

SUPPLEMENTARY FIGURE. Phylogenetic tree displaying the 36 positive SARS-CoV-2 specimens from surveillance of sentinel emergency departments (EDs), conducted March 1–20, 2020, with select full genome sequences available from the Global Initiative on Sharing All Influenza Data (GISAID), as of April 1, 2020. Red dots represent the positive SARS-CoV-2 specimens from sentinel EDs; identical sequences have overlapping red dots. Pink dots are selected New York state sequences from GISAID.

