Supplementary Fig.3 Validation the expression trends of the ten randomly selected lncRNAs (five upregulated and five downregulated at 75dpc point) by qRT-PCR.

Note: The GAPDH was used as reference genes, the fold change was calculated using the 2-△△Ct method. (Mean±SD，n=3，\*p < 0.05, \*\*p < 0.01).