



Supplementary Figure 1: TRAILR2 and CDH17 RNA expression in related normal tissues. mRNA-expression (TPM, RNAseq) levels for TRAILR2 (y-axis) and CDH17 (x-axis) in non-diseased colon, esophagus, pancreas and stomach from the Genotype-Tissue Expression (GTEx) project database. The number of samples included in the analysis for each group is indicated in parenthesis. In these normal tissues, medium to low number of samples showed high levels of expression of both CDH17 and TRAILR2. This was mainly driven by the reduced expression of CDH17. In colon, 221 out of 539 samples (41%) showed levels >10TPM for both targets. In stomach 1 out of 272 (0.4%), in esophagus 2 out of 1101 samples (0.2%), and in pancreas none of the 268 samples available showed levels >10TPM for both targets.