

Supporting Information

Alteration of chromatin high-order conformation associated with oxaliplatin resistance acquisition in colorectal cancer cells

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Supplementary figures

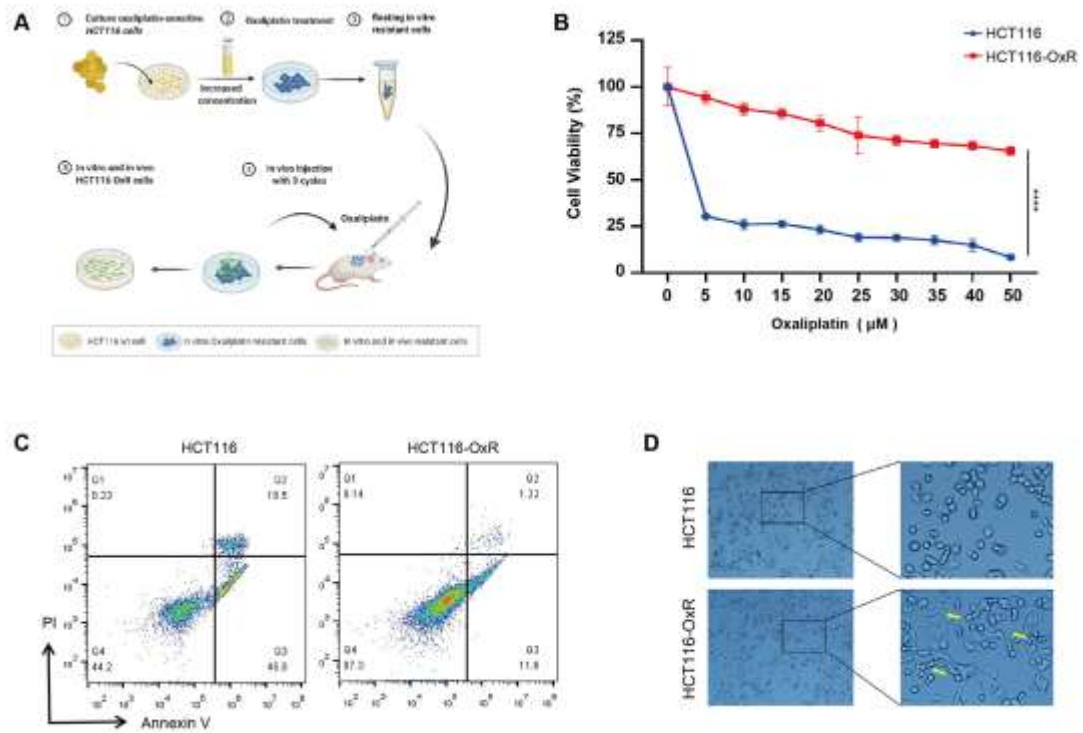


Figure S1. Establishment and validation of oxaliplatin resistant HCT116 cells.

A Construct oxaliplatin-resistant HCT116-OxR cell using concentration gradient method and nude mouse injection method. **B** HCT116-OxR was significantly insensitive to oxaliplatin by CCK8 assay, *** $P < 0.001$. **C** The percentages of live cells (Annexin V⁻/PI⁻), and apoptotic cells (Annexin V⁺/PI⁻ & Annexin V⁺/PI⁺) was evaluated among the HCT116 and HCT116-OxR cells pre-treated with oxaliplatin (100 μM , 24 h). **D** Compared with HCT116, the morphology of HCT116-OxR cells became fusiform and the polarity was lost.

