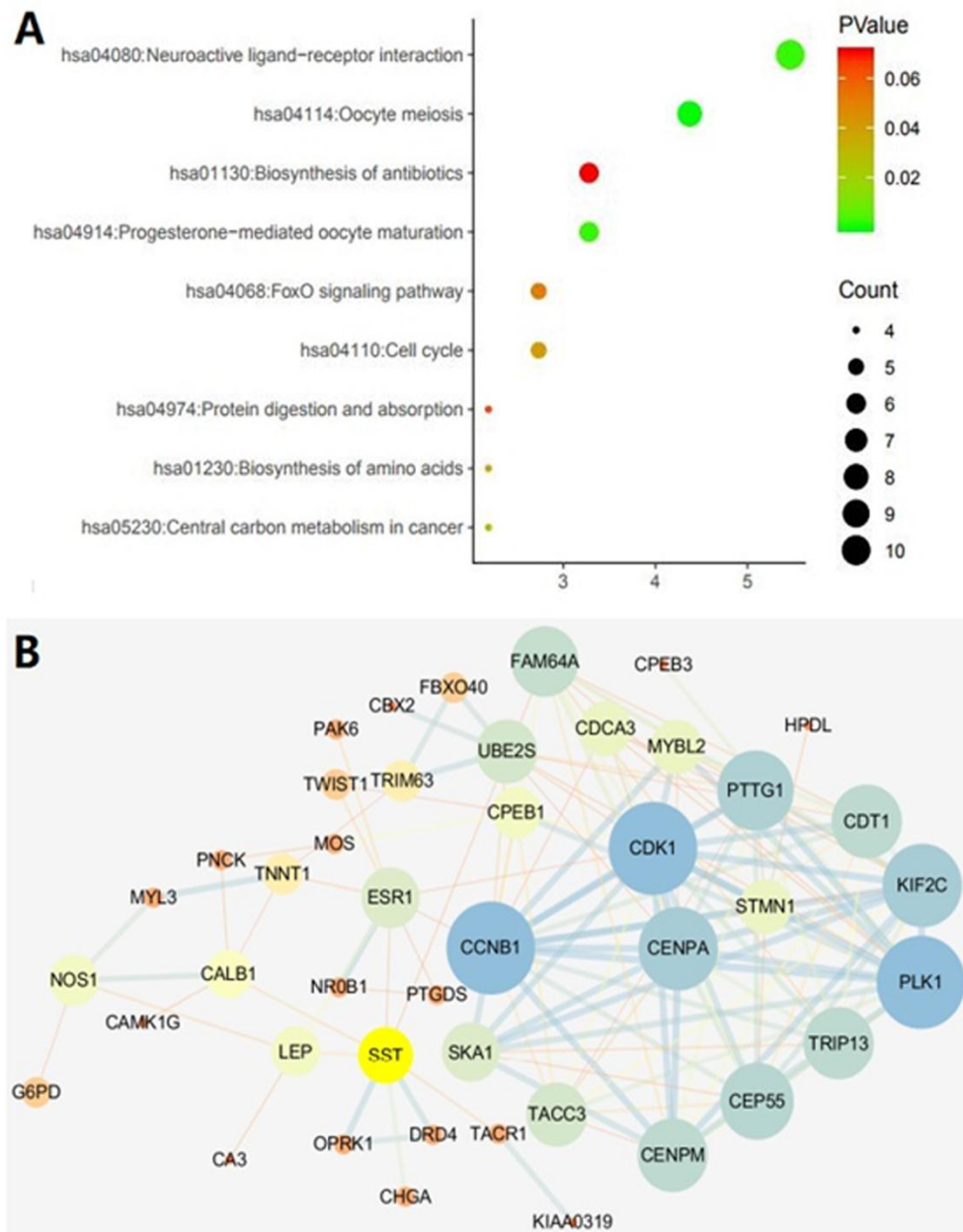


SUPPLEMENTARY FIGURE



Supplementary Figure 1. The differentially expressed genes between high-risk and low-risk patients related to the gene signature. (A) Enrichment plots of differentially expressed genes, **(B)** Networks regulated by risk score-associated genes in HCC constructed by Cytoscape. (Low degree to bright colors for map nodes color, low degree to small sizes for map nodes size, Low combined-score to small sizes for map edge size.)