

Fig. S1 The lung scRNA-seq data analysis results of 8 donors (a-h). For each donor, the cell cluster co-expressing ACE2, SFTPB and SFTPC are identified as AT2.

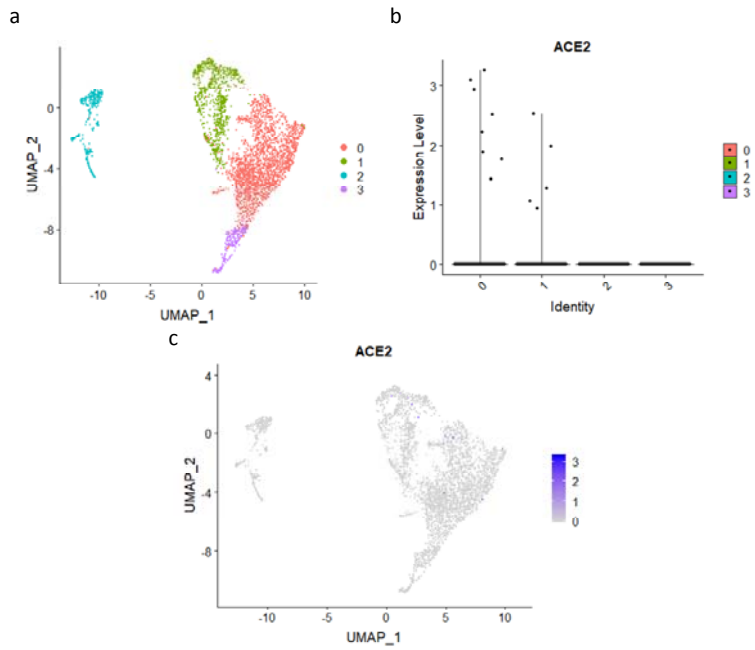


Fig. S2 Nasal mucosa scRNA-seq data analysis showed few cells expressing ACE2. a) The cells were categorized into 4 clusters. b) Violin plot showing the ACE2 expression distribution of different cell clusters. c) The ACE2 expression distribution demonstrated in scatter plot.

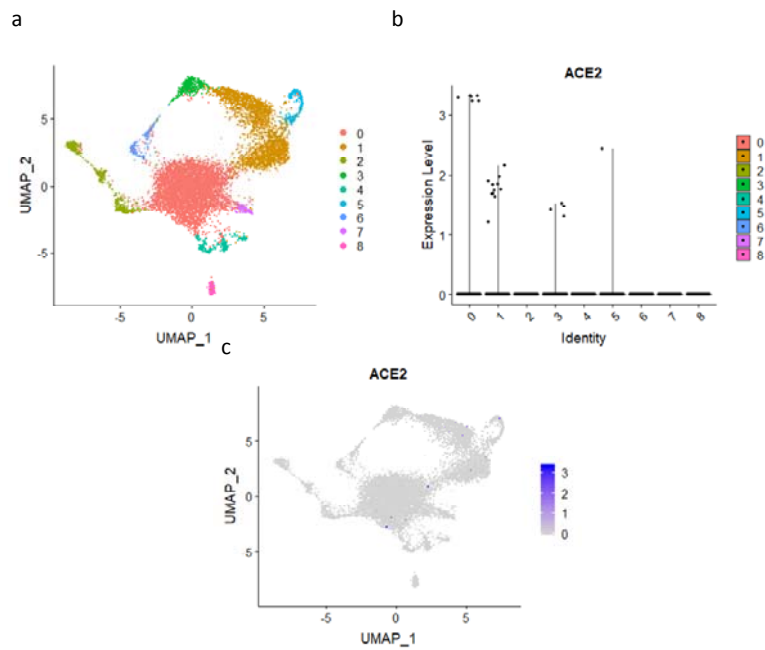


Fig. S3 Bronchus scRNA-seq data analysis showed few cells expressing ACE2. a) The cells were categorized into 8 clusters. b) Violin plot showing the ACE2 expression distribution of different cell clusters. c) The ACE2 expression distribution demonstrated in scatter plot.

