

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: RNAseq analysis of SARS-CoV-2 infected versus mock infected Vero-TMPRSS2+ cell at 8 hpi. n=3 biologically independent samples were examined. The P value and log fold change in the table were generated with DESeq2 under default settings with Wald test and normal shrinkage, respectively.

File Name: Supplementary Data 2

Description: Quantitative spent media analyses of SARS-CoV-2 infected versus mock infected Vero E6 TMPRSS2+ cell at 8 hpi. n=6 biologically independent samples were examined. P-values were generated with two-tailed P value from Student's t-test.

File Name: Supplementary Data 3

Description: Intracellular metabolite profiling of SARS-CoV-2 infected versus mock infected Vero E6 TMPRSS2+ cell at 8 hpi. n=6 biologically independent samples were examined. P-values were generated with two-tailed P value from Student's t-test.

File Name: Supplementary Data 4

Description: . Oligonucleotides used in the study.