Bulk and single-cell transcriptomics identify tobacco-use disparity in lung gene expression of ACE2, the receptor of 2019-nCov

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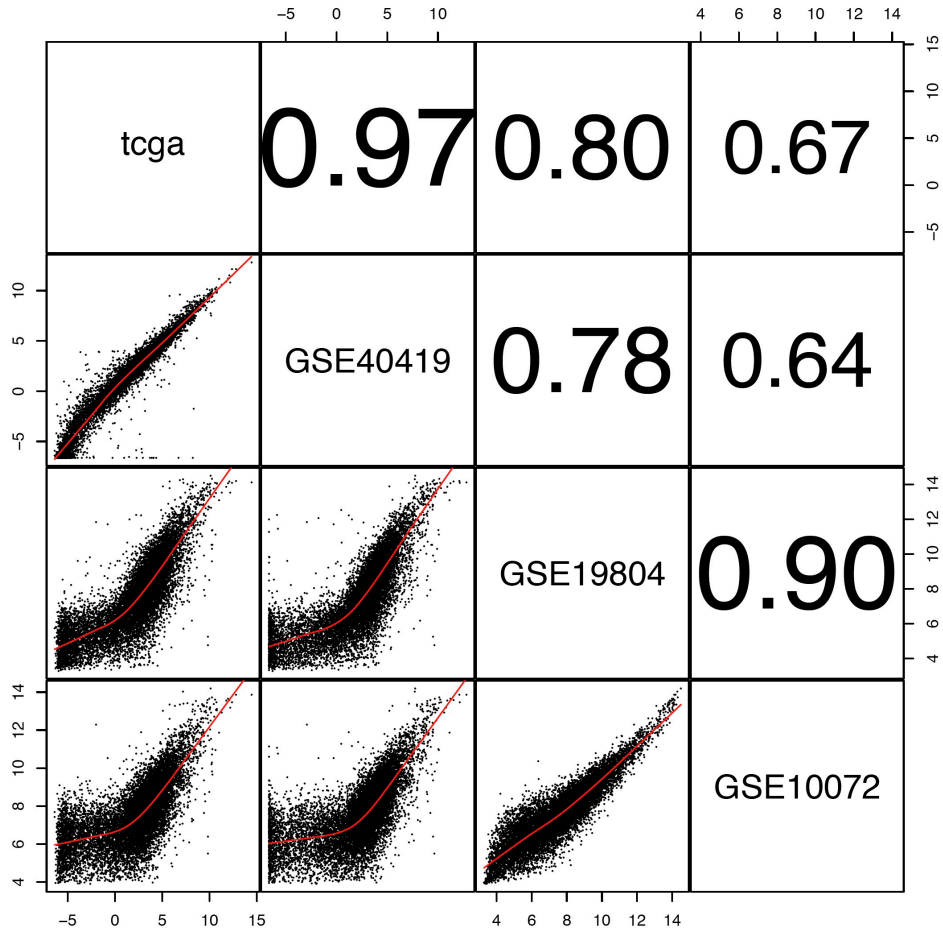
Figure S1. Correlation of four datasets.
Lower panel shows pairwise scatter plots of data mean across samples in each dataset.

Upper panel shows their corresponding Pearson correlation coefficients.
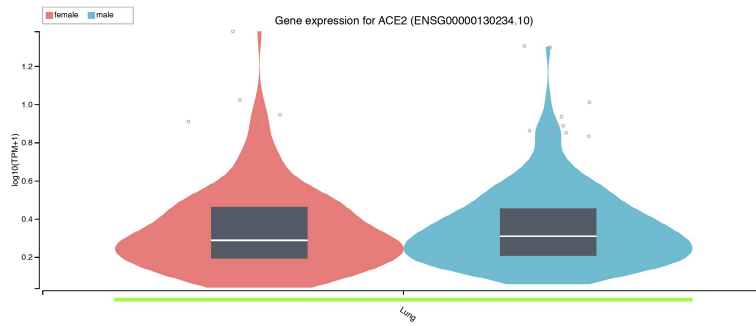
Figure S2. *ACE2* **gene expression in GTEx female and male lung tissues.**
y-axix shows the log10 scaled RNA-seq Transcript Per Million (TPM) values.
