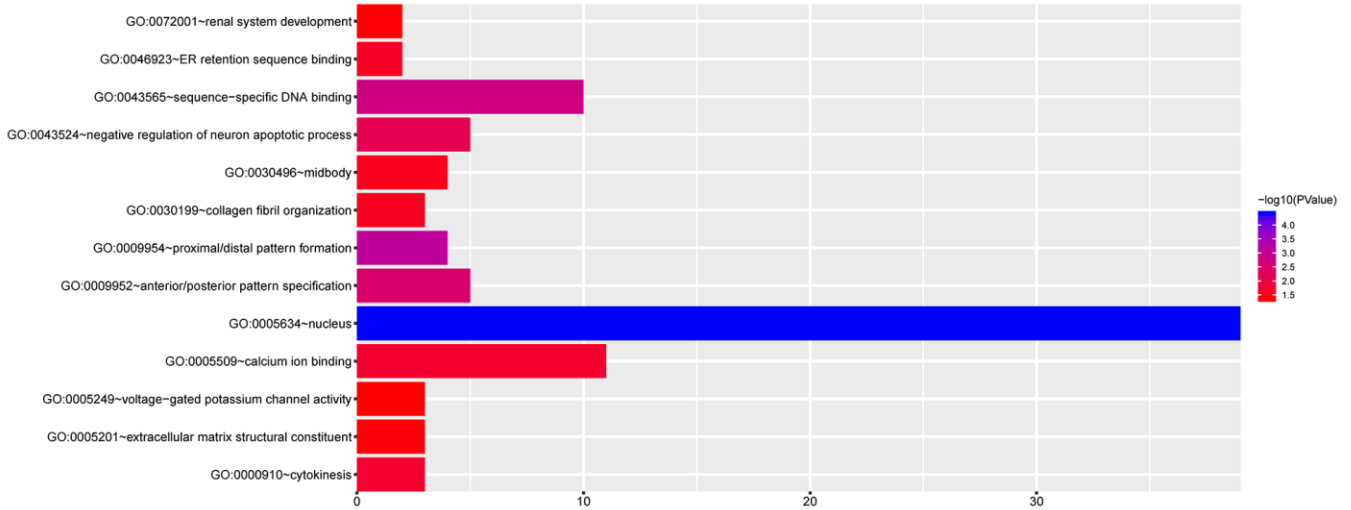
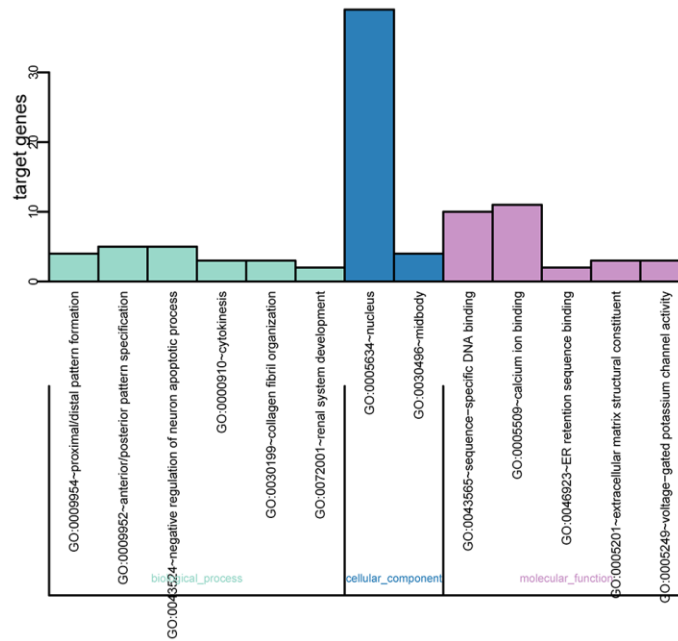


