

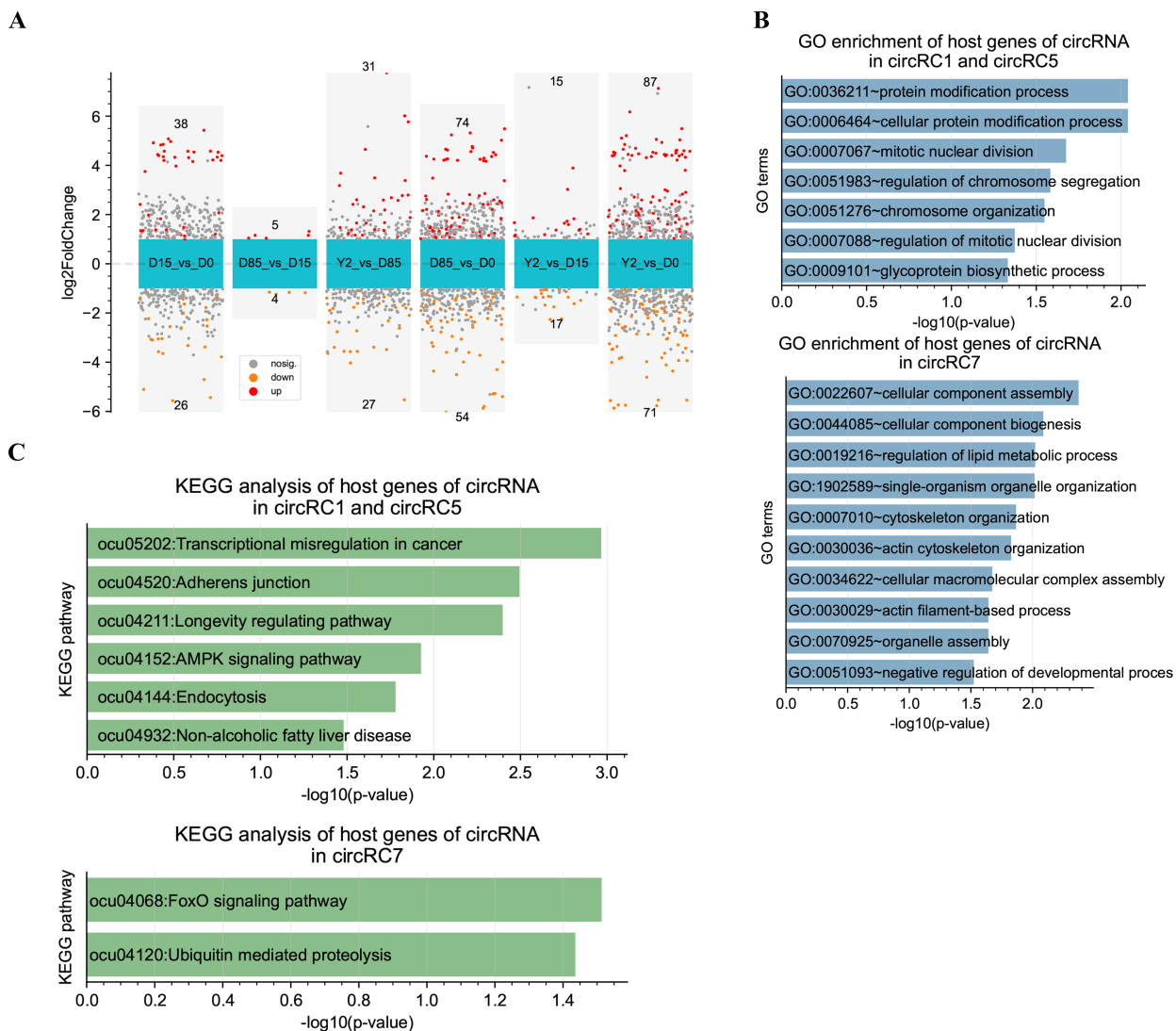


Figure S6. Differential analysis of circRNAs and host gene analysis. (A) The number of detected differentially expressed circRNAs (DECs) in different comparisons. The scatter plot showed circRNAs with $|\log_2(\text{fold-change})| > 1$ in the six comparisons. The red and orange points show the circRNAs with a $p\text{-value} < 0.05$. (B) GO-BP enrichment of host genes of DECs. (C) KEGG pathway analysis of host genes of DECs.