**Supplementary Table 1** mRNA differentially expressed in tumors of 30 NSCLC patients with low muscularity compared with the gene expression of 59 NSCLC patients with high muscularity.





The transcriptomic data from “NSCLC-Radiomics-Genomics” TCIA collection were accessed using GEO (GSE58661). RefSeq: NCBI reference sequence; TCIA: The Cancer Imaging Archive; GEO: Gene Expression Omnibus.