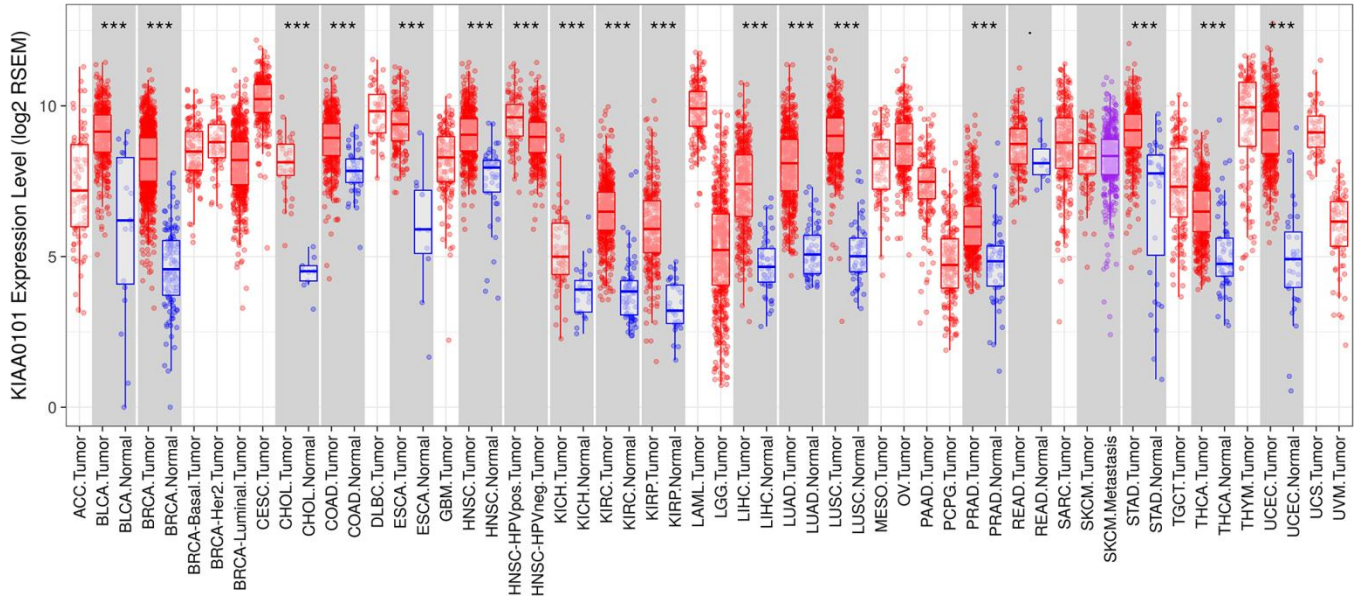
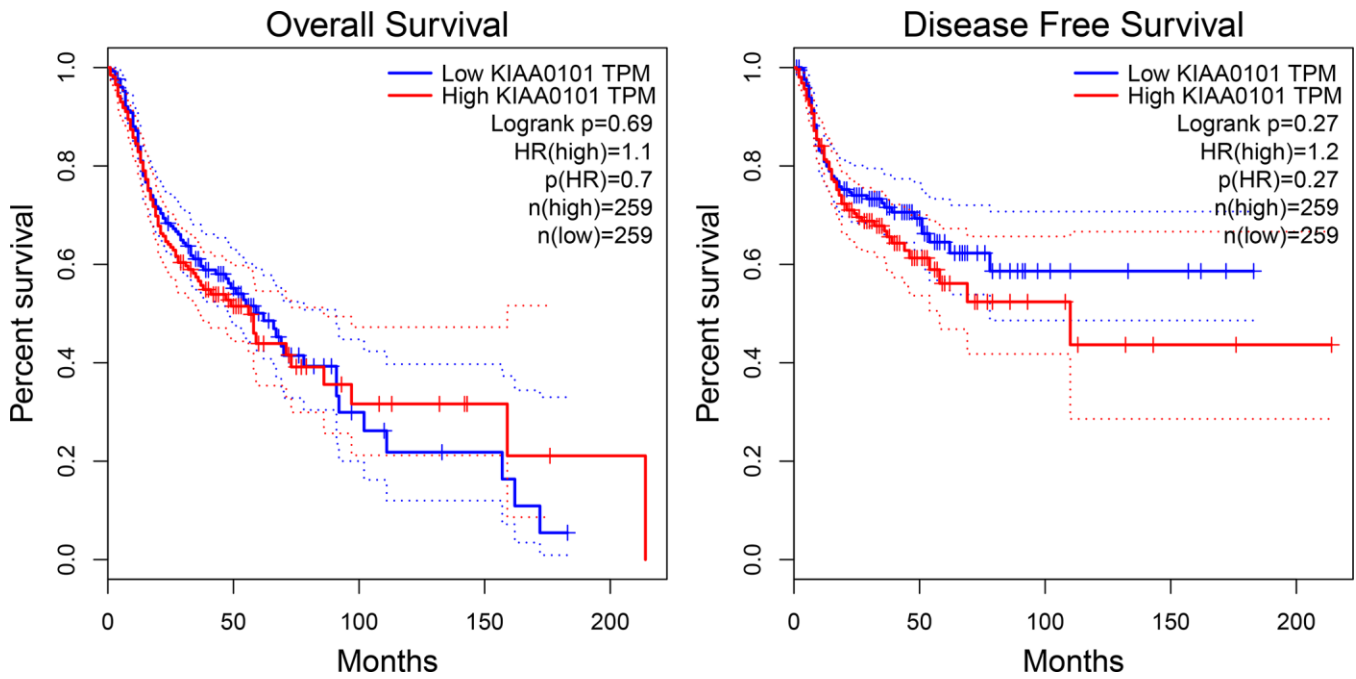


