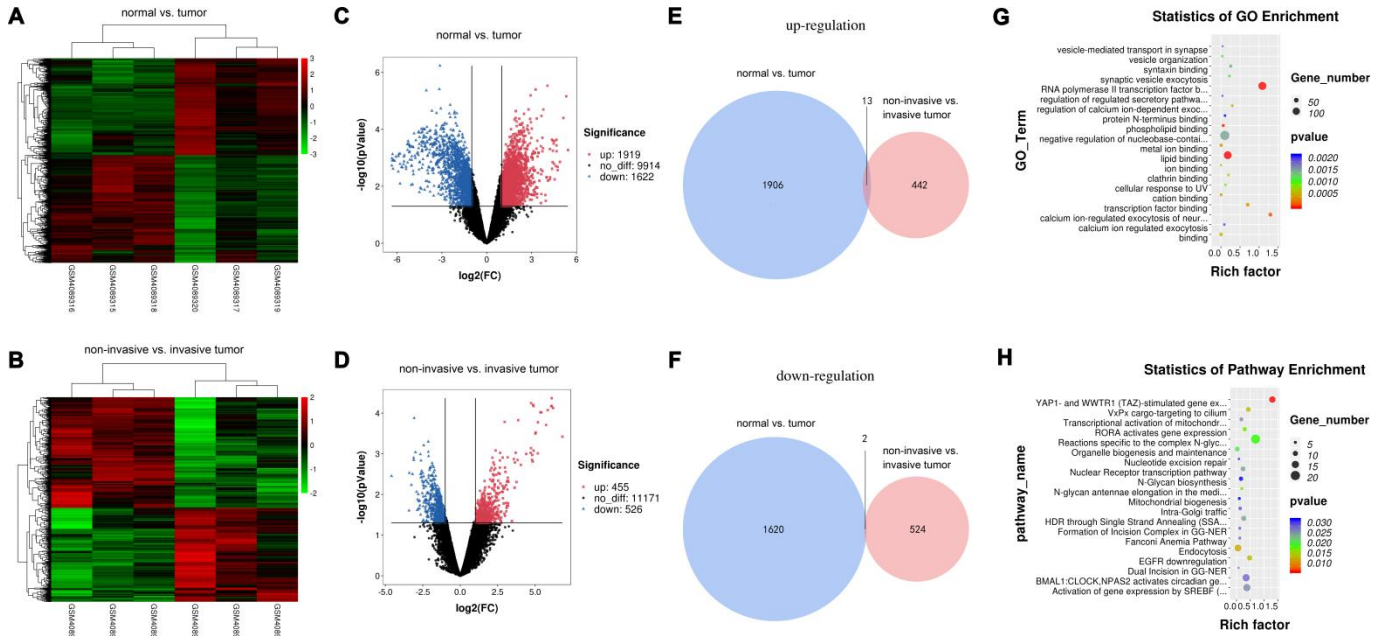


SUPPLEMENTARY FIGURE



Supplementary Figure 1. Screening of ccRCC-related circRNAs in multiple microarrays. (A, B) Hierarchical clustering analysis and volcano plots of significantly differentially expressed circRNAs in the tumor or adjacent normal tissues from three patients with ccRCC. (C, D) Hierarchical clustering analysis and volcano plots of significantly differentially expressed circRNAs in the invasive tumor or non-invasive tumor tissues from three patients with ccRCC. (E, F) The Venn diagrams demonstrated the up-regulated and down-regulated circRNAs that overlapped during the comparison between tumor and adjacent normal tissues, or between invasive tumor and non-invasive tumor tissues from ccRCC. (G, H) GO enrichment and pathway analysis for dysregulated circRNAs gene symbols.