



Figure S1. Gene expression analysis in SARS-CoV-2 infected vs not-infected Calu3 cells up to 72 hours-post-infection (hpi). (A) *NBAS* and core genes of the NMD pathway; (B) *STAU1* and known NMD targets; and (C) components of the syntaxin 18 complex known to interact with *NBAS*. Each bar represents the mean of at least four technical replicates \pm standard error of the mean (SEM); analysis of the outliers was performed using the ROUT method and $Q=1\%$. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$. One-way ANOVA with the Bonferroni multiple comparison test.