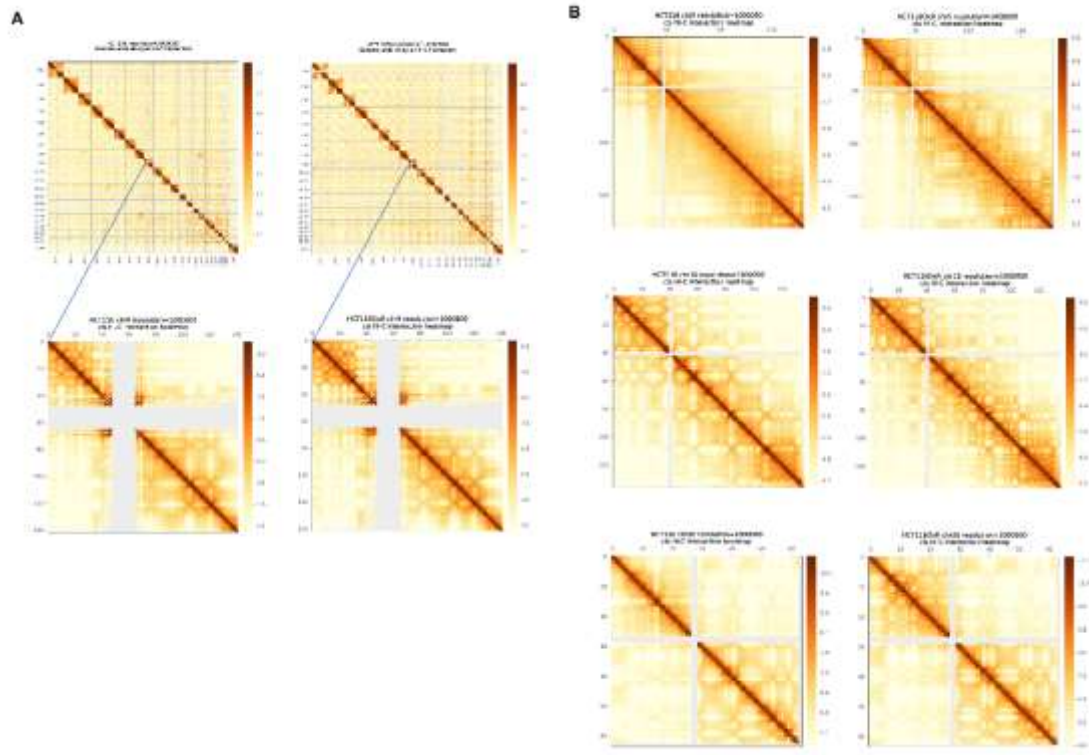


Figure S2. The repeatability chromatin remodeling and merged interaction heatmap in HCT116 and HCT116-OxR cells.

A Whole genome interaction maps show that intrachromosomal interactions are more frequent. **B** The heat map of the interaction of multiple chromosomes between the two groups of cells showed no significant difference.

