



Fig. 1suppl. Correlation of microarray data with data obtained by qRT-PCR on 4 genes that displayed significant expression changes, showing a high correlation in the obtained transcriptional profiles. The y-axis is Z-normalized log2 expression (array) and 2-normalized QT value (qPCR). The x-axis shows biopsy file names. Spearman correlation coefficient was used for data comparison; DLX2: $r_s = 0.918$, $p = 0.00043$; DDX25: $r_s = 0.945$, $p = 0.00014$; MAGEA4: $r_s = 0.880$, $p = 0.000026$; EGR4: $r_s = 0.838$, $p = 0.0038$.