

Supplementary Fig. 1. Expression analysis of EMT-associated genes and microRNAs in the invasion front of colorectal liver metastases. Average, relative fold-change expression of A, Cadherin-11, B, Collagen 1A1, C, Collagen 3A1, D, vimentin, E, matrix-metalloproteinase 3, F, hypoxia-inducible factor 1A, G zinc finger E-box binding homeobox 1, H, zinc finger E-box binding homeobox 2, I, fibroblast-specific protein, J, microRNA 141, K, microRNA 200a, L, microRNA 200b, M, microRNA 200c. Average, relative expression fold-change expression in liver (L), the liver invasion front (LI) and the tumor invasion front (TI) compared to the tumor center (T), respectively. Means  $\pm$  SD are shown. P-values were evaluated using the Wilcoxon signed rank test.