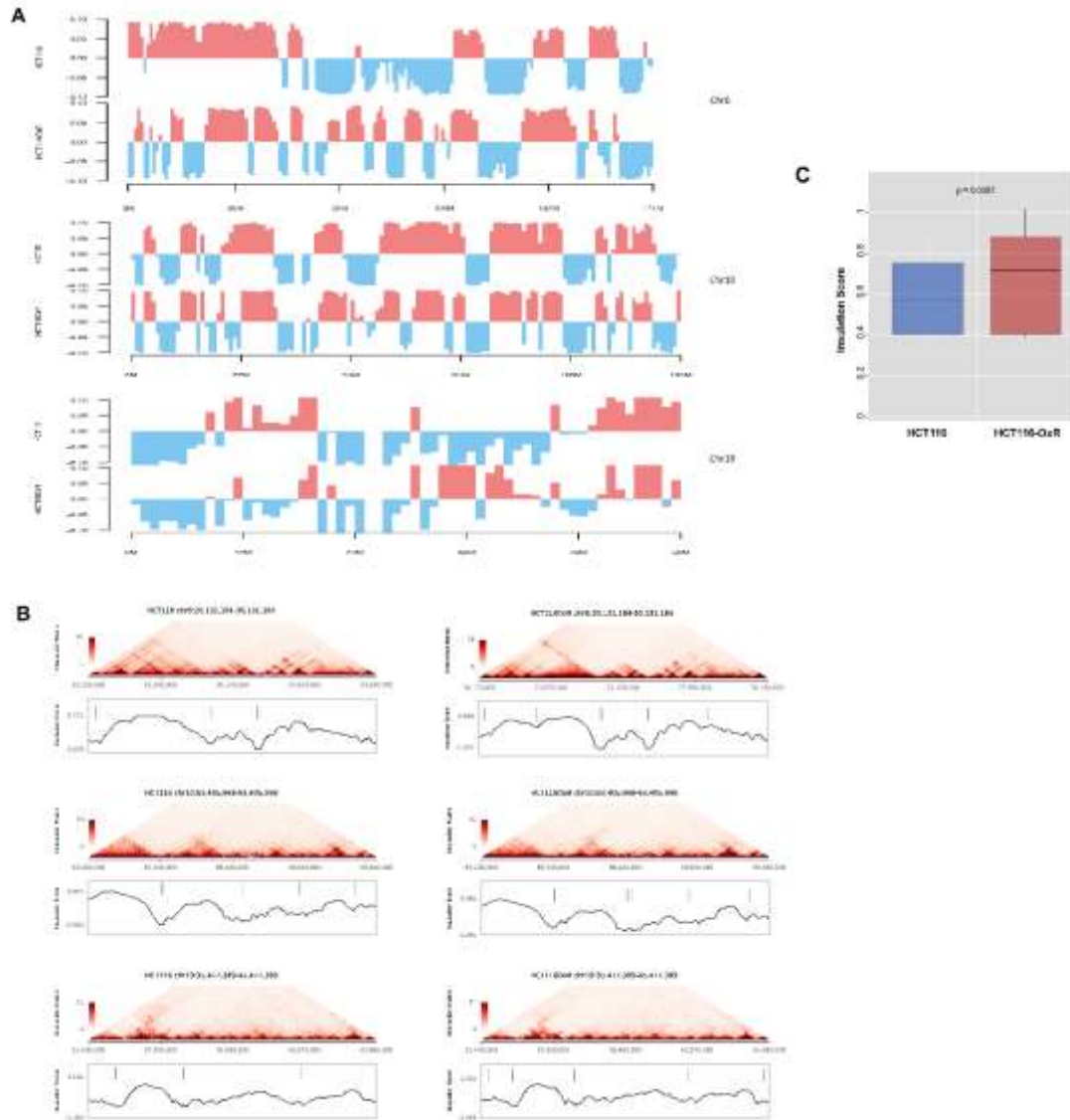


Figure S3. The difference between the A/B area and TAD structure of the two groups of cells.

A In the process of drug resistance, there are more genes in multiple chromosomes with A/B compartment transformation. **B** The structure of TAD is highly conservative during the occurrence of drug resistance. **C** The insulation score comparison between HCT116 and HCT116-OxR cells surrounding the gene PRDX6 at chr 1: 167,800,000–182,840,000. The sliding window was defined as 40 kb, and regions with low score were insulating.

