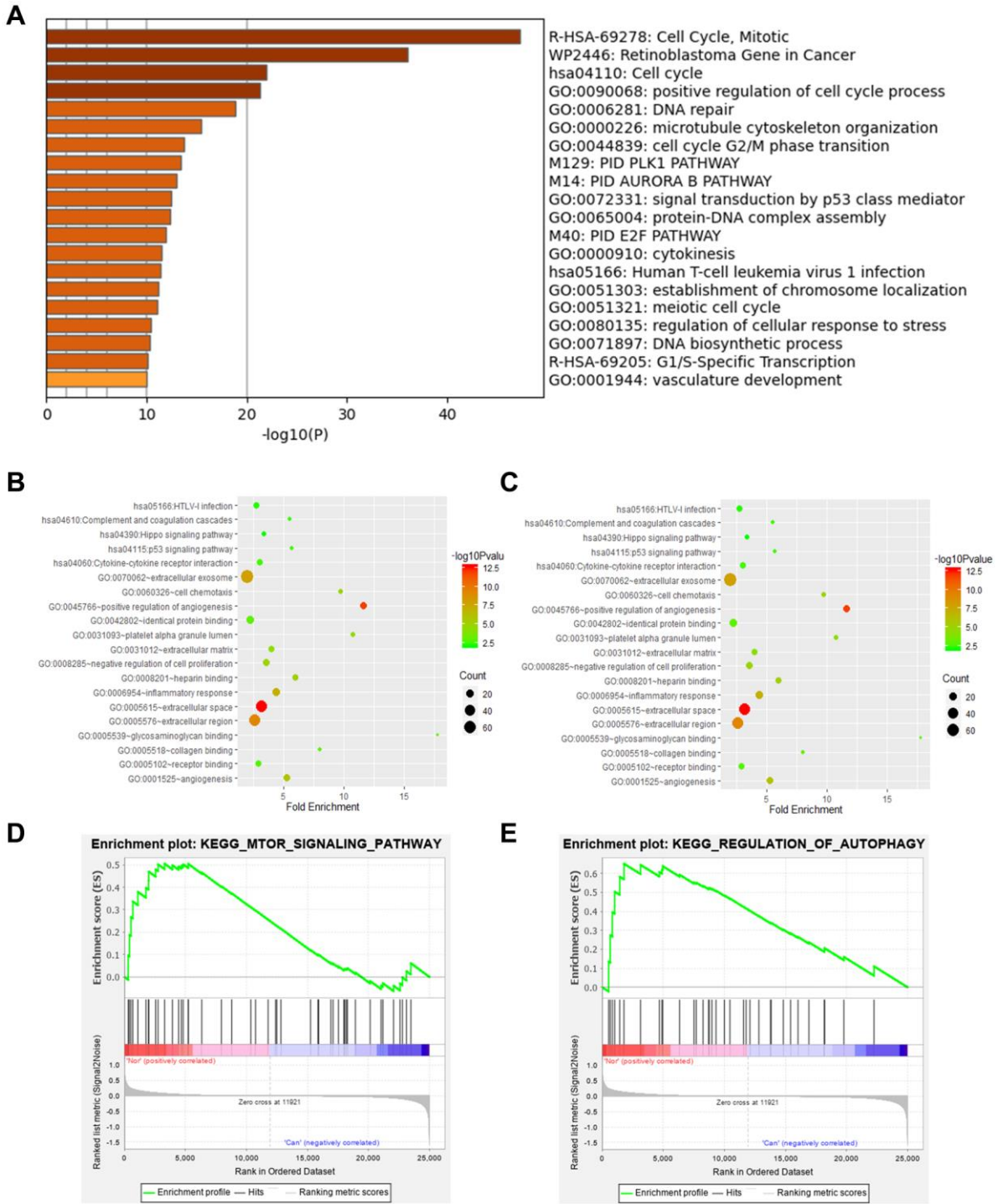
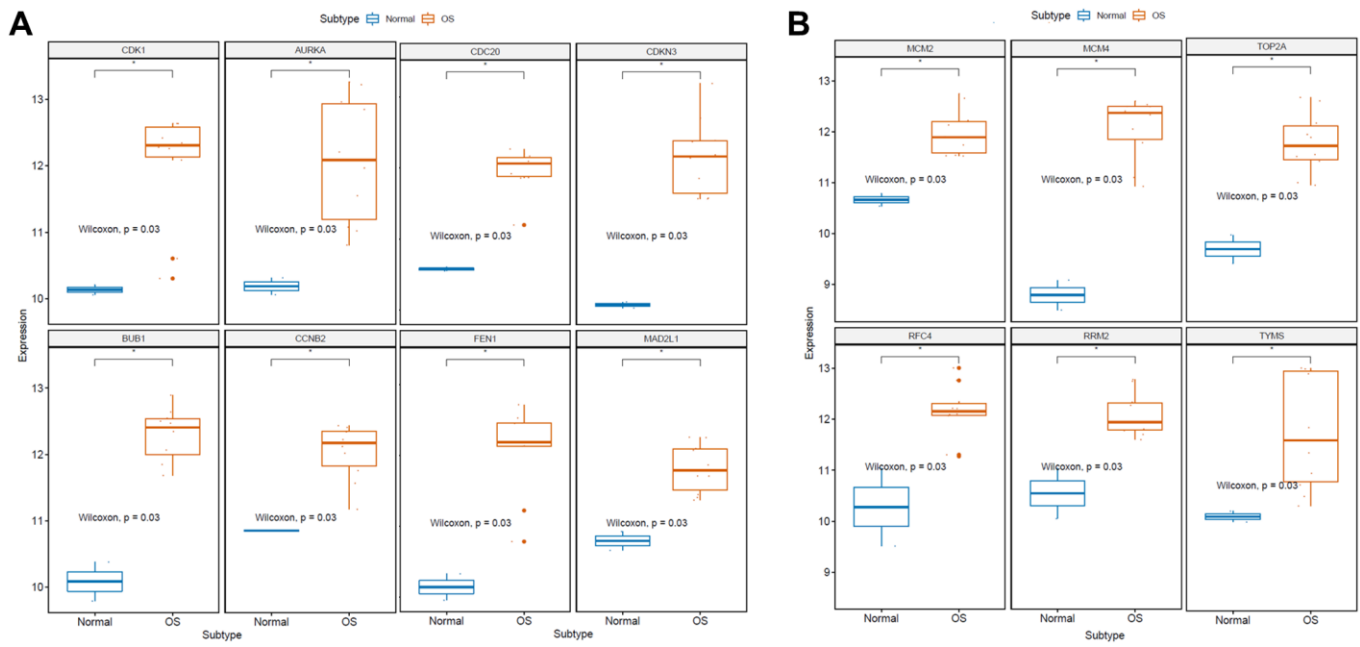


SUPPLEMENTARY FIGURES



Supplementary Figure 1. Functional and pathway enrichment analyses. (A) GO terms and enriched KEGG pathways of the DEGs. (B) Functional annotation and enrichment of the upregulated genes. (C) Functional annotation and enrichment of the downregulated genes. (D, E) GSEA results.



Supplementary Figure 2. The expression of hub genes in the GSE14359 dataset.