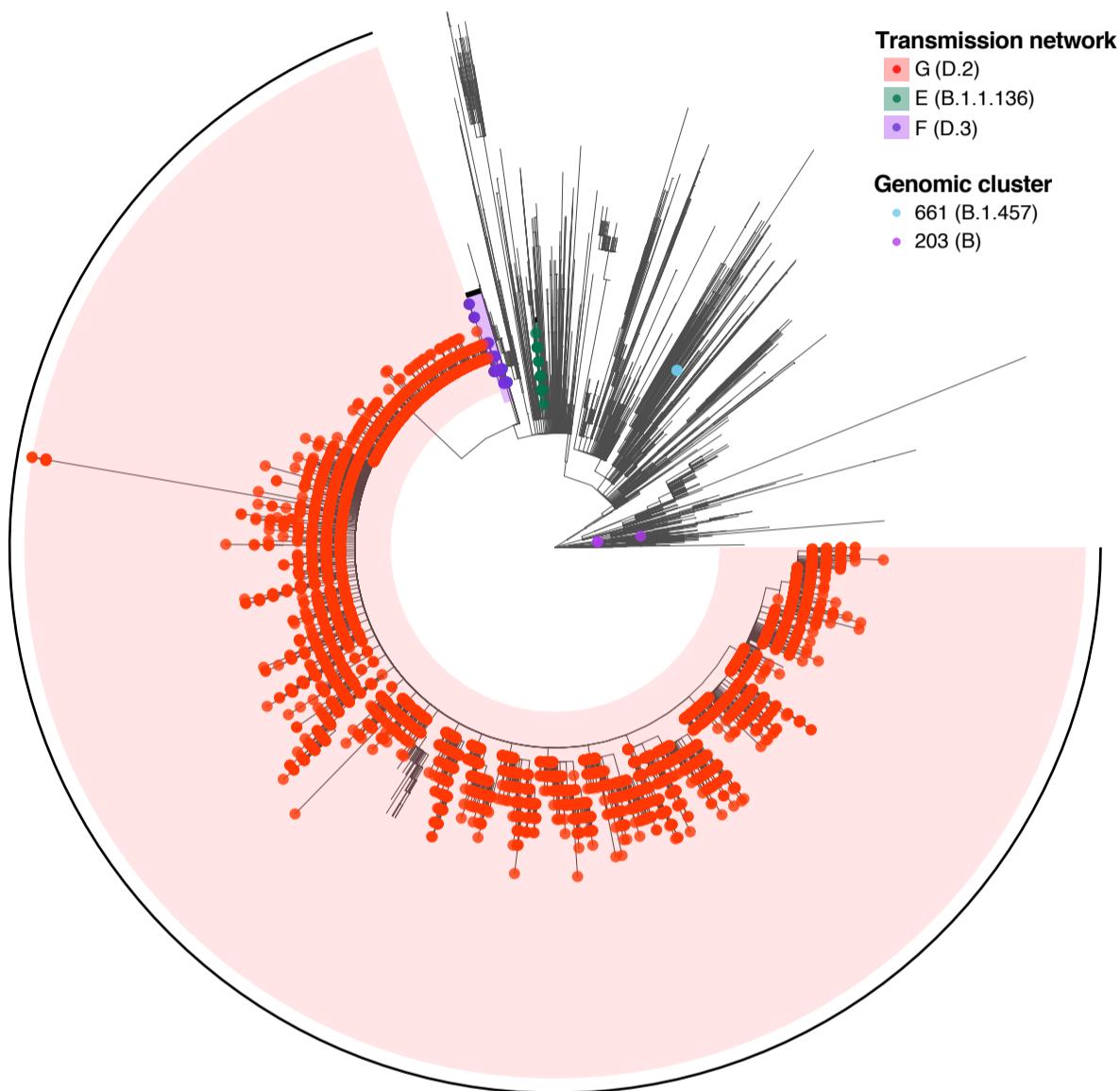


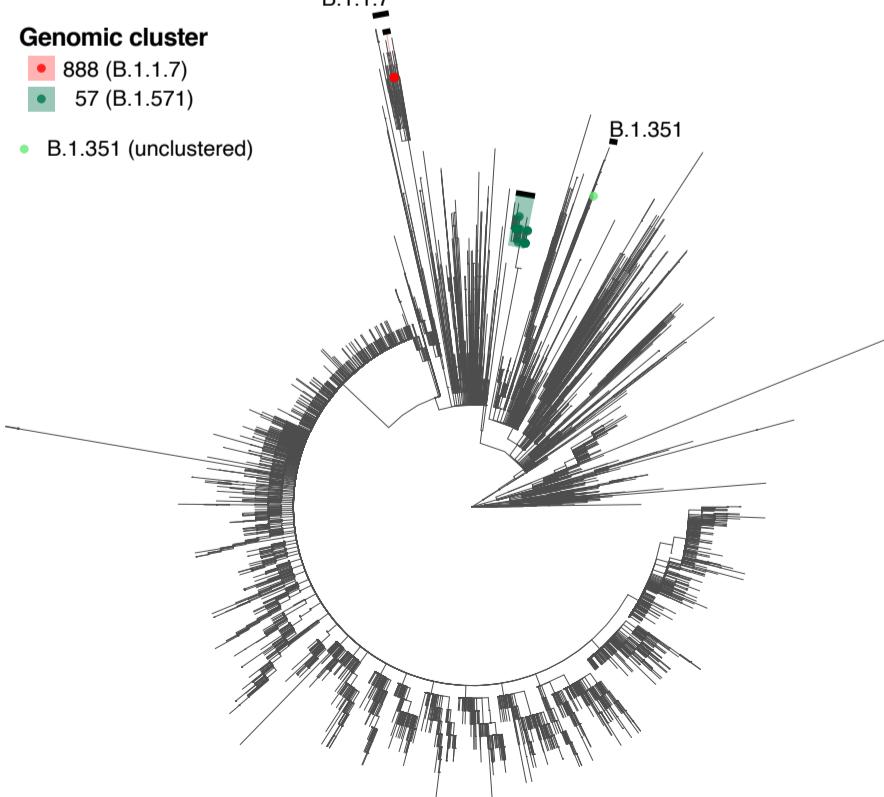
Figure 3: Maximum likelihood phylogenetic trees of Australian SARS-CoV-2 samples

Samples from COVID-19 cases in Victoria diagnosed between Jan 25, 2020, and Jan 31, 2021, and interstate samples from cases uploaded to AusTrakka before Feb 11, 2021, are included. (A) Sequences from Victorian cases diagnosed during period 3 (May 31, 2020, to Dec 6, 2021) identified as within local transmission networks or genomic clusters indicated. Travel-only clusters and unclustered sequences have not been indicated. Further information on genomic clusters 661 and 203 provided in Supplementary Figure 2 (appendix 1, p 7). (B) Sequences from Victorian cases diagnosed during period 4 (Dec 7, 2020 to Jan 31, 2021) identified as within local-transmission genomic clusters or variants of concern are indicated. Regions of the phylogenetic tree containing B.1.1.7 and B.1.351 variants of concern are also labelled. Travel-only clusters and unclustered sequences have not been indicated. (C) Sequences from interstate samples (excluding Victoria) identified within transmission network G.

A



B



C

