



Supplementary Figure 3. EMT gene expression is enriched in pre-clinical models highly expressing AKT3 across multiple carcinoma cell lines. (a) GSEA on PDX models. Cut-off for AKT3 high and low expression was set to 2Log expression >3.5 and <1, respectively; (b) GSEA on 50% high versus 50% low AKT3 expressing carcinoma cell lines included in the CCLE panel, separated per cancer type.