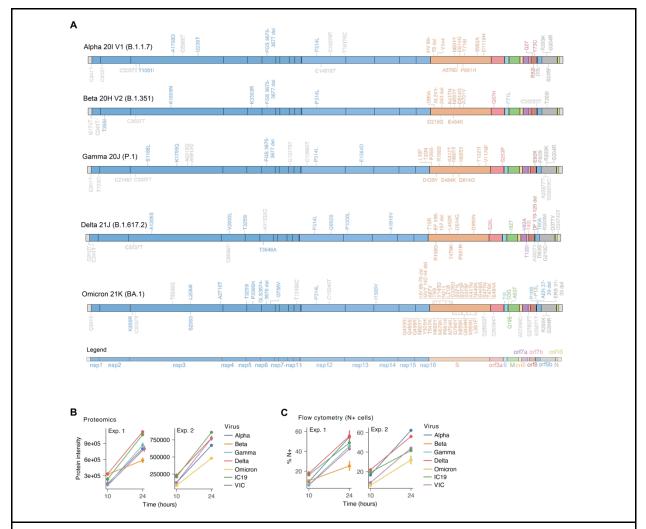
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Genomic mutation map and replication kinetics of the SARS-CoV-2 variants of concern.

- (A) Annotated map of protein-coding (colored) and non-coding (light gray) mutations for each SARS-CoV-2 variant of concern. Color corresponds to the viral gene as depicted in the legend (bottom).
- (B) Quantification of the sum of non-structural protein intensities from abundance proteomics for each virus in each experiment (1 or 2) at 10 and 24 hours post infection in Calu-3 cells. Error bars depict the standard error (SE).
- (C) Flow cytometry assessing the percentage of cells positive for nucleocapsid (N) staining for each virus at 10 and 24 hours post infection in Calu-3 cells. Error bars represent the standard error (SE).