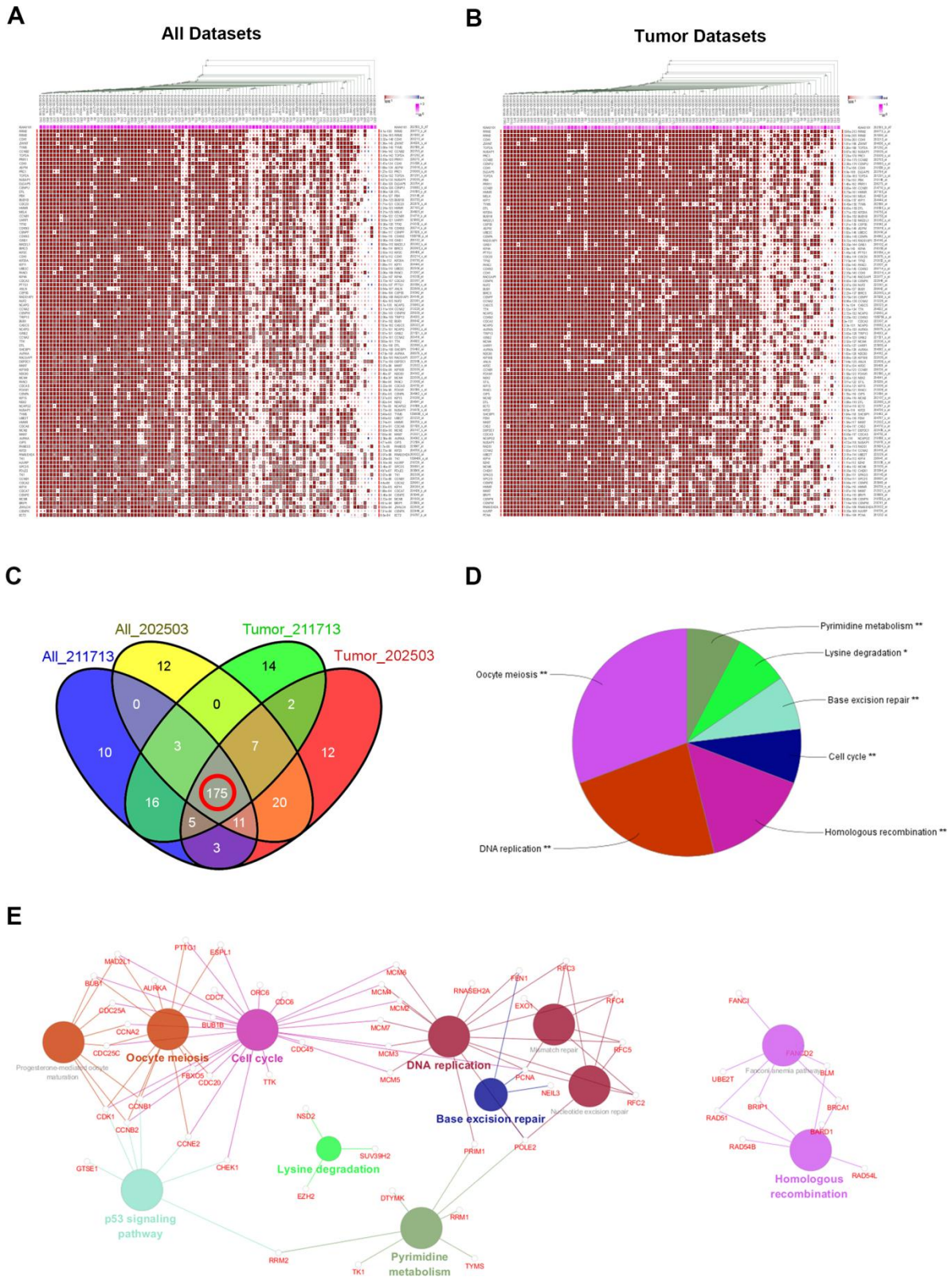
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Highly expressed KIAA0101 in several human cancer types.

