

## **Description of Additional Supplementary Files**

File Name: Supplementary Data 1

Description: An Excel Sheet of the microRNA (miRNASeq) and mRNA (RNASeqv2) pre-processed source data that were downloaded from the Cancer Genome Atlas Project (TCGA) publicly available at Broad Institute (<http://gdac.broadinstitute.org/>) for patients with lung adenocarcinoma (LUAD). The clinical information in the Excel Sheet was downloaded from cBioPortal (<http://www.cbioportal.org/>). Data was processed as explained in the “Methods” section of the manuscript (page # 38; TCGA) and utilized to generate the graphs in Figure 6c,d and Supplementary Figure 5.