

Figure S1. Genome distribution map of four samples.

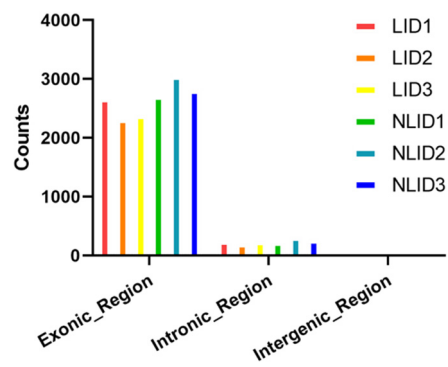


Figure S2. The classification of predicted circRNAs.

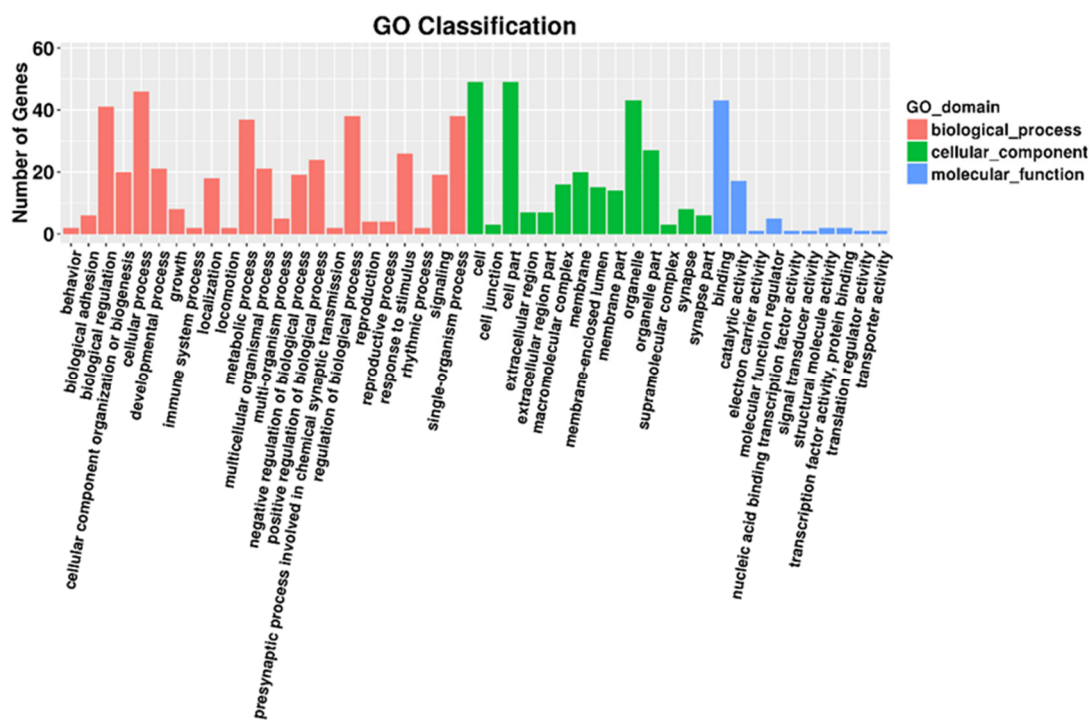


Figure S3. Go functional classification of the host genes of DEcircRNAs

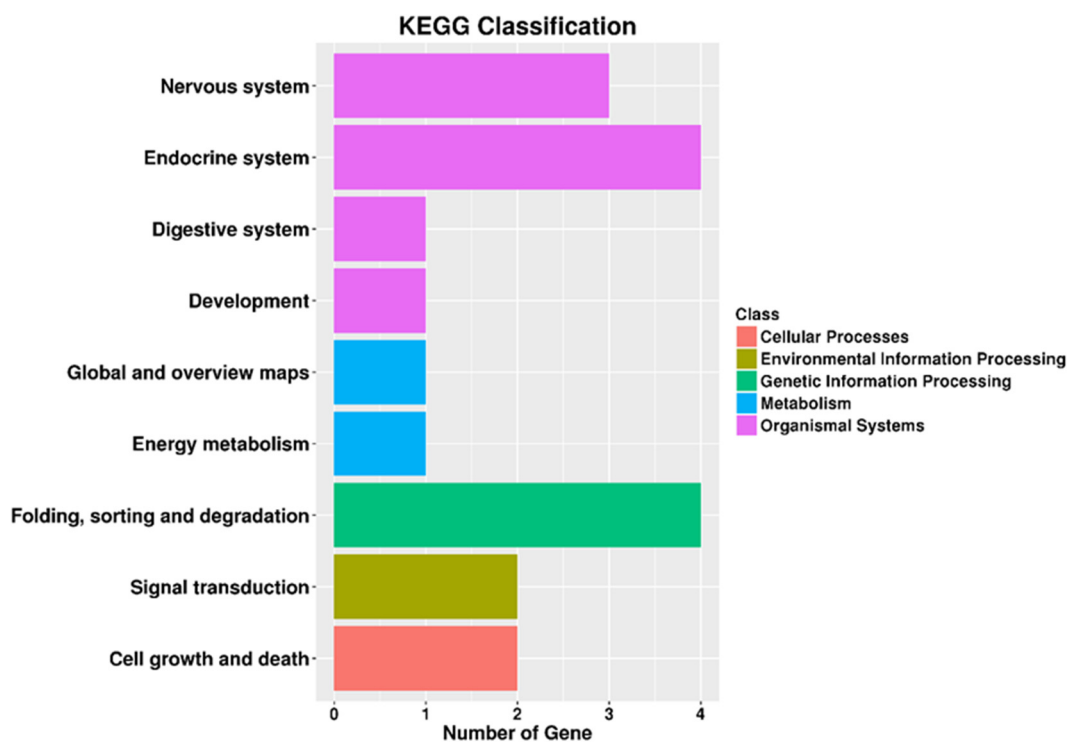


Figure S4. KEGG classification of the host genes of DEcircRNAs