Extended Data Fig. 6. SARS-CoV-2 5’ end svRNAs are present in exosomes secreted from infected cells. (a) Schematic of exosome purification and RNA-seq in this study. (b) Immunoblot analysis of CD63 and SARS-CoV-2 spike protein with RNAs from input supernatants and exosome-immunoprecipitates. (c) Fraction of svRNA categories in exosomes released from SARS-CoV-2-infected cells. The fraction of each svRNA category in the svRNA population of each infection is shown, according to the color legend in the figure. (d) Read map of 5’ UTR svRNAs from exosomes. Reads were mapped to the positive sense RNA. Read counts were quantified for each nucleotide. (e) Size distribution of 5’ UTR svRNAs in the exosomes. Representative svRNA fragment with the highest count is also shown.