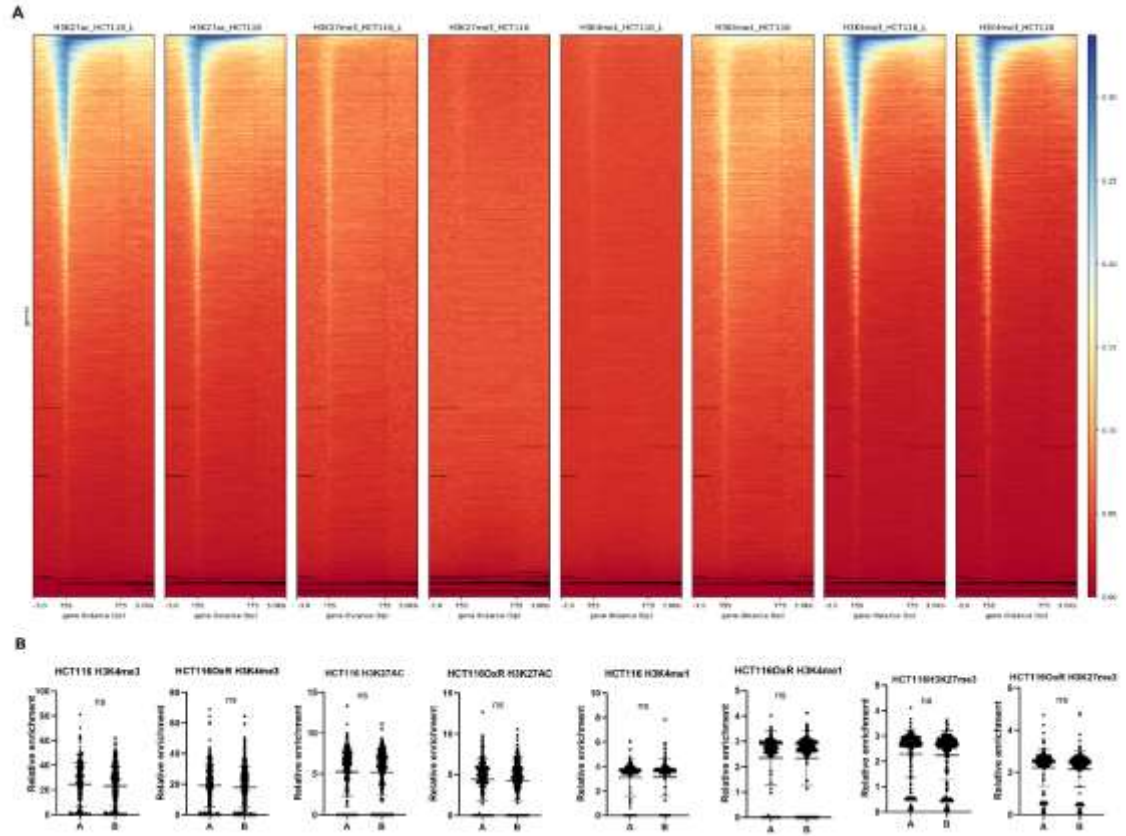


Figure S4. Modification of H3K27ac, H3K4me1, H3K4me3 and H3K27me3 in HCT116 and HCT116-OxR cells.

A H3K27ac, H3K4me1, H3K4me3 and H3K27me3 modified gene distribution in two cells. **B** The relative enrichment of indicative histones in A or B compartments (%) in HCT116 and HCT116OxR cells.

