

Figure S1 The differentially expressed genes (DEGs) and related signaling pathways based on TCGA data. FC, fold change; PPI, protein-protein interaction; TCGA, The Cancer Genome Atlas.

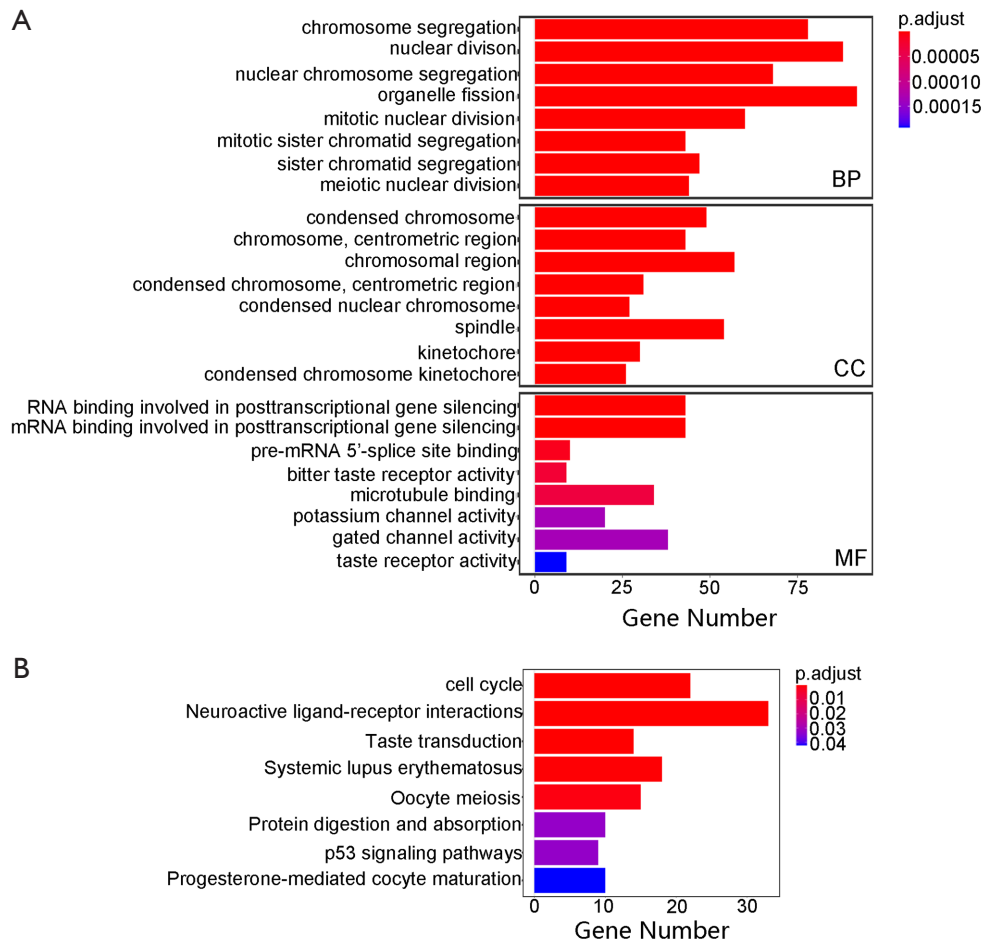


Figure S2 The correlation between TTK expression and tumor progression-associated signaling pathways was dissected through KEGG and GO analysis. KEGG, Kyoto Encyclopedia of Genes and Genome; GO, Gene Ontology. TTK, tyrosine threonine kinase; BP, Biological Process; CC, Cellular Component; MF, Molecular Function.