

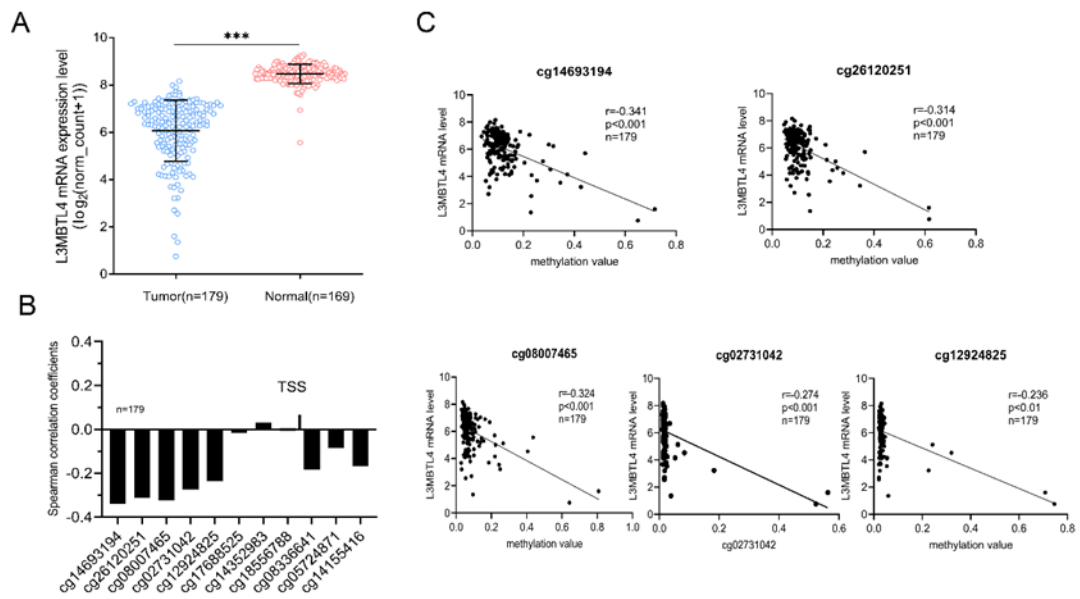


Figure S1. The expression and methylation status of L3MBTL4 in TCGA and GTEx database. (A) The expression levels of L3MBTL4 in pancreatic cancer tissue and normal tissue samples. *** $P < 0.001$. (B) The coefficients of L3MBTL4 mRNA expression and methylation status of 11 CpG sites in the promoter region of L3MBTL4 by Spearman analysis. (C) Scatter plots representing a reverse association between L3MBTL4 mRNA expression and L3MBTL4 CpG sites methylation.