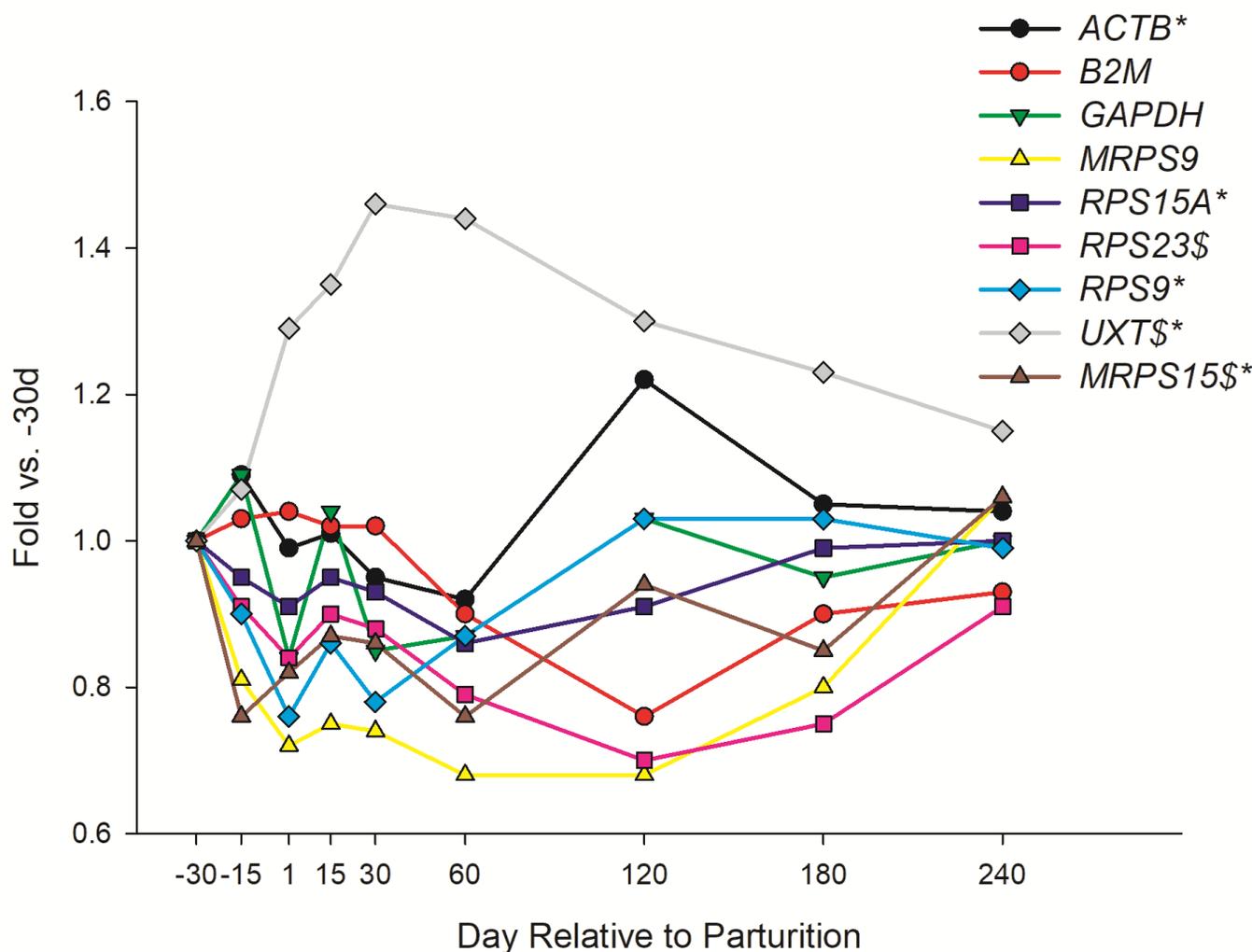




Supplementary Figure S1. Comparison of selected genes of Microarray (M: blue) and qPCR (Q: red) data in yak mammary gland during lactation.



Suppl. Figure 2. Evaluation of the transcription pattern as detected by microarray on previously assessed potential internal control genes for RTqPCR. \$ indicate prior reference genes used to normalize RTqPCR data from the same samples used in the present manuscript (Jiang et al., 2016).* indicate transcripts that were statistically affected with a $FDR < 0.05$ by day relative to parturition in yak mammary tissue in the current experiment (see Suppl. File 1)