

Epitranscriptomic N^6 -methyladenosine profile of SARS-CoV-2-infected human lung epithelial cells

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Supplemental Tables (three separate Excel files)

Table S1: All differentially m^6a -modified transcripts in SARS-CoV-2-infected A549-hACE2 cells vs. mock-infected control cells.

Table S2: All pathways associated with at least one differentially methylated mRNA in SARS-CoV-2-infected A549-hACE2 cell vs. mock-infected control cells.

Table S3: Primer sequences used for qPCR amplification of selected targets.