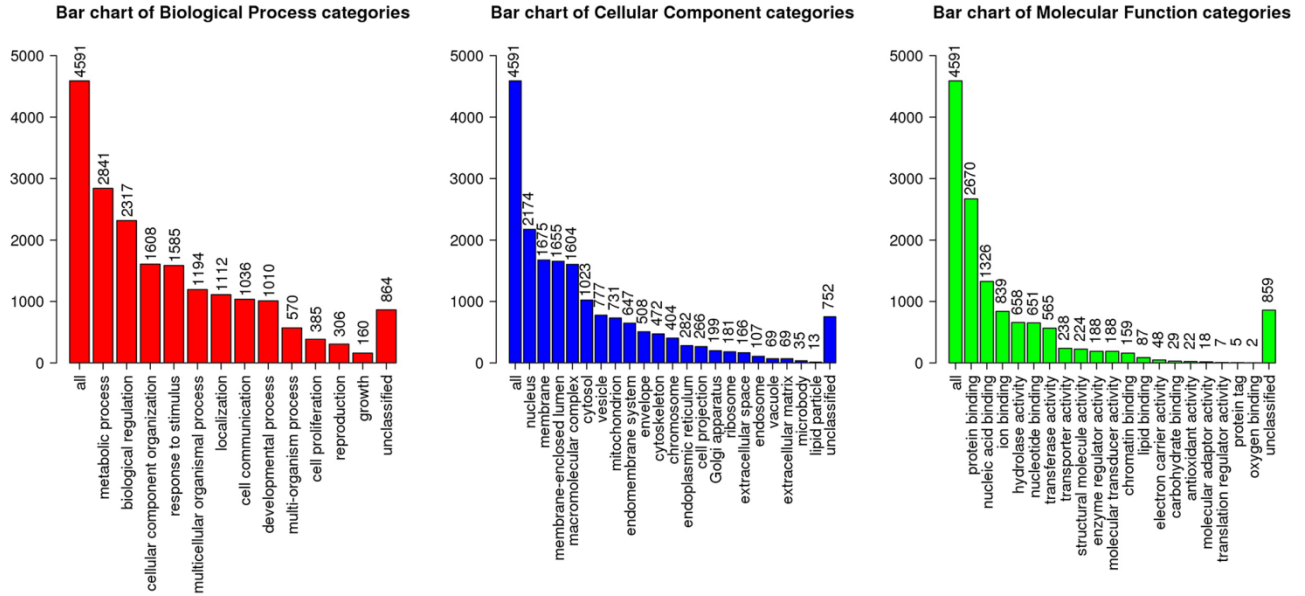


Supplementary Figure 2. Overall and disease-free survival analysis of KIAA0101 in TCGA HNSC samples.