SUPPLEMENTARY FIGURES

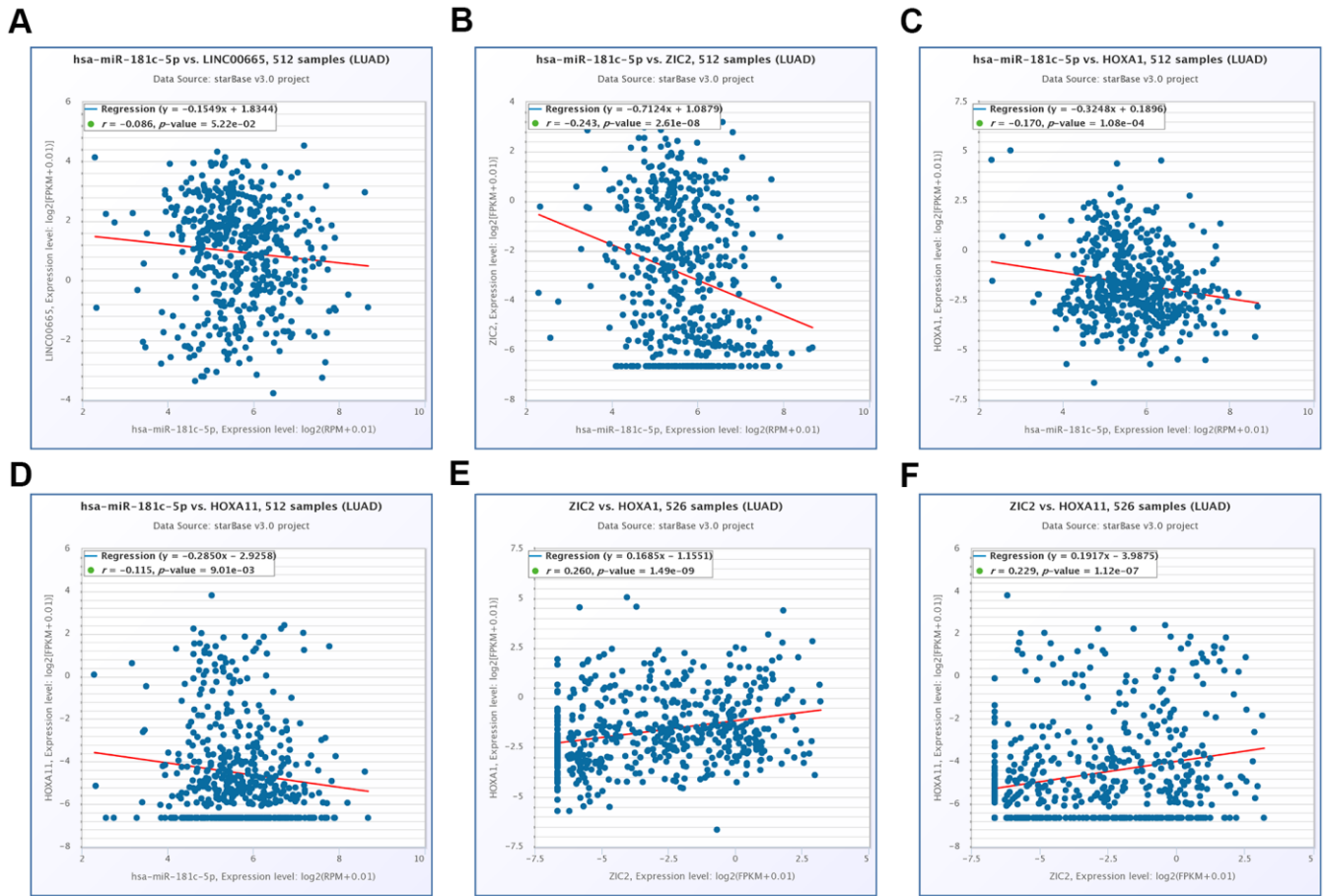
A



B



Supplementary Figure 1. GO analysis. (A, B) GO pathway analysis of the 151 genes which were upregulated in LUAD and also targeted by miR-181c-5p.



Supplementary Figure 2 Analysis of the correlation of the expression levels of LINC00665, ZIC2, HOXA1, HOXA11 in LUAD samples. (A–F) StarBase database was applied to analyze the correlations between the expression levels of miR-181c-5p and LINC00665, ZIC2, HOXA1, HOXA11, as well as ZIC2 and HOXA1/HOXA11.