



Supplementary Figure S2: Expression-levels of candidate biomarkers and pair-wise correlation of underlying samples. Plot A shows the absolute expression of candidate biomarkers for each sample. Plot B shows the correlation of the samples to identify outliers or incoherent clustering. The expression-correlation of the samples shows a clustering according to malignant and benign subtype, respectively, however, the expression levels of the biomarker candidates between and within groups vary and show considerable heterogeneity as shown in Plot A.