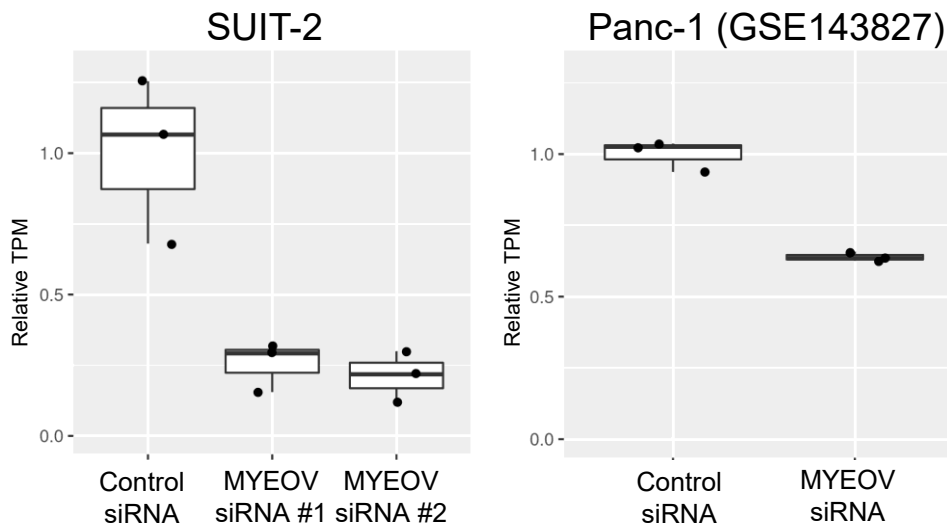


A



B

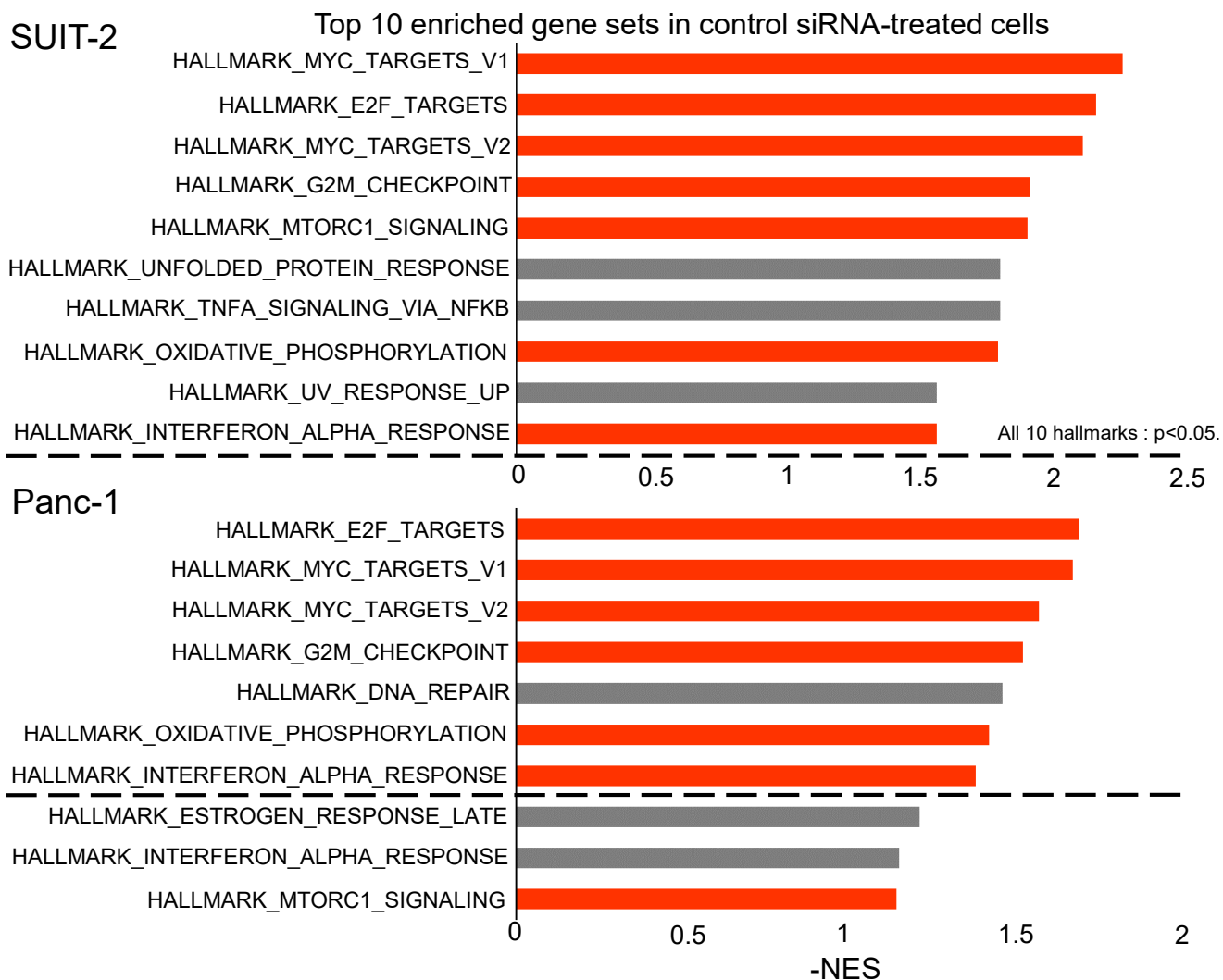


Fig. S4 The comparison of two RNA-Seq data in *MYEOV*-knockdown pancreatic cancer cells. A, Knockdown efficiency of *MYEOV* in two RNA-Seq datasets. Relative values based on TPM are presented. B, GSEA analysis was performed on the two indicated datasets. The bar graph shows top 10 enriched gene sets in control siRNA-treated SUIT-2 (upper panel) and Panc-1 (lower panel) cells. Gene sets enriched in both datasets were presented as red bars. In lower panel, hallmarks above the horizontal dashed line indicate statistically significant enrichment (nominal p value < 0.05) in Panc-1 cells.