# SARS-CoV-2 State Introductions

## Importation Summary for states

Last modified: 27 Jun 2021

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# 1 Summary

This notebook plots figures about the dataset, applies the importation lag model to the Germany transmission lineage TMRCAs and plots figures with lineage importations.

### 1.1 Data and Method

- GISAID tree until???? as initial tree.
- The tree contains ??? Germany sequences.
- The tree is time-calibrated by TreeTime.
- Sankoff algorithm is used to assign location (Germany and non-Germany) to inner vertices of the tree.

## 2 Germany Sequenced Genomes

#### ## #BE0F34 #B0B0B0

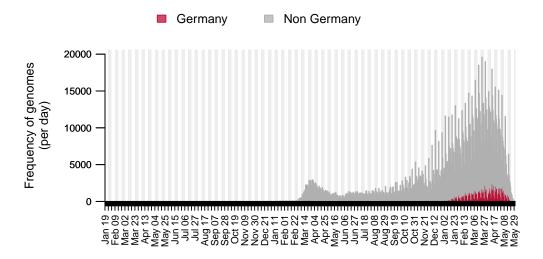


Figure 1: Collection dates of the nrow(metadata) genomes analysed here (left-hand axis). Genomes are coloured by sampling location.

#### ## #BE0F34

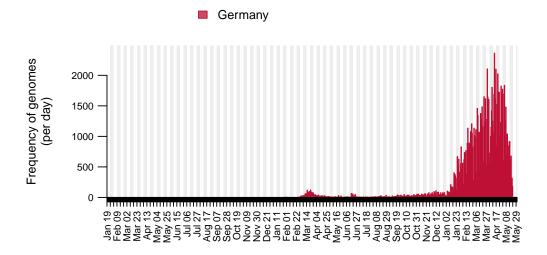


Figure 2: Collection dates of the nrow(metadata) genomes analysed here (left-hand axis). Genomes are coloured by sampling location.

# 3 Lineage importation distribution (shifted TMRCA distribution)

- GISAID tree until 2021-06-24 as initial tree.
- The tree is time-collibrated by TreeTime.
- Sankoff algorithm is used to assign location (Germany and non-Germany) to inner vertices of the tree, for each state sparately.
- Dataset contains Germany: 248 transmission lineages (2 or more sequences), comprising Germany: 32802 sequences from the Germany, as well as a further Germany: 88005 singletons.
- Mean and SD of the importation (shifted TMRCA) distribution: Germany : NA  $\pm$  NA days (singletons excluded).
- Median and interquartile range of the importation (shifted TMRCA) distribution Germany: NA [Germany: 2020-09-21, Germany: 2021-01-03] (singletons excluded).
- 80% of importations fall in [Germany: 2020-08-15, Germany: 2021-02-11].

Table 1: Estimated importation lags for multistate transmission lineages of different sizes. Importation lag is the waiting time between importation date and the TMRCA of the sampled genomes in the transmission lineage. Detection lag is the waiting time from the importation date to the sampling time of the oldest (first) sampled genome in the transmission lineage.

Lineages of size	No. of lineages of Germany	$\begin{array}{c} \text{Importation lag} \\ \text{(mean} \pm \text{SD)} \\ \text{Germany} \end{array}$	Importation lag (median and IQR) Germany	Detection lag (mean $\pm$ SD) Germany	Detection lag (median and IQR) Germany
All	248	$3.75 \pm 4.2$	1.79 [0.99-4.33]	$NA \pm NA$	NA [27-105]
2 to 10	80	$8.65 \pm 4.27$	7.95 [4.72-10.36]	$NA \pm NA$	NA [21.25-56.5]
11 to	102	$1.78 \pm 0.59$	1.64 [1.29-2.24]	$66.4 \pm 49.18$	55 [30-92]
100					
101 to	60	$0.86 \pm 0.08$	0.85 [0.8 - 0.94]	$96.28 \pm 51.68$	93 [60.75-130]
1000					
Bigger	6	$0.73 \pm 0$	0.74 [0.73 - 0.74]	$159.5 \pm 15.33$	160.5 [148-170.75]
than					
1000					

Table 2: 3. Estimated importation and detection lags for Germany transmission lineages ordered by importation date and aggregated by epi-week. Importation lag is the waiting time between importation date and the TMRCA of the sampled genomes in the transmission lineage. Detection lag is the waiting time from the importation date to the sampling time of the oldest (first) sampled genome in the transmission lineage. All statistics show means and standard deviations computed from the MCC trees.

Week		Estimated no. of		Importation lag	Detection lag
start-	Epi-	importations of	Lineage sizes (median	$(\text{mean} \pm \text{SD})$	$(\text{mean} \pm \text{SD})$
ing	week	Germany	and IQR) Germany	Germany	Germany
19 Jan 19	3	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jan	4	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
26 19 Feb	5	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
02 19 Feb	6	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
09 19 Feb	7	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
16 19 Feb	8	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
23 19 Mar	9	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
02 19 Mar	10	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
09 19 Mar	11	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
16 19 Mar	12	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
23 19 Mar	13	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
30 19 Apr	14	2	NA [NA-NA]	$NA \pm NA$	$\mathrm{NA}\pm\mathrm{NA}$
06 19 Apr	15	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
13 19 Apr	16	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 19 Apr	17	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
27 19 May	18	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
04 19 May	19	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
11 19 May	20	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
18					
19 May 25	21	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jun 01	22	2	NA [NA-NA]	$NA \pm NA$	NA ± NA
19 Jun 08	23	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$

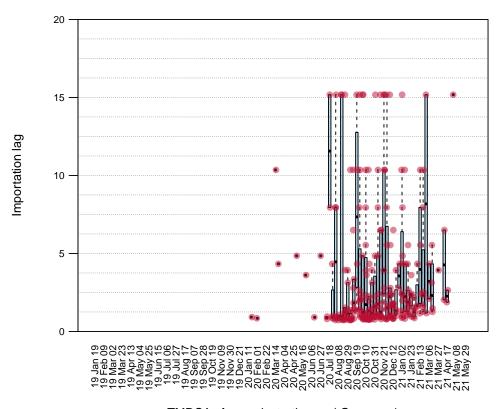
Week start-	Epi-	Estimated no. of importations of	Lineage sizes (median	Importation lag (mean $\pm$ SD)	Detection lag $(\text{mean} \pm \text{SD})$
ing	week	Germany	and IQR) Germany	Germany	Germany
19 Jun	24	2	NA [NA-NA]	$NA \pm NA$	NA ± NA
15 19 Jun 22