

Table S1 The relative sequences used for the PCR

Targeted gene	Forward primer 5'-3'	Reverse primer 5'-3'
<i>AC122108.1</i>	GGGTCAGTAGCAGTGAACC	GTGGGCTCTGATGCAGAACT
<i>Si-AC122108.1</i>	GCCAAUAUGUCUCUAGAAATT	UUUCUAGAGACAUUUGGCTT
<i>ALDOA</i>	GAGGCGTCCATCAACCTCAA	GCAGCCTTCAGGTTCTCCTT
<i>GAPDH</i>	TGACTTCAACAGCGACACCCA	CACCTGTTGCTGTAGCCAAA

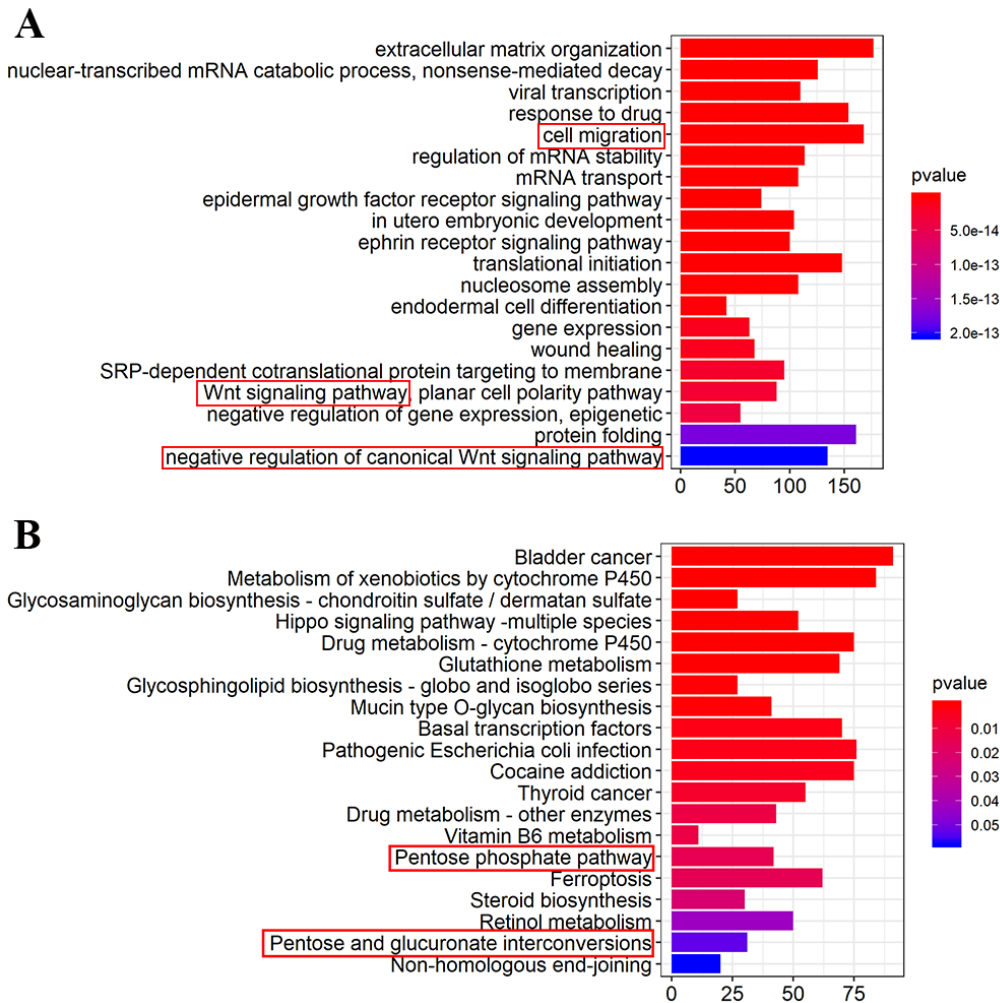


Figure S1 The transcriptional changes were displayed at the overall level by the GO and KEGG classification analysis. (A) The GO significant enrichment analysis showed that these differently expressed genes are mainly involved in the Wnt signaling pathway. (B) The KEGG significant enrichment analysis showed that these differently expressed genes are mainly involved in cancers and the pentose phosphate pathway.