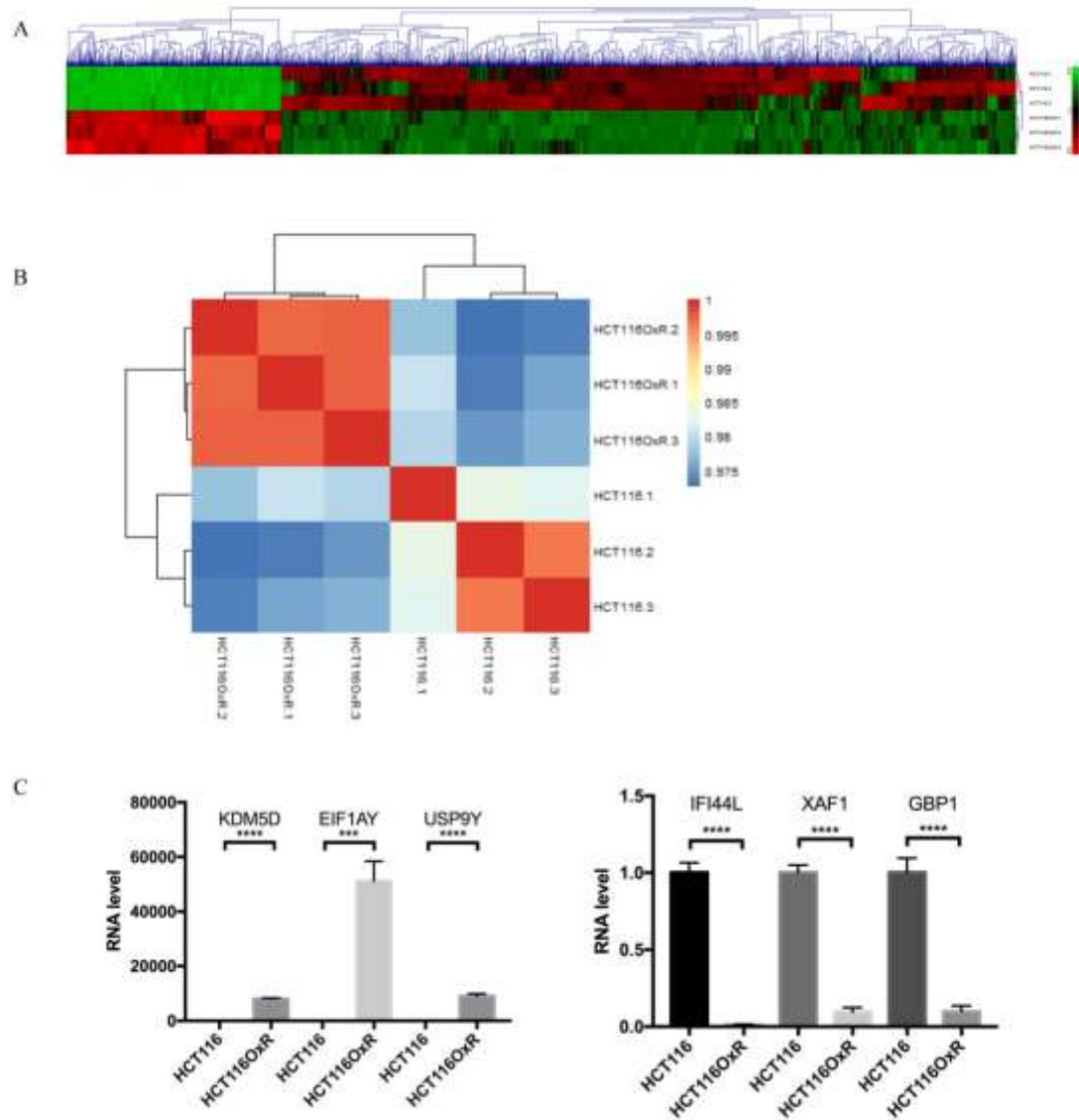


Figure S5. Differential genes detected by RNA-seq.

A Cluster analysis of differentially expressed genes, red indicates high expression genes, green indicates low expression genes. **B** The correlation coefficient heatmap between samples were established by using the identified 32044 transcripts. The results showed that the 3 samples were highly repeatable. **C** qRT-PCR was performed to verify 3 elevated and 3 decreased genes.

