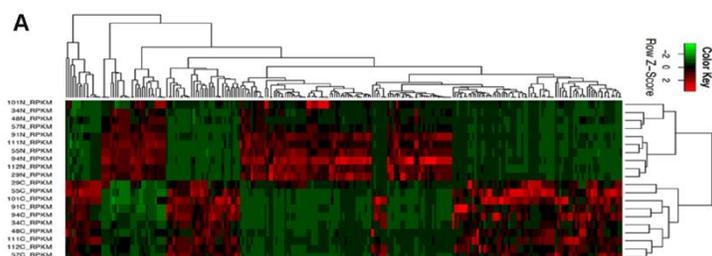
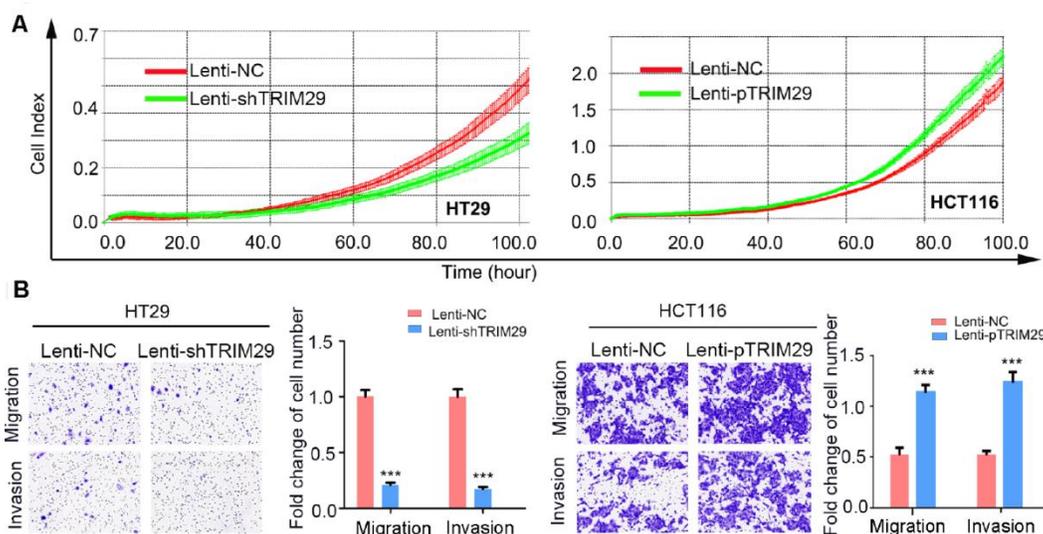


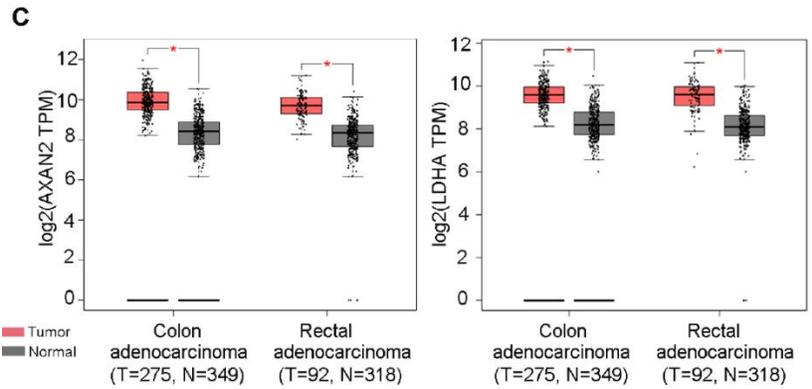
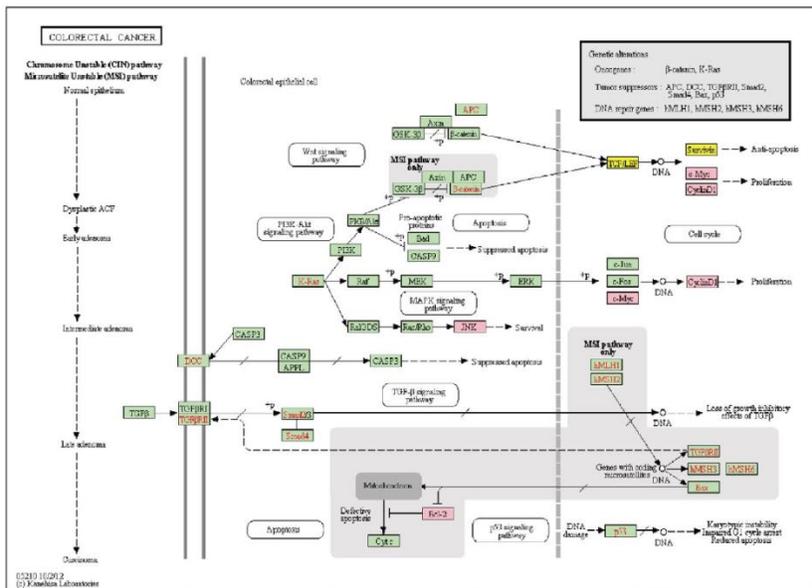
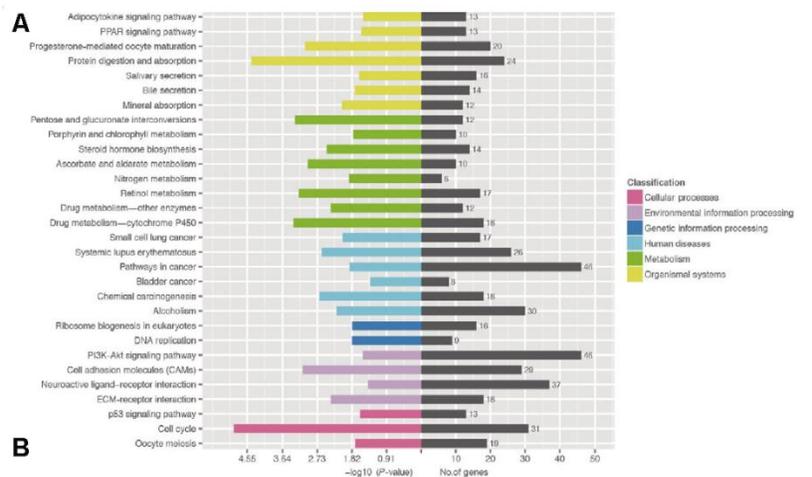
SUPPLEMENTARY FIGURES