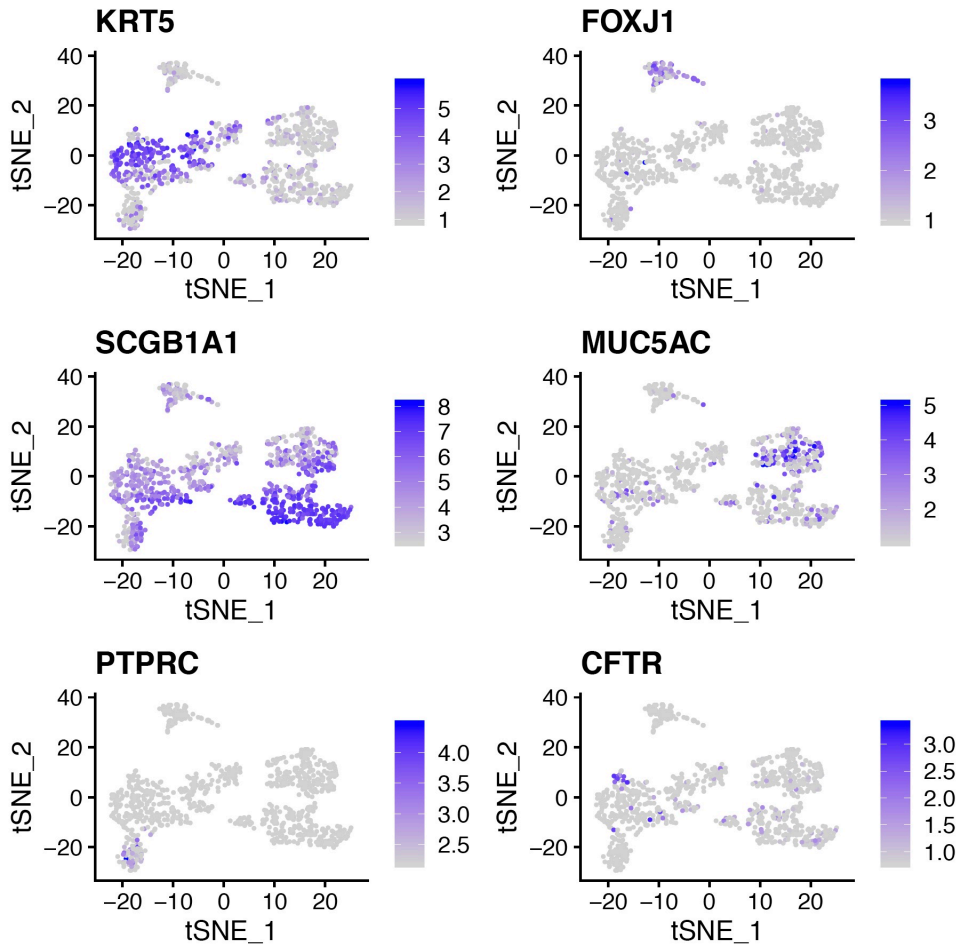
Figure S3. Expression profile of bronchial epithelium cell markers.
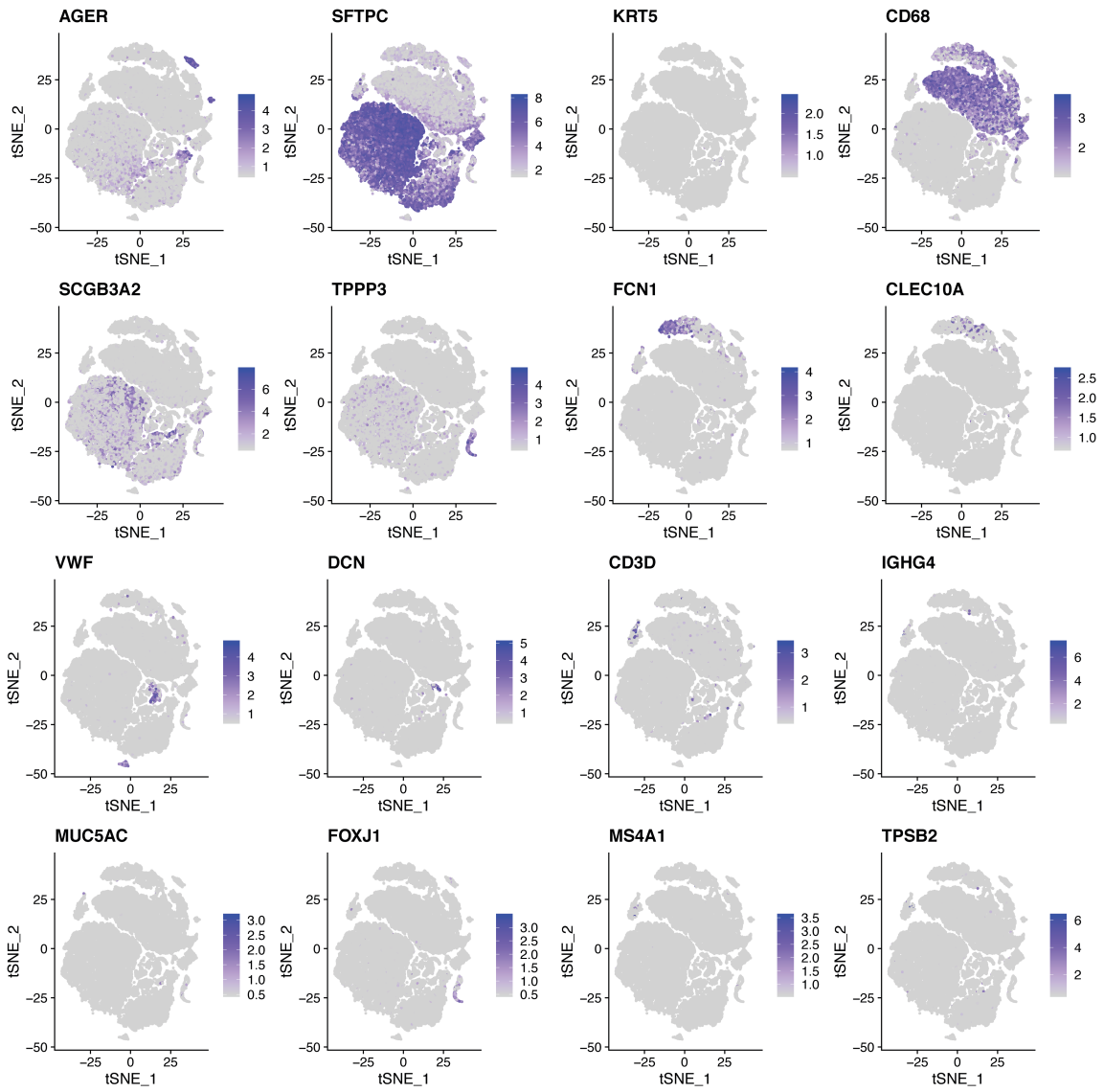