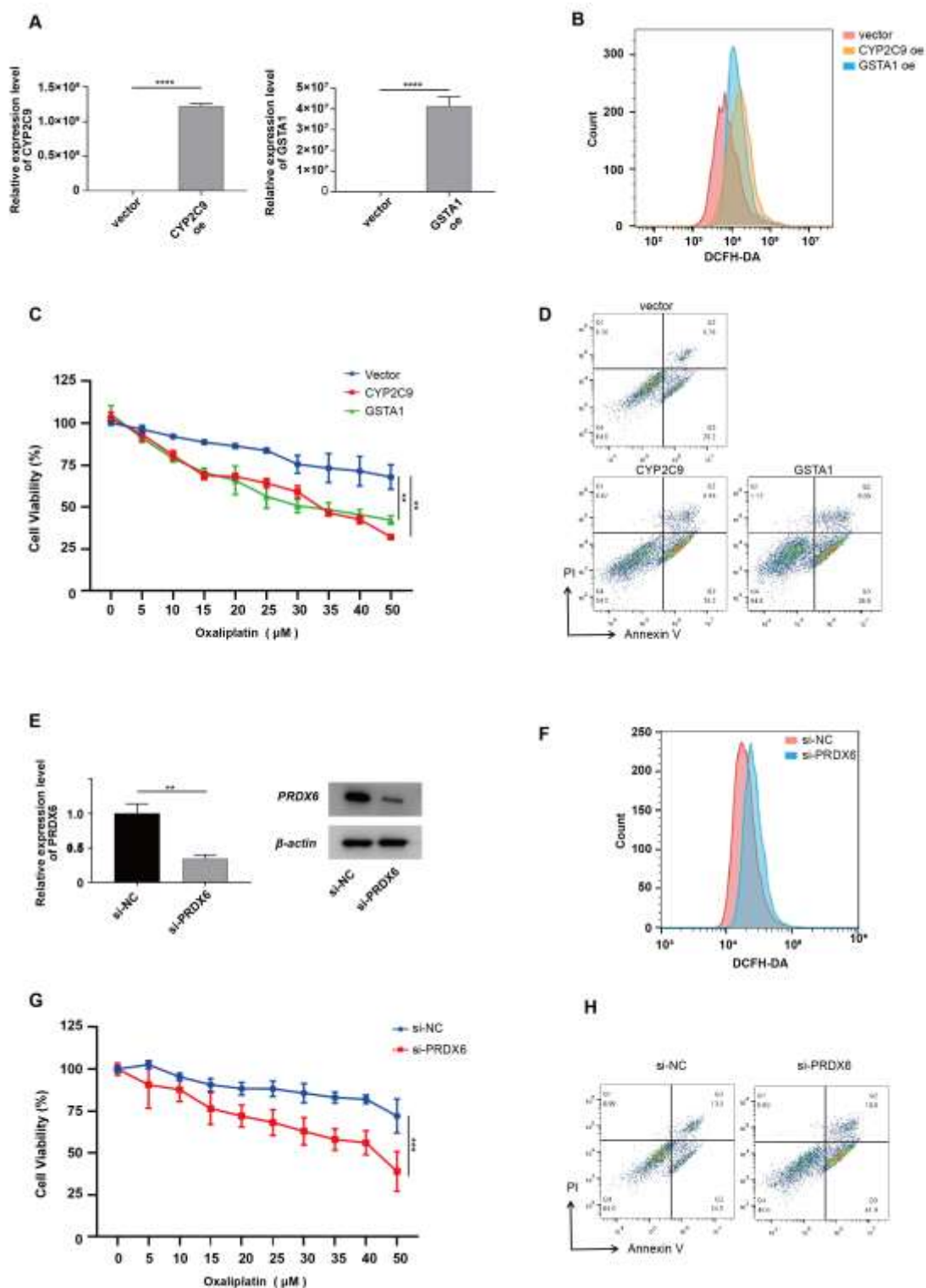


Figure S6. Compartment switching genes can affect cellular response to oxaliplatin by regulating ROS homeostasis.

A The expression of CYP2C9 and GSTA1 mRNA level in HCT116-OxR cells after transduction with indicated overexpression plasmid or vector plasmid was detected by qRT-PCR, *** $P < 0.001$. **B** The level of ROS in indicated cell lines. **C** The responsiveness of HCT116-OxR cells after transduction with indicated overexpression plasmid or vector plasmid to oxaliplatin recorded by CCK8 assay,

****P<0.01. D** The percentages of live cells and apoptotic cells was evaluated in indicated cell lines pretreated with oxaliplatin (100 μ M, 48 h). **E** The knockdown efficiency of si-PRDX6 in HCT116-OxR was detected by qRT-PCR (left) and Western Blot (right). **F** The level of ROS in HCT116-OxR cells treated with si-PRDX6 or corresponding negative control (si-NC). **G** The responsiveness of HCT116-OxR cells treated with si-PRDX6 or si-NC to oxaliplatin was recorded by CCK8, *****P<0.001. H** The percentages of live cells and apoptotic cells was evaluated in indicated cell lines pretreated with oxaliplatin (100 μ M, 48 h).

