Electronic Supplementary Material

Long Term Culture of Human Kidney Proximal Tubule Epithelial Cells Maintains Lineage Functions and Serves as an *Ex vivo* Model for Coronavirus Associated Kidney Injury

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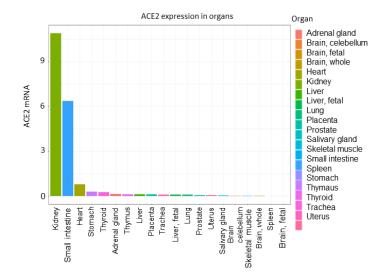


Fig. S1 RNA-sequencing analysis of ACE2 expression level from public datasets. Bar plot of ACE2 expression level in different tissues from 20 human tissues reported in SRP056969.

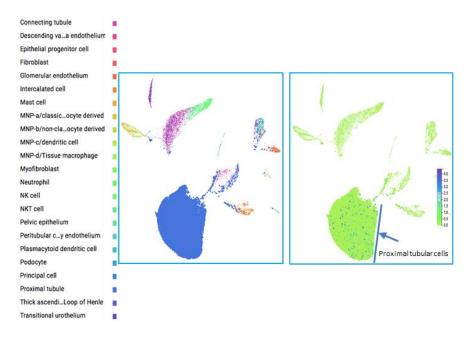


Fig. S2 Single cell RNA-sequencing (scRNA-seq) analysis of ACE2 expression in kidney cells from public datasets. ACE2 expression is predominantly in proximal tubule epithelial cells. This data can be visualized and assessed through a special website portal (www.covid19cellatlas.org).