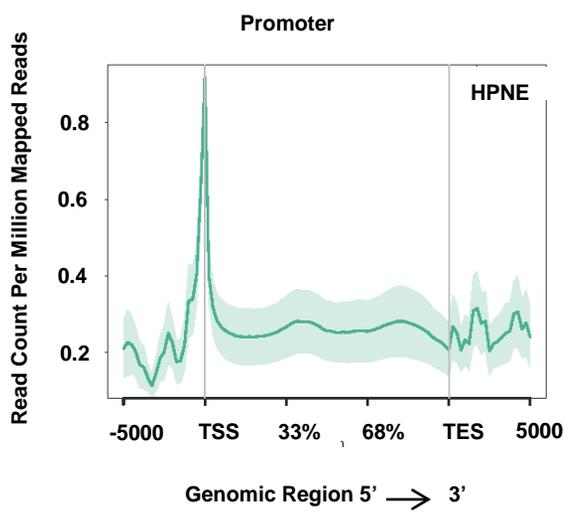
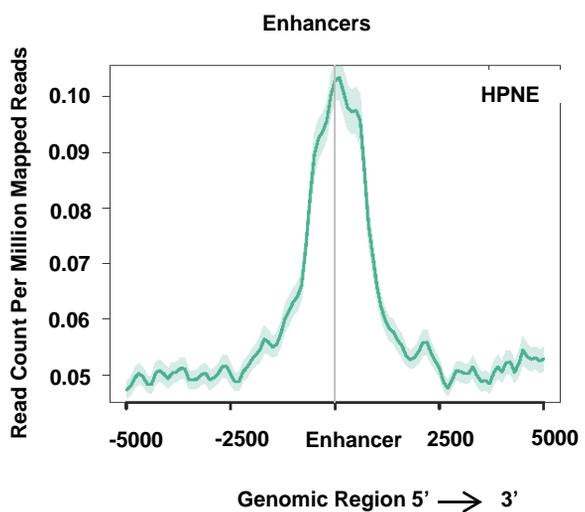


**Supplemental\_Fig\_S3.** Chromatin accessibility patterns from ATAC-seq in pancreatic cells define regulatory regions of the genome. ATAC-seq peaks in HPNE pancreatic cells are enriched in (A) Gene Bodies, (B) known enhancers, (C) known **DNase** hypersensitivity regions and (D) CpG islands.

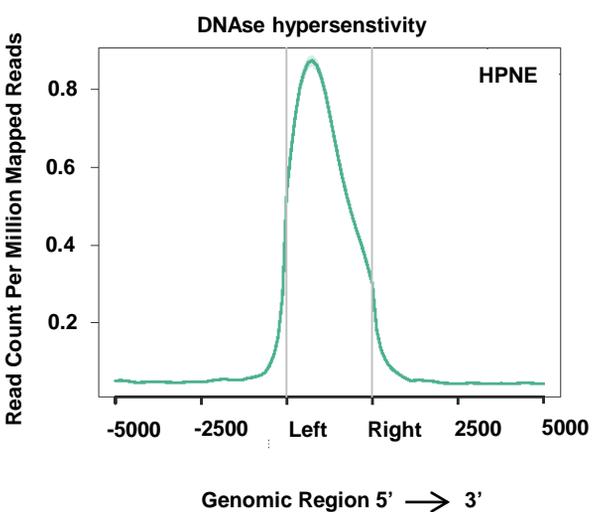
A



B



C



D

