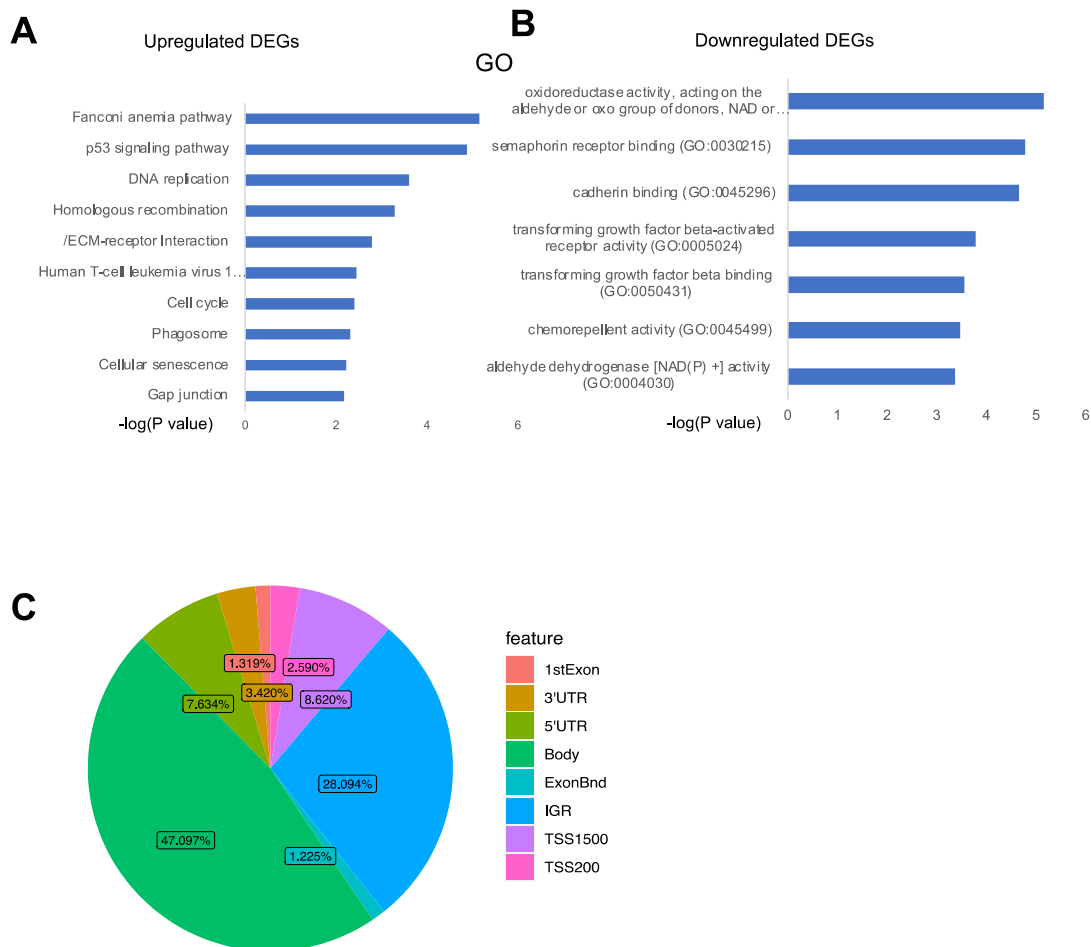


Supplemental Material



Supplemental Figure S2. NTX301 induced transcriptome and methylome reprogramming in OC cells. (A,B) Gene Ontology (GO) analyses of upregulated (A) and downregulated (B) DEGs show top significantly enriched pathways in OVCAR5 cells treated with NTX301 (100nM) vs. DMSO (n=3 replicates per group). (C) The distribution of DMPs in response to treatment with NTX301 across the genome.