



**Figure S3: Flow chart depicting the generation of the EMT-core gene list.** Overview of the meta-analysis, the generation of the 365 genes list, the EMT-core list as well as the single enrichment analysis and the correlation with clinical data. GEO, Gene expression Omnibus; AE, Array Express; EMT, epithelial to mesenchymal transition; FC, Fold change; Up, upregulated; Down, downregulated.