Table SII. Fold changes and statistical significances (P-values in brackets) of investigated comparisons with the Wilcoxon signed-rank test.

Comparison	*L1TD1*	*SLITRK6*	*ST6GALNAC1*	*TCEA3*
Central part vs. Invasive front	-2.34	1.12	-2.64 (P=0.041)[a]	-1.22 (P=0.035)[a]
Central part vs.lymph node metastasis	-52.95 (0.002)	-13.45 (0.001)	-12.82 (P<0.001)	-2.68 (P=0.003)
Central part vs.liver metastasis	-28.12	4.37	-2.18	1.06

[a]Comparison falls above the corrected borderline significance P=0.017 after additional Bonferroni correction for the investigated comparisons. ΔCq data obtained by normalization to the geometric mean of *B2M* and *IPO8* reference genes, $\Delta\Delta$Cq data obtained by normalization to central part of primary tumor. Fold changes calculated as $2^{-\Delta\Delta Cq}$.

Table SIII. Fold changes and statistical significances (P-values in brackets)of investigated comparisons with the Wilcoxon signed-rank test for the investigated miRNAs.

Comparison	miR-199a-3p	miR-425-5p	miR-1225-3p	miR-1233-3p	miR-1303
Central part vs.invasive front	1.73 (0.040)[a]	-1.88	-1.24	1.09	1.95
Central part vs.lymph node metastasis	46.00 (P <0.001)	26.01 (P <0.001)	17.15 (0.004)	16.84 (P <0.001)	5.83 (0.007)
Central part vs.liver metastasis	75.43 (0.002)	1520.56 (0.004)	1.63	-3.16 (0.006)	-1.60

[a]Comparison falls above the corrected borderline significance P=0.017 after additional Bonferroni correction for the investigated comparisons. miRNA/miR, microRNA. ΔCq data obtained by normalization to the geometric mean of RNU6B and miR-1274b reference genes, $\Delta\Delta$Cq data obtained by normalization to central part of primary tumor. Fold changes calculated as $2^{-\Delta\Delta Cq}$.