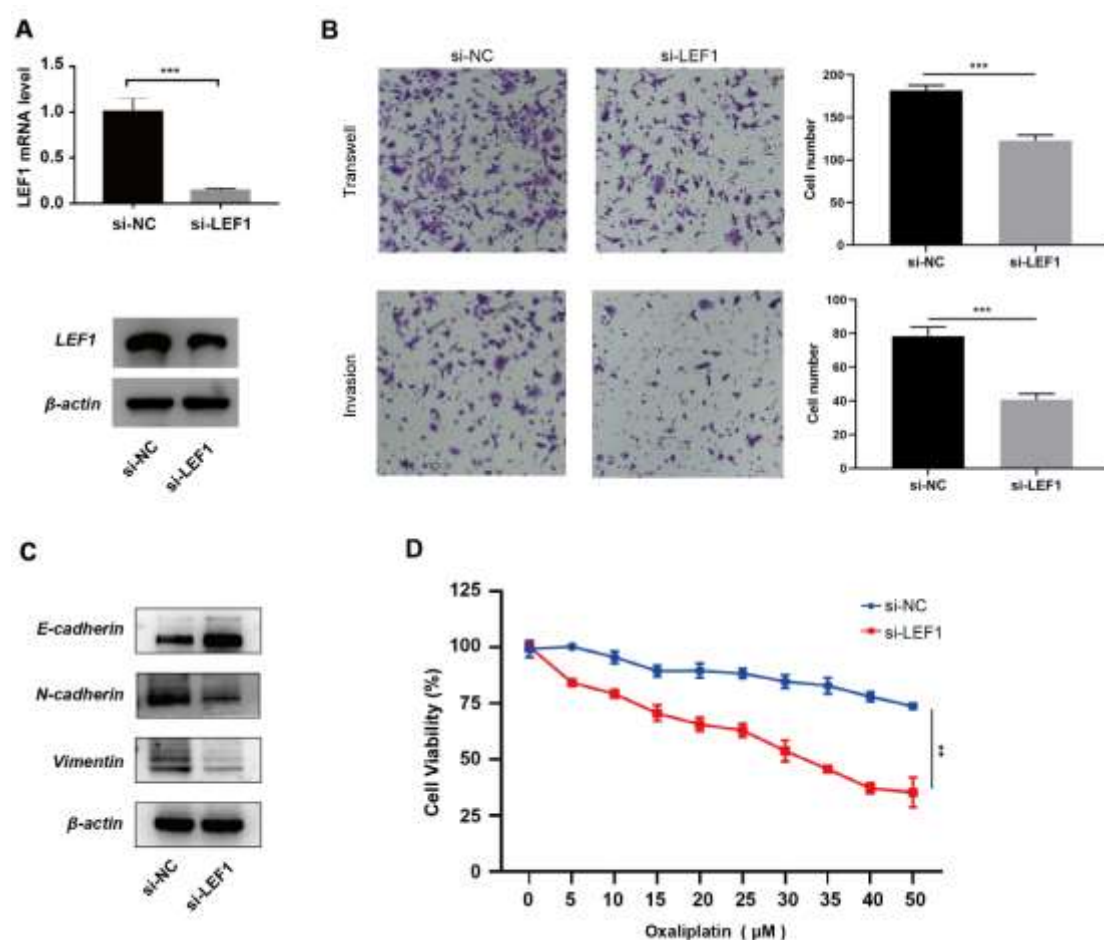


Figure S7. Reducing the expression of LEF1 could weaken the invasion and migration ability of drug-resistant cells.

A The knockdown efficiency of si-LEF1 in HCT116-OxR was detected by qRT-PCR (up) and Western Blot (bottom), *****P<0.001** **B** Cell migration and invasion experiments detected that reducing the expression of LEF1 could reduce the invasion and migration ability of HCT116-OxR cells, ****P<0.01. C** By detecting the expression of EMT markers, it is proved that reducing the expression of LEF1 can reduce the EMT characteristics of drug-resistant cells, *****P<0.001. D** The responsiveness of HCT116-OxR cells treated with si-LEF1 or si-NC to oxaliplatin was recorded by CCK8, ****P<0.01.**