

### Figure 3\_updated: Genomic clustering of Victorian COVID-19 cases diagnosed between 21st February and 11th August 2020, with sequence data reported as of 14th August 2020.

Each graph point represents a single case, with date of diagnosis represented on the X-axis and reported genomic cluster on the Y-axis. Cases are coloured orange if the case is suspected to have acquired COVID-19 through overseas travel (data provided by DHHS) and as indicated in the Figure legend.



**Note:** A transmission network, in this context, represents a group of very closely related genomic clusters with a most recent common ancestor. Each transmission network is thought to represent a separate but single importation of the virus into Victoria, with genomic diversity, which is well supported but small in magnitude, arising in Victoria as the virus has circulated within the community and resulting in multiple closely related genomic clusters.