



Figure S1 The differential expression of mRNAs between PC samples and normal tissues was analyzed. (A) Volcano plots were constructed using fold-change values and adjusted P. The red point in the plot is the over-expressed mRNAs, and the blue point shows the down-expressed mRNAs with statistical significance. (B) Hierarchical clustering analysis of mRNAs, which were differentially expressed between tumor and normal tissues. (C-E) The results of differentially expressed genes of KEGG-DOWN, GO-UP, and GO-DOWN.