A — Variant proportion over time

B — Mean size of clusters of identical sequences over time

C Testing for a difference in transmissibility between variant and non-variant from the size distribution of clusters of identical sequences

Variant legend:
- Other
- D614G
- Alpha
- Beta
- Gamma
- Delta
- Epsilon
- BA.1
- BA.2 (21L)
- BA.2.12.1
- BA.4
- BA.5
- BA.2.75
- BQ.1
- XBB

R MLE estimates consistent with variant transmission advantage
- ● Yes
- ▲ No

Panel C facets: D614G, Epsilon, Alpha, Delta, Omicron (BA.1), Omicron (BA.2), Omicron (BA.4/BA.5)

x-axis (Panel C): Days after 10 variant sequences sampled
y-axis (Panel C): p-value