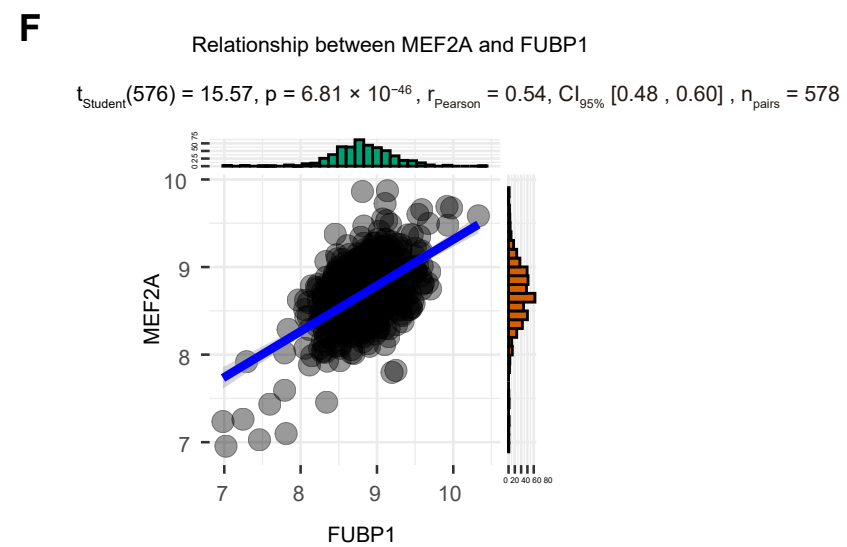
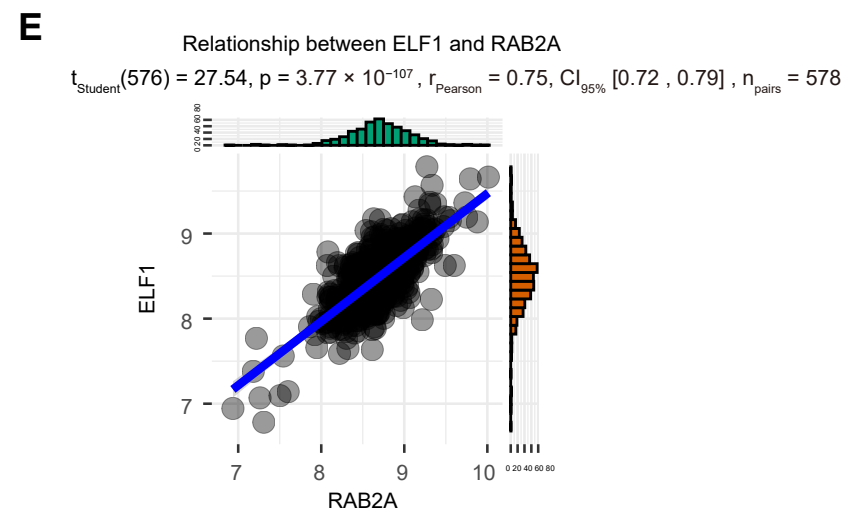
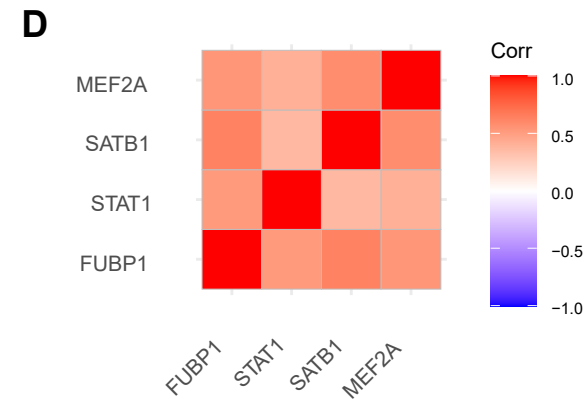
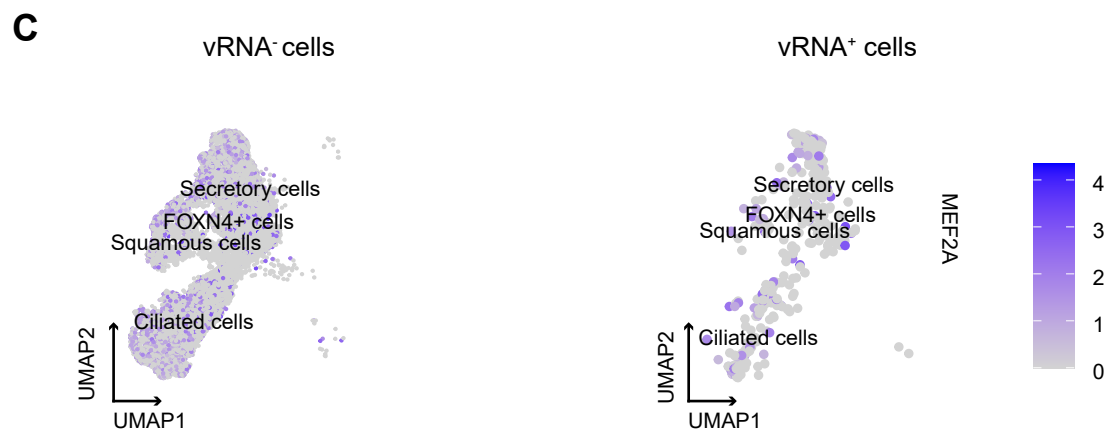
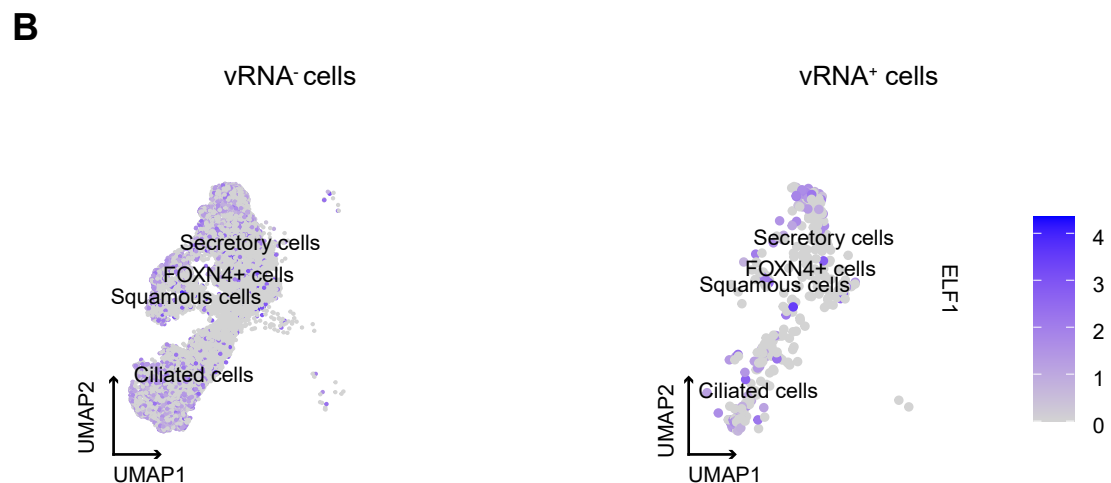
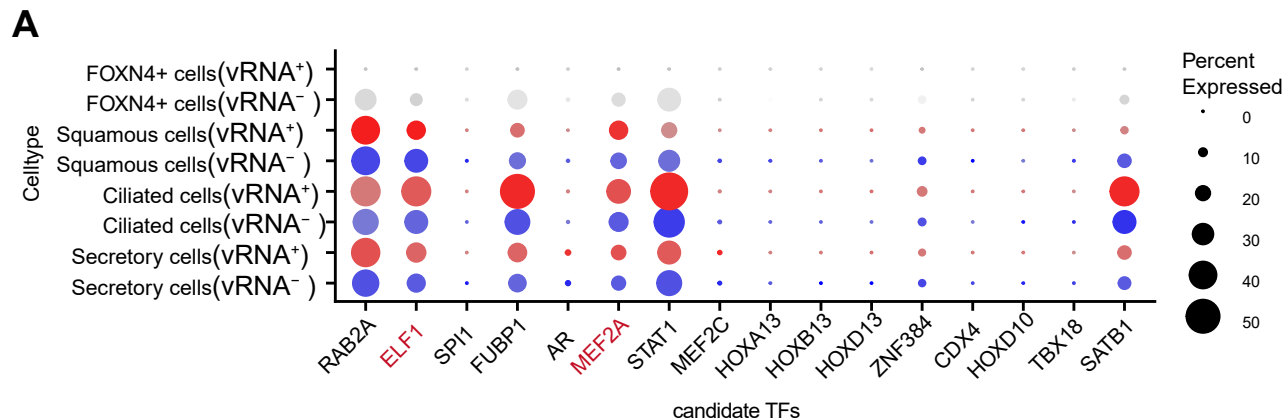




Supplementary Figure S4 Expression of candidate TFs in single-cell RNA sequencing data and relationship between TFs and RBPs Dotplot showing the number of cells expressing candidate Transcription factors in both vRNA+ and vRNA? groups. (B) heatmap showing the correlation between RBPs and candidate TFs. Featureplots show expression levels of (B) ELF1, (C) MEF2A in scRNA-seq data grouped by SARS-CoV-2 RNA status. Correlationplot showing relationship between (E)ELF1 and RAB2A and (F) MEF2A and FUBP1.