

Figure S1 *C. tropicalis* promotes the chemotherapy resistance of colon cancer to oxaliplatin in mouse xenograft tumor model.

(A) Different treatment in mouse CRC xenograft model. (B) Proportions of macrophage (CD11b+F4/80+), MDSCs (CD11b+Gr-1+), G-MDSCs (CD11b+Ly6G+) and M-MDSCs (CD11b+Ly6C+) in spleen were detected by flow cytometry. (C) Cell viability was detected using CCK8 in SW480 co-cultured with different multiplicity of infection (MOI) of *C. tropicalis*. (D-E) Cell apoptosis was measured by flow cytometry in SW480 co-cultured with *C. tropicalis* (MOI = 1). (F) Wound Healing assay was performed to detect the cell migration in SW480 treated with MOI=1 *C. tropicalis*. Data with error bars are represented as mean \pm SD. Each panel is a representative experiment of at least three independent biological replicates. *p < 0.05, **p < 0.01 and ***p < 0.001 as determined by unpaired Student's t test.

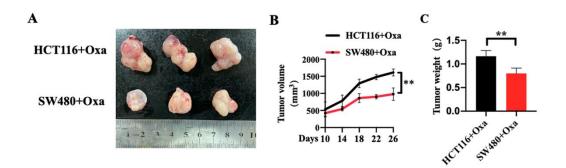


Figure S2 MMR is inhibited in *C. tropicalis*-induced CRC chemotherapy resistance.

(A) Representative image of tumors in mice injected with 5×10^6 HCT116-male cells or 1×10^7 SW480-male cells followed by oxaliplatin. (B and C) Statistical analysis of tumor volumes (B) and weights (C) in mice xenograft with HCT116 or SW480-male cells, n =3/group. Data with error bars are represented as mean \pm SD. Each panel is a representative experiment of at least three independent biological replicates. *p < 0.05, **p < 0.01 as determined by unpaired Student's t test.

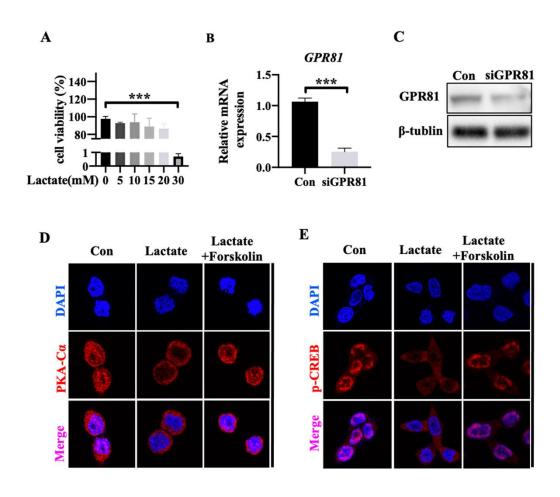


Figure S3 Lactate reduces the expression of MLH1 via GPR81-cAMP-PKA-CREB axis.

(A) Cell viability was detected using CCK8 in SW480 treated with different concentrations of lactate. (B-C) The knockdown function of GPR81 siRNA was verified by testing mRNA and protein levels of GPR81 in SW480. (D) The nucleus translocation of PKAC α in the SW480 cells treated with lactate and forskolin was detected by immunofluorescence. (E) The content of intranuclear p-CREB in lactate and forskolin-treated SW480 were tested by immunofluorescence. Data with error bars are represented as mean \pm SD. Each panel is a representative experiment of at least three independent biological replicates. *p<0.05, **p<0.01 and ***p<0.001 as determined by unpaired Student's t test.