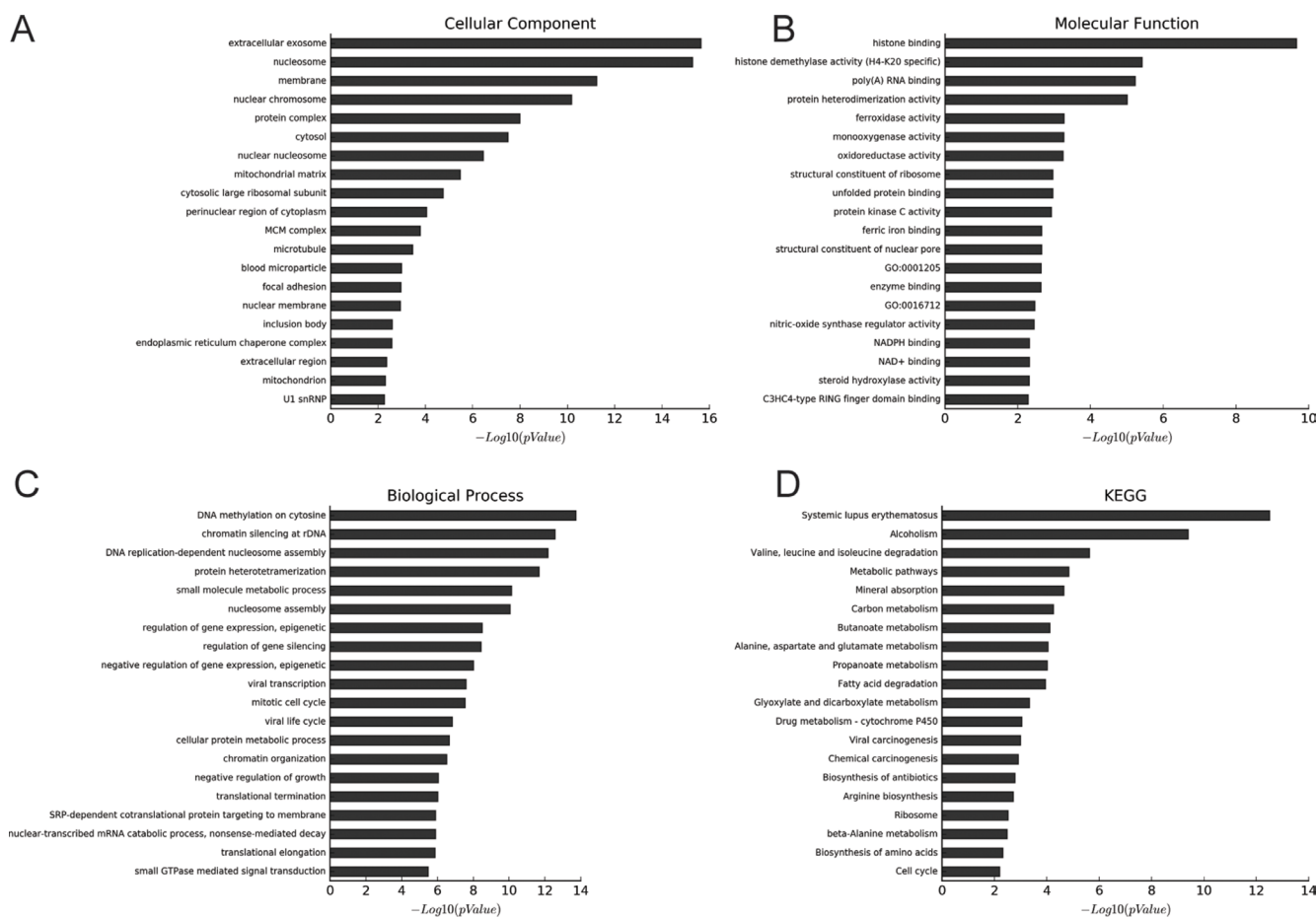
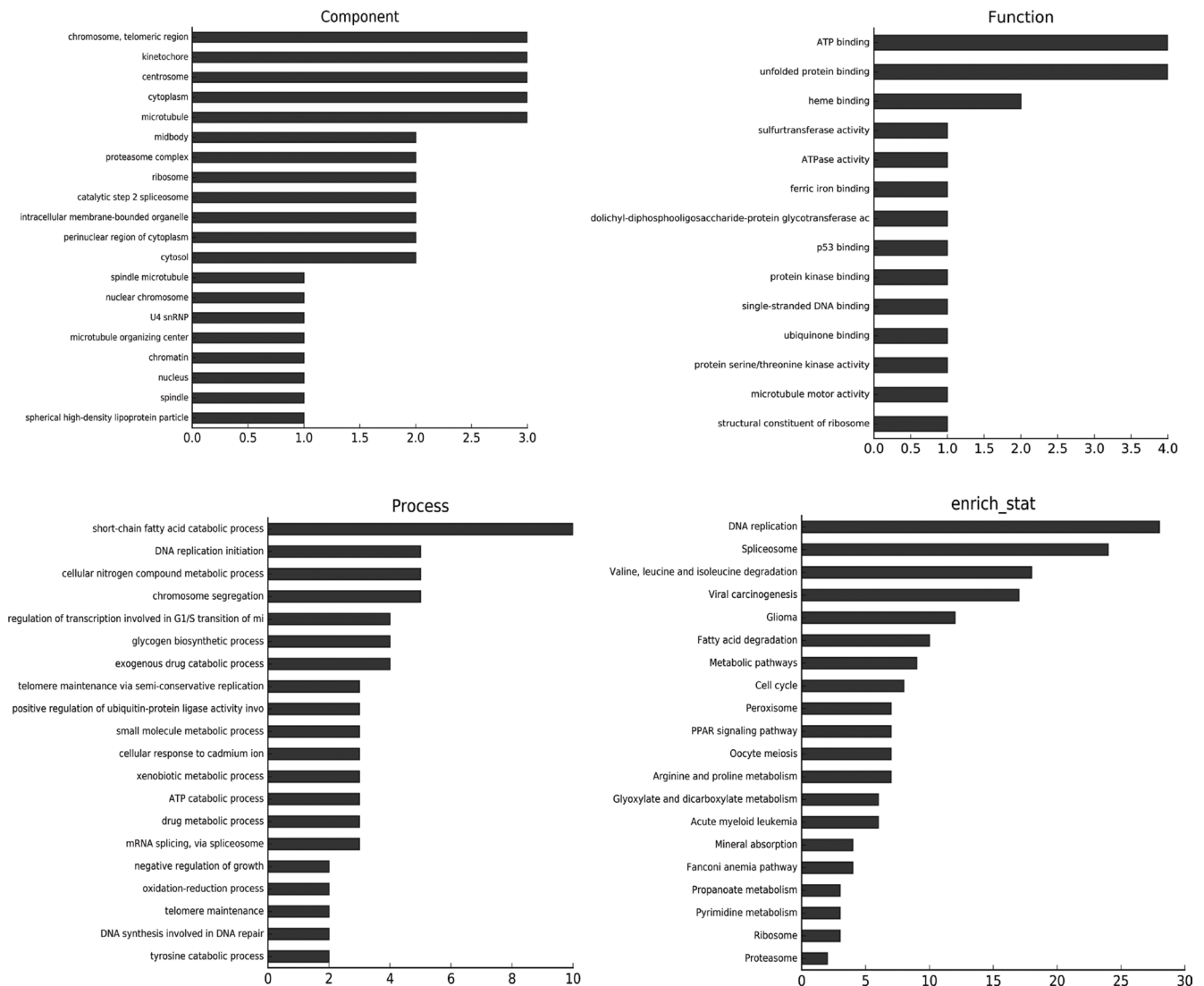


LncRNA-SVUGP2 suppresses progression of hepatocellular carcinoma

Supplementary Materials



Supplementary Figure 1: GO analysis and KEGG pathway analysis were performed to evaluate the functional categories and involved pathways of the dysregulated coding genes in HCC.



Supplementary Figure 2: GO analysis and KEGG pathway analysis for the top 200 co-expressed mRNAs of the lncRNA-SVUGP2.