

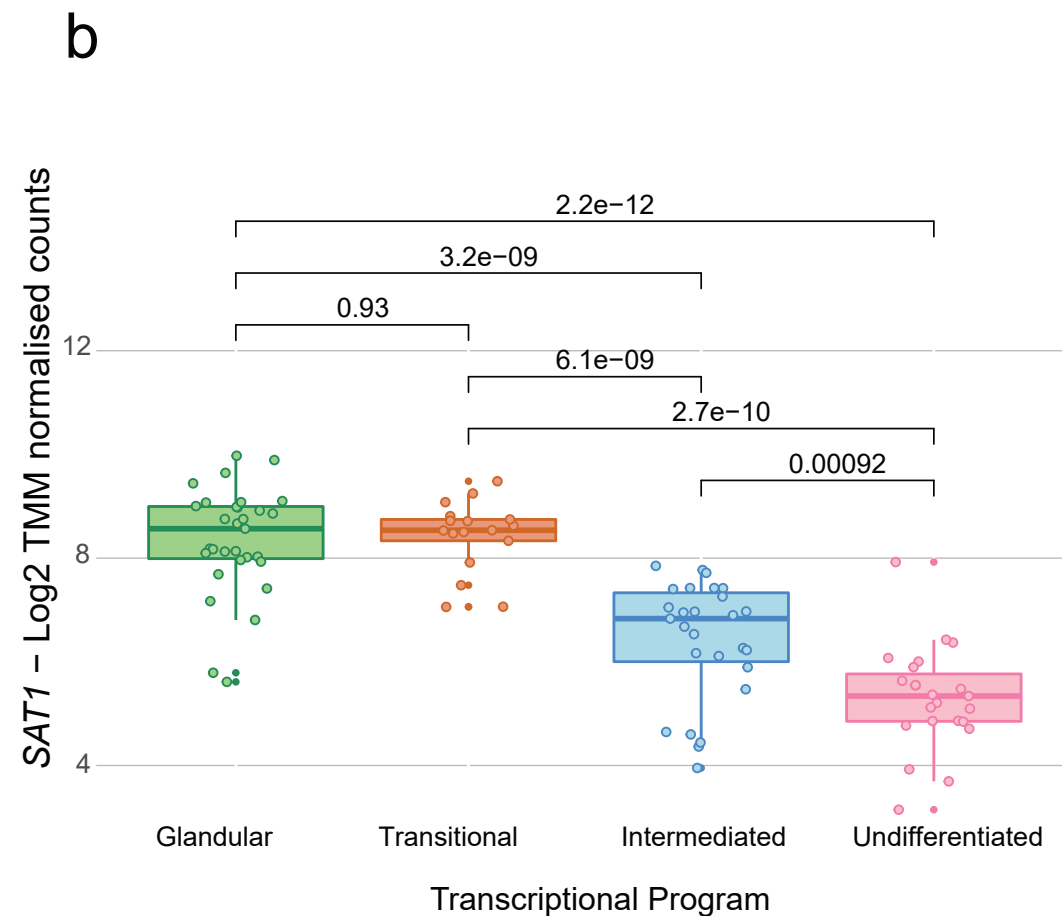
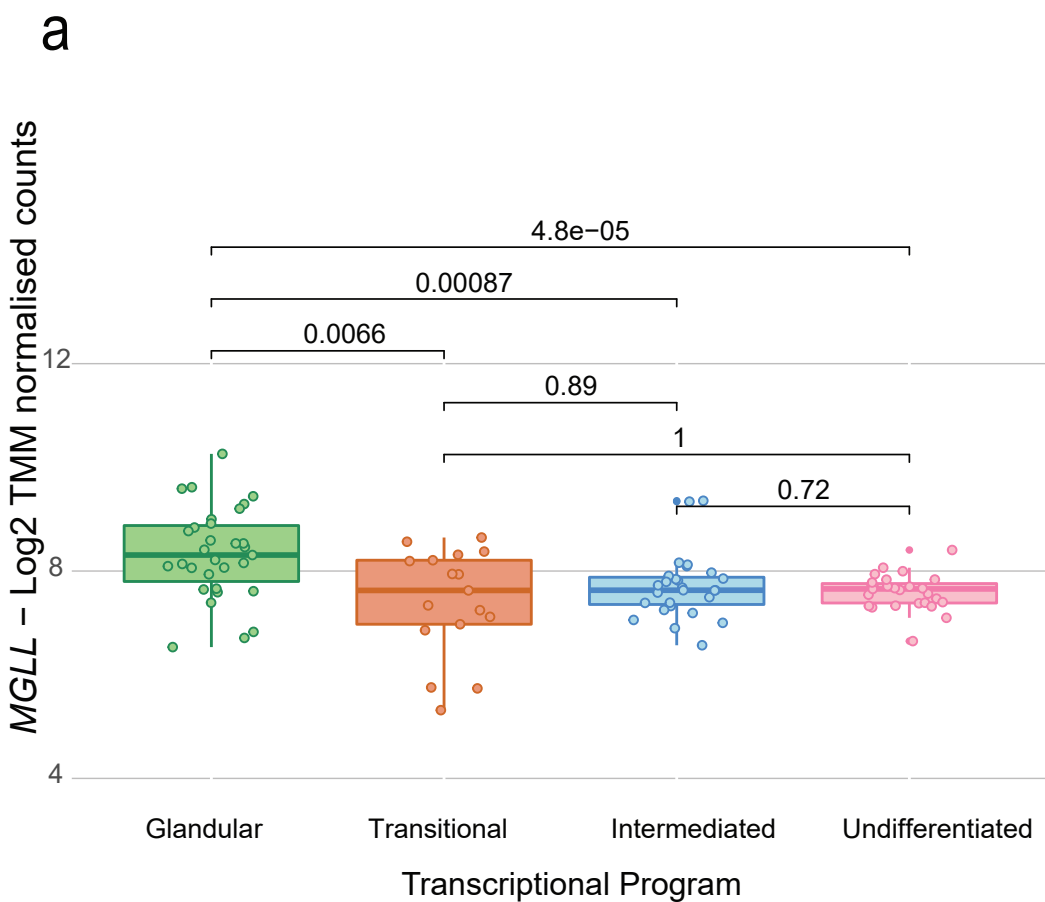


Figure S3. *MGLL* and *SAT1* expression in laser macro-dissected PDAC cancer cells RNASeq data(21). Raw counts were downloaded from Gene Expression Omnibus (GSE208732) and TMM normalised. a) *MGLL* gene expression in cancer regions enriched for each of the four transcriptional programs described by Di Chiaro et al (21). b) *SAT1* gene expression across the four transcriptional programs. *MGLL* and *SAT1* gene expression was higher in cancer cells with "*Granular*" and "*Transitional*" transcriptional programs, which were showed to be enriched in the PNI TME(21). This data support our findings that *MGLL* and *SAT1* was up-regulated in the cancer compartment of PNI compared to non-PNI foci.