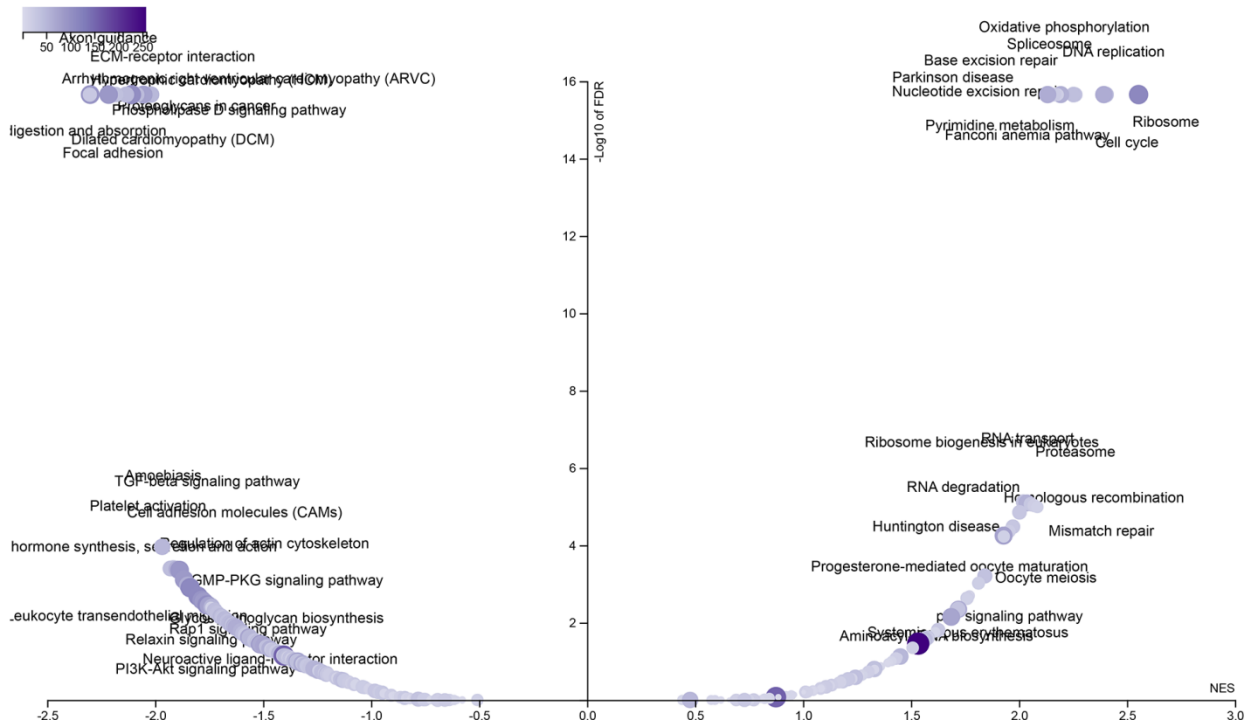


Supplementary Figure 3. MEM co-expression analysis of KIAA0101 reveals DNA replication and cell cycle pathways participate in tumor process. (A, B) Heatmap analysis of all and tumor datasets showing top 100 co-expressed genes with KIAA0101 probe 202503_S_AT. **(C)** Overlap venn diagram of co-expressed genes in all and tumor datasets queried with KIAA0101 probes 202503_S_AT and 211713_S_AT. **(D, E)** Pie chart and pathway enrichment analysis of 158 common genes with Cytoscape CluGO KEGG (17 same genes deleted).

A



B



Supplementary Figure 4. LinkedOmics GSEA analysis results of co-expression genes with positive or negative correlations with KIAA0101 in TCGA HNSC samples.