Phylodynamic Analysis

# Location: Nsw

# Most recent sample: 2020-03-17

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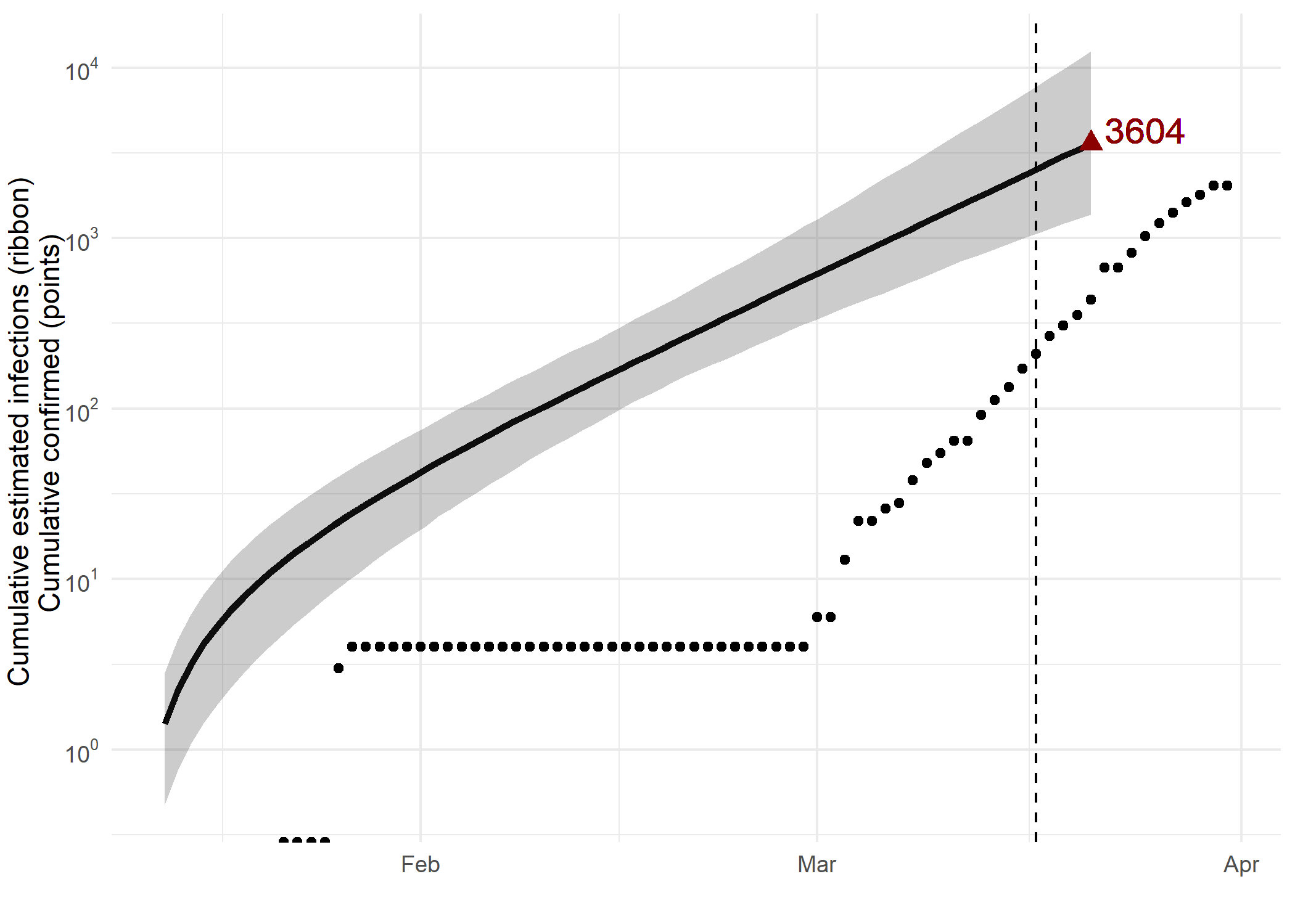
## Background information

#### This is analysis is based on :

* **43 whole genomes** sampled from **within Nsw**
* **62 whole genomes** sampled from outside of **Nsw**
* Samples within Nsw were collected between **2020-01-18** and **2020-03-17**

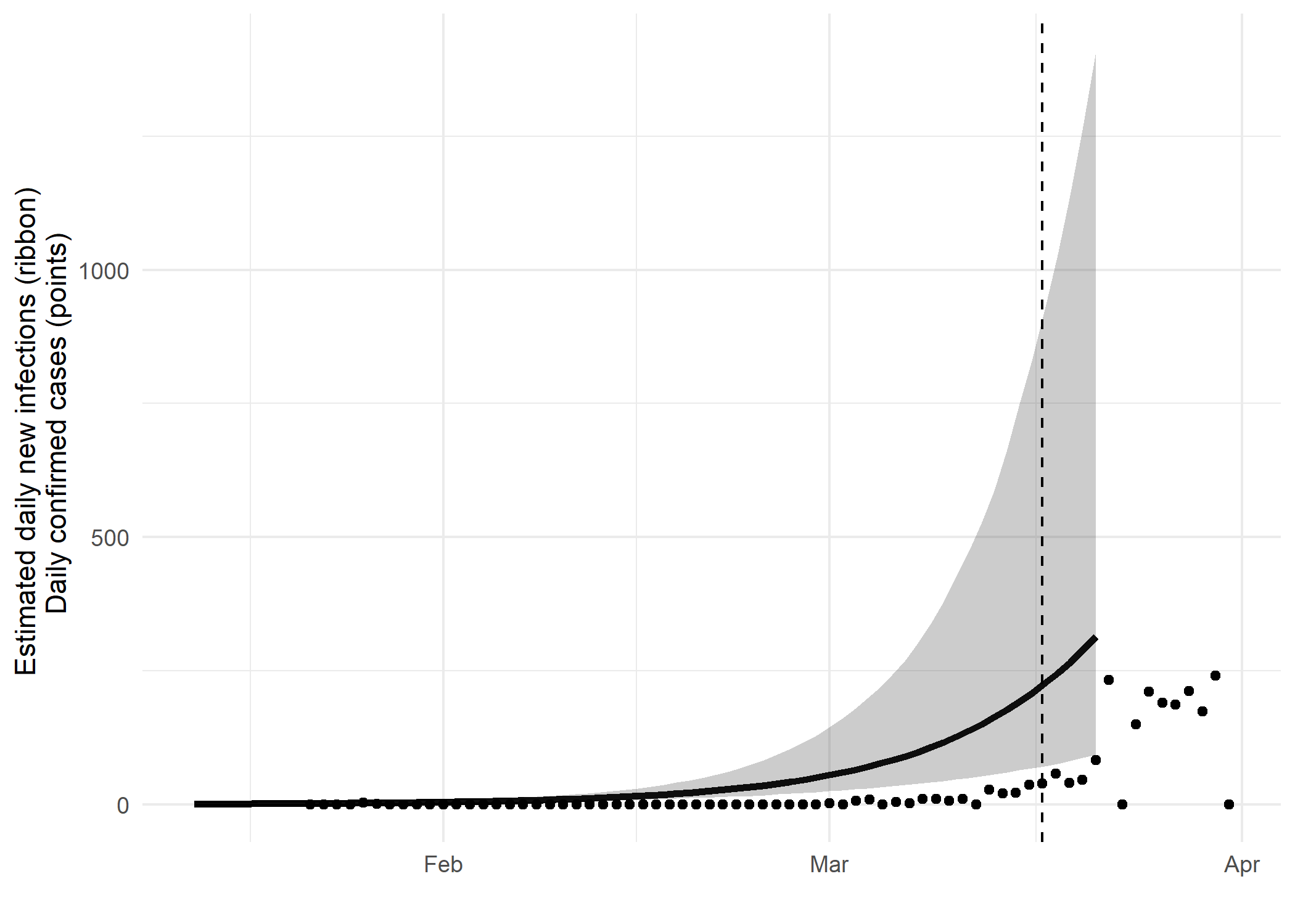
##### To add: [optional plot of sample distribution through time]

## How many are infected in Nsw?

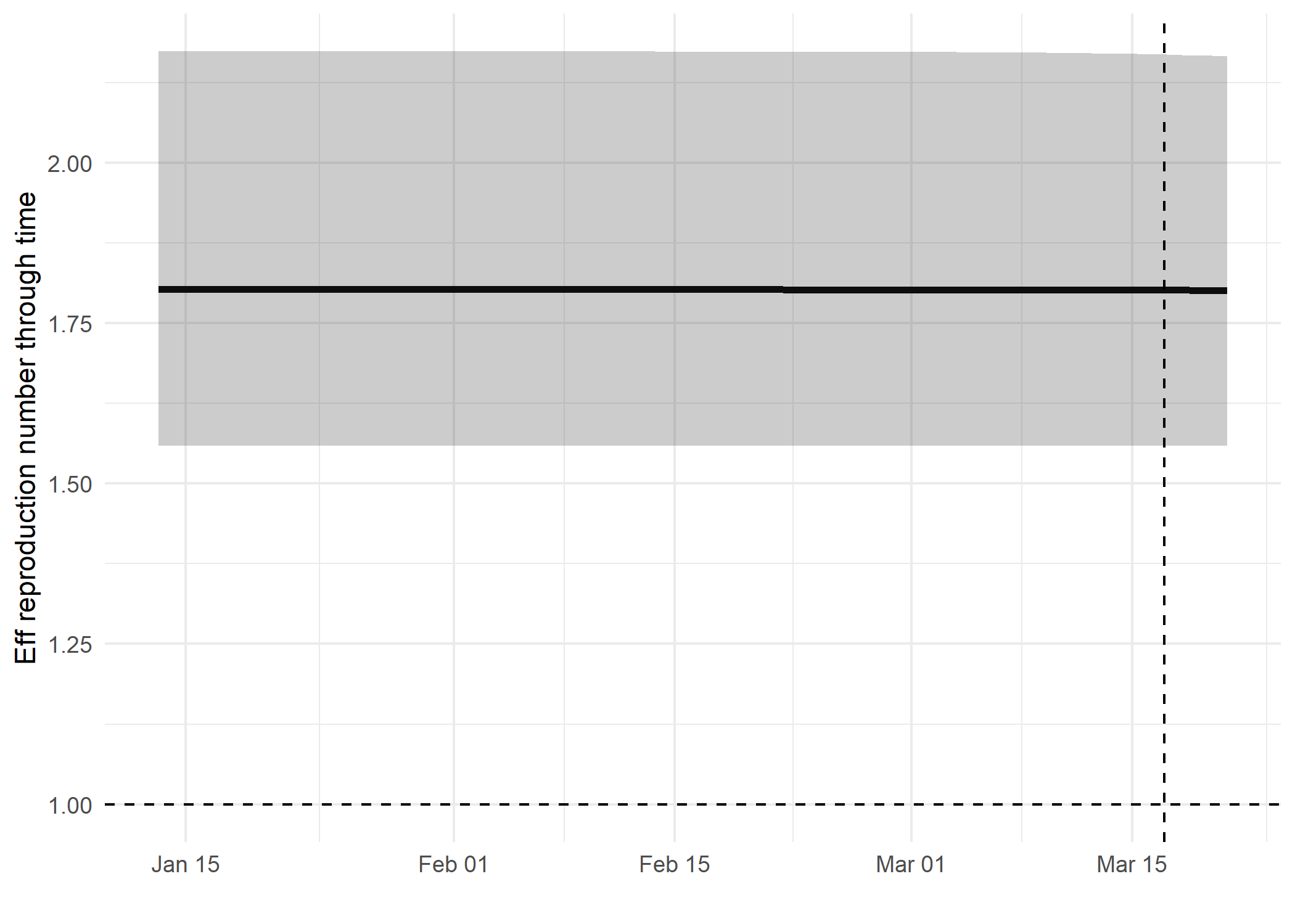


*Figure 1: Cumulative estimated infections through time. Points represent reported cases in Nsw. The dashed line indicates the date of last sample in Nsw in this analysis.*

* Estimated cumulative infections at last sample (2020-03-17): **3604 [1374-12429]** median [95%CI]
* Cumulative confirmed infections reported at 2020-03-17: **210**
* Cumulative number of active infections at 2020-03-17:



*Figure 2: Daily estimated infections through time. Points represent reported cases in Nsw. The dashed line indicates the date of last sample in Nsw in this analysis.*



*Figure 3: Reproduction number through time. The dashed line indicates the date of last sample in Nsw in this analysis.*

