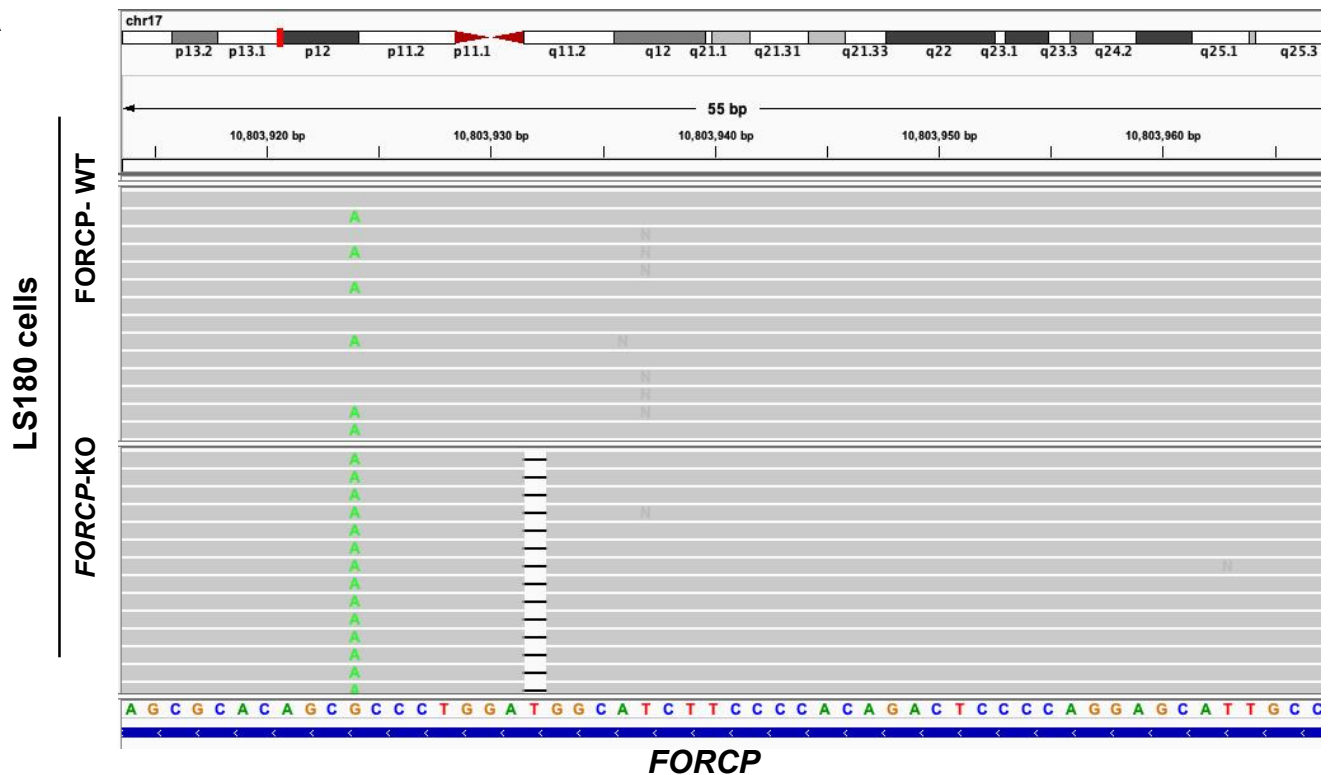
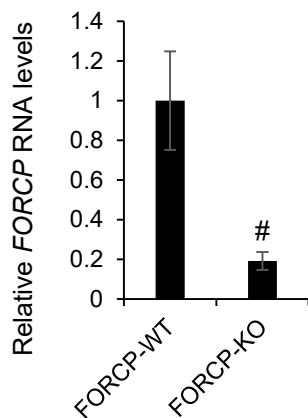
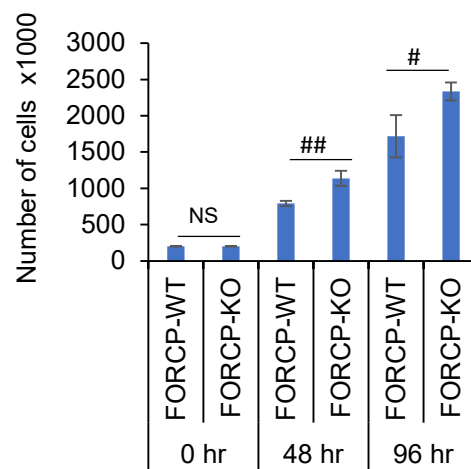
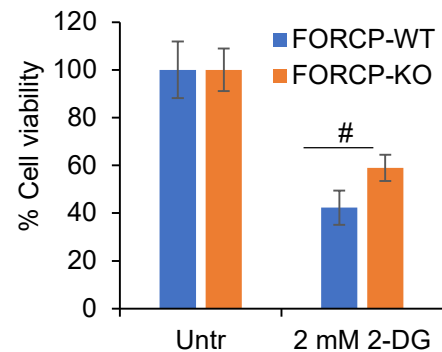


A**B****C****D****Figure 4-figure supplement 3.**

(A) IGV visualization of targeted amplicon sequencing data from *FORCP* wild type (WT) and isogenic *FORCP*-KO (knockout) LS180 cells. In the *FORCP*-KO cells, a 1 bp deletion is observed in one of the alleles within the sgRNA target site while it is likely a large deletion was generated in the other allele which had removed the PCR primer binding sites. **(B)** qRT-qPCR was performed from *FORCP*-WT and *FORCP*-KO LS180 cells. *GAPDH* was to normalize the data. # $p < 0.005$, ## $p < 0.001$. **(C)** *FORCP*-WT cells and *FORCP*-KO cells were seeded in 6-well plates and trypan blue exclusion cell count assays were performed at the time of seeding (0 hr) and after growing the cells for 48 and 96 hr, respectively. **(D)** Cell viability assays were performed from *FORCP*-WT and *FORCP*-KO LS180 cells that were left untreated or treated with 2-DG to inhibit glycolysis. Error bars represent SD from 3 experiments. # $p < 0.01$, ## $p < 0.001$.