrichment analysis. (E) Gene Set Enrichment Analysis (GSEA) analysis.