



Supplementary Fig. 5. Genomic and epigenomic data of hepatocellular carcinoma (HCC) patients. (A) Diacylglycerol O-acyltransferase 2 (DGAT2) expression level of patients paired with adjacent normal tissue. (B) Methylation level of CpG island on DGAT2. (C) Copy number variant of DGAT2 in HCC patients. (D) Number of genes according to the number of patients with mutation. DGAT2 is silently mutated in two patients. Data represent mean \pm standard deviation. NS, not significant; SNP, single nucleotide polymorphism; TCGA, The Cancer Genome Atlas. ^a $P < 0.01$, ^b $P < 0.0001$: two-tailed Student's *t*-test in (B, C); two-tailed Mann-Whitney *U* test in (A).