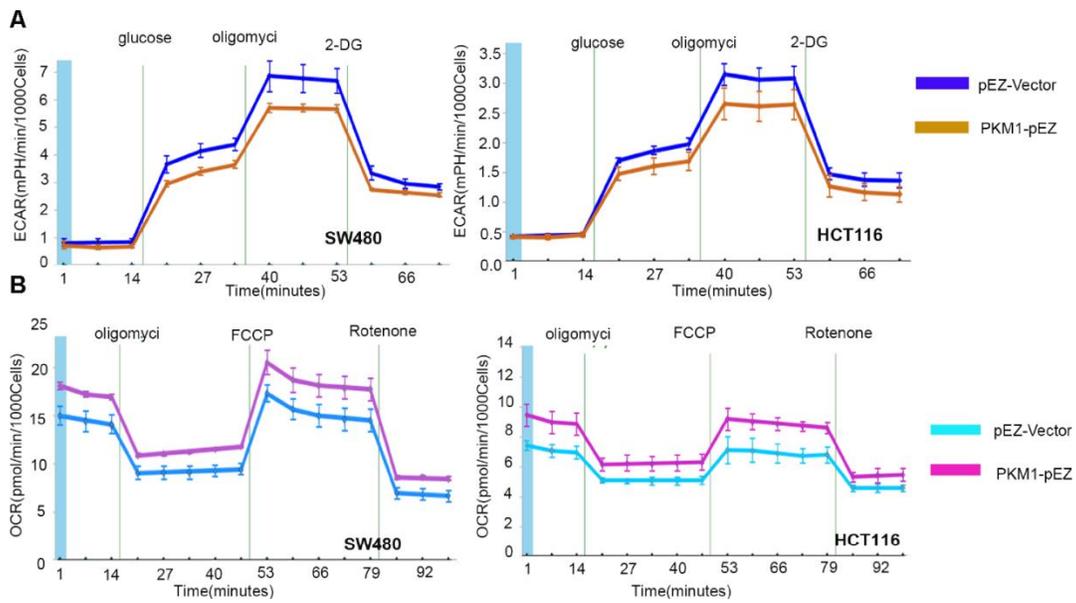
Supplementary Figure 1. (A) Heat map of the mRNA profile showing DEGs in CRC.



Supplementary Figure 2. HT29 cells were stably infected with lenti-NC or lenti-shTRIM29 and HCT116 cells were stably infected with lenti-NC or lenti-pTRIM29. (A) RTCA-MP was performed. (B) Transwell assays were performed. The statistical analysis was performed using the two-tailed Student's t-test. ** $P < 0.01$, *** $P < 0.001$.



Supplementary Figure 3. (A) Enrichment statistics of KEGG pathways in a sample-classification map. (B) The CRC signaling pathways. (C) The expression of AXAN2/LDHA. Data were collected from the TCGA and GTEx databases using GEPIA (<http://gepia.cancer-pku.cn/index.html>).



Supplementary Figure 4. (A) The ECAR and (B) OCR of cells transfected with NC or Si2 was measured. The statistical analysis was performed using the two-tailed Student's t-test. $**P < 0.01$, $***P < 0.001$.