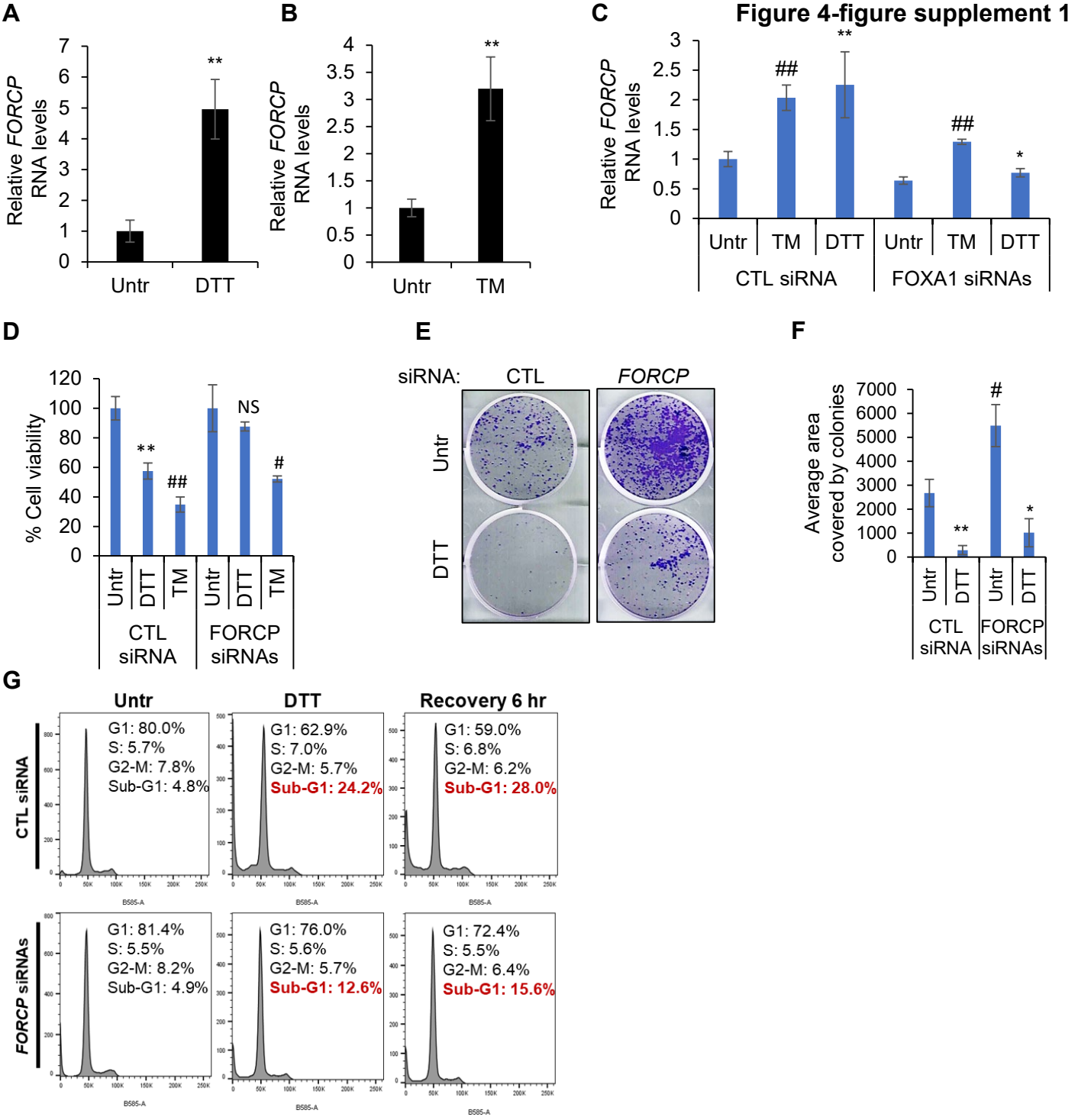

Figure 4-figure supplement 1.

(A, B) qRT-PCR was performed from LS180 cells after induction of ER stress using 0.5 mM DTT (A) or 4 ug/ml TM (B) to determine the effect of ER stress on *FORCP* RNA levels. Data was normalized with *GAPDH*.

(C) The effect of FOXA1 knockdown on *FORCP* induction upon ER stress was determined by qRT-PCR after transfection of LS180 cells with CTL siRNA or FOXA1 siRNAs for 48 hr followed by treatment with DTT for 2 hr.

(D) Cell viability assays were performed from LS180 cells transfected with CTL siRNA or *FORCP* siRNAs under untreated (Untr), DTT treated or Tunicamycin (TM) treated conditions. Cell viability was measured using CCK-8 assay. P-values were calculated by comparing DTT or TM treated samples with untreated (Utr).

(E, F) Colony formation on plastic assays were performed from LS180 cells following *FORCP* knockdown and treatment with ER-stress inducing agents (DTT or TM). Quantification of colonies was performed using ImageJ software. **(G)** PI staining and FACS analysis was performed from SW1222 cells transfected for 48 hr with CTL siRNA or *FORCP* siRNAs and then left untreated or treated with DTT for 2 hr or after 6 hr recovery from DTT treatment. Error bars represent SD from 3 experiments. *p<0.05, #p<0.01, **p<0.005, ##p<0.001.