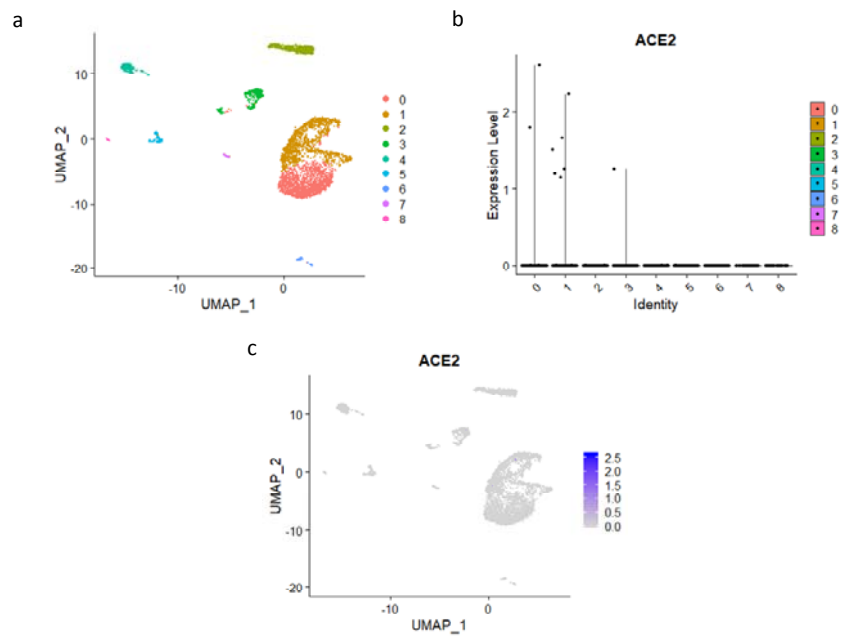


Fig. S4 Stomach scRNA-seq data analysis showed very few cells expressing ACE2. a) The cells were categorized into 9 cell types. b) Violin plot showing the ACE2 expression distribution of different cell clusters. c) The ACE2 expression distribution demonstrated in scatter plot.

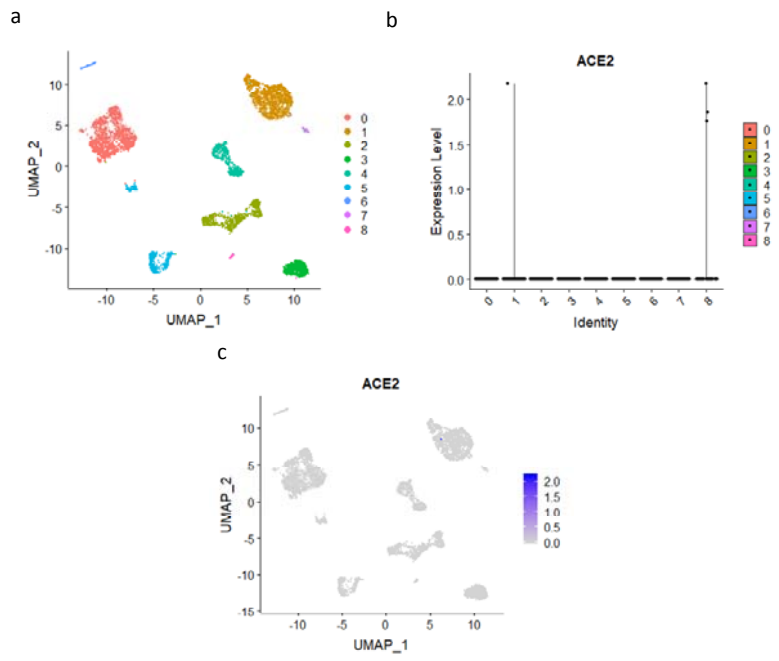


Fig. S5 Liver scRNA-seq data analysis showed almost no cell expressing ACE2. a) The cells were categorized into 9 cell types. b) Violin plot showing the ACE2 expression distribution of different cell clusters. c) The ACE2 expression distribution demonstrated in scatter plot.