Figure S4. Expression profile of lung tissue cell markers.
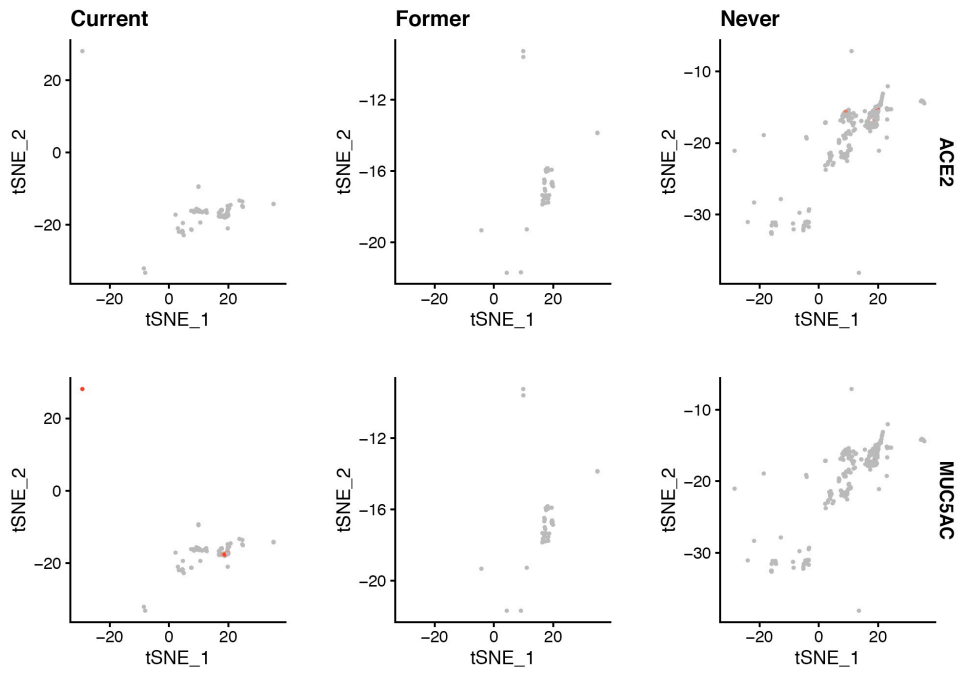
Figure S5. Expression profile of *MUSC5AC* (top) and *ACE2* (bottom).