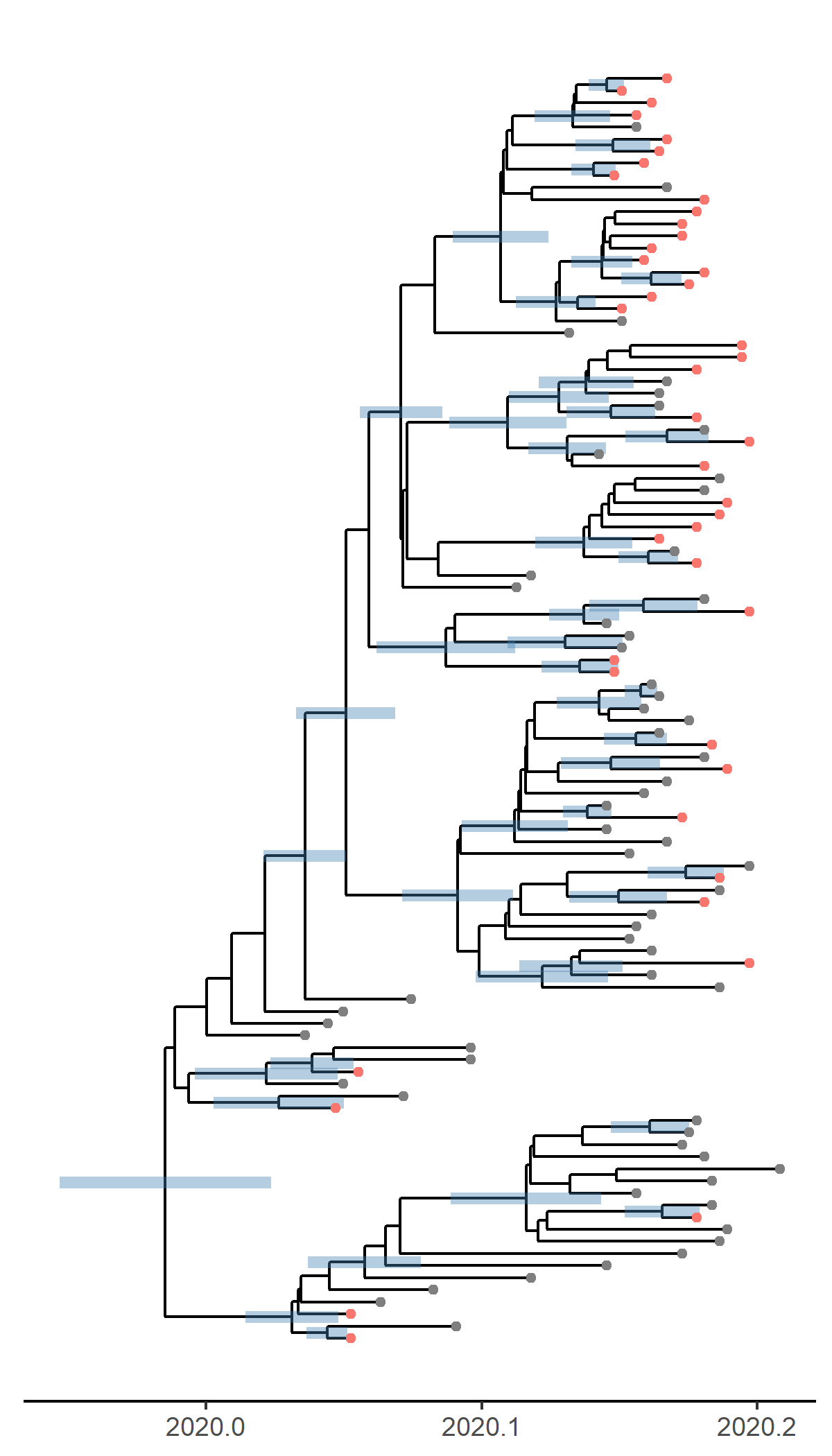
Reproduction number at last sample (2020-03-17): **1.8 [1.56-2.17]** median [95% CrI]

## How quickly has the epidemic in Nsw grown?

Table 1: Reproduction number, growth rate and doubling times

|  |  |  |  |
| --- | --- | --- | --- |
| Quantile | Reproduction number | Growth rate (per day) | Doubling time (days) |
| 50% | 1.81 | 0.0875 | 7.92 |
| 2.5% | 1.56 | 0.0629 | 5.69 |
| 97.5% | 2.2 | 0.122 | 11 |

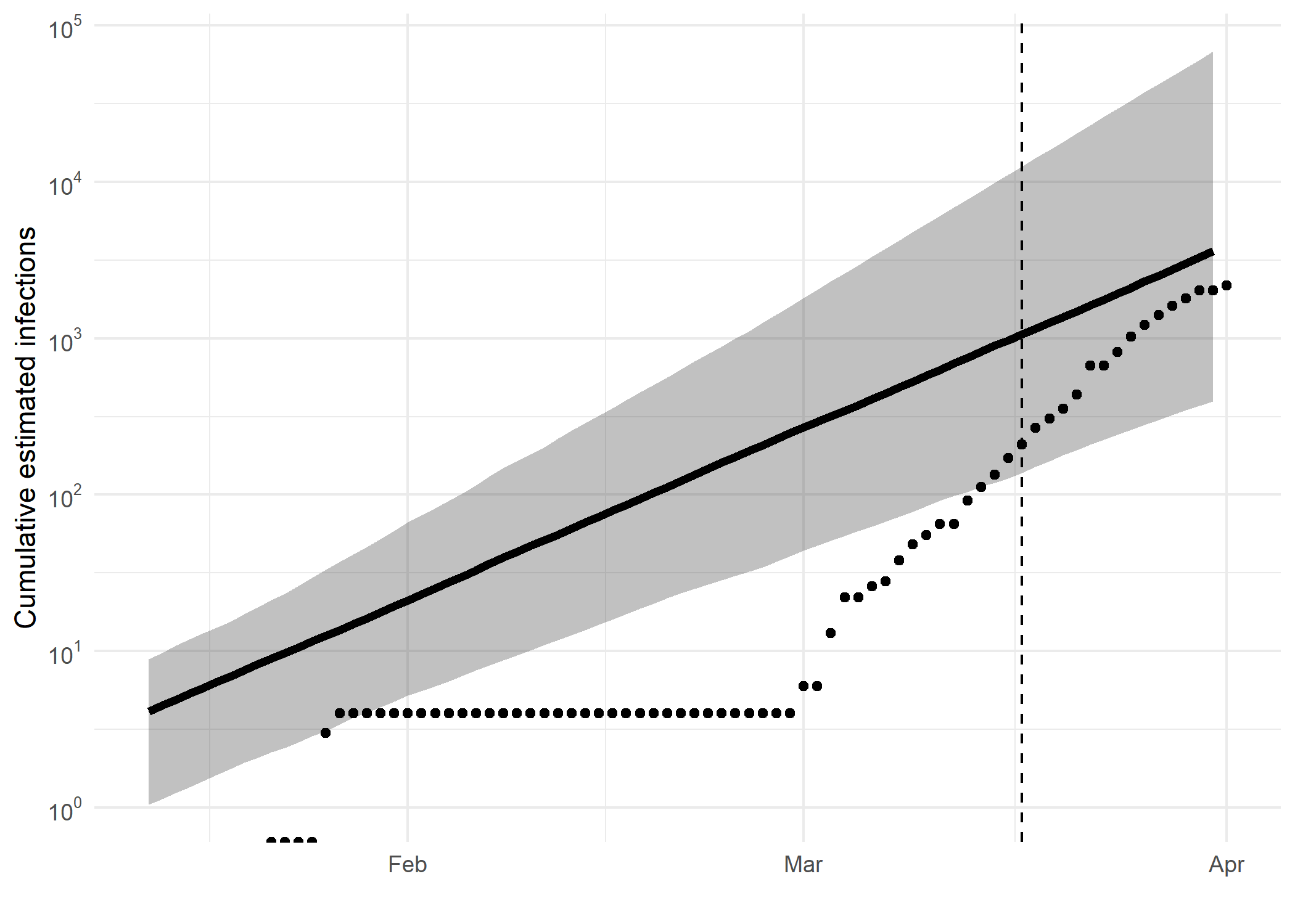
## How has SARS-CoV 2 evolved in Nsw?



*Figure 4: Time scaled phylogeny co-estimated with epidemiological parameters. The colour of the tips corresponds to location sampling; red tips were sampled from within Nsw, blue tips from outside.*

##### Molecular clock rate of evolution: **0.00168 [0.00131-0.0021]** median [95% CrI]

## Predicted cumulative infections over next 14 days (assuming exponential growth):



*Figure 5: Cumulative estimated infections. The dashed line indicates the date of last sample in Nsw in this analysis. The points represent reported cases in Nsw.*

Based on an estimated growth rate of 0.0875 [0.0629 - 0.122] median [95% CrI]:

We estimate cumulative number of infections at last sample (2020-03-17) as: 1063 [138 - 12582]

We estimate number of infections at 2020-03-31 (14 days after last sample) as: 3604 [395 - 68142]

## Methods summary

Details on methods and priors can be [found here](http://whoinfectedwhom.org/seijr0.1.0_methods.pdf).

Model version: seijr0.1.0

Report version: 20200407-223937-ce202008

## Acknowledgements

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Sequence data were provided by [GISAID](http://www.epicov.org) and [these laboratories](http://whoinfectedwhom.org/gisaid_cov2020_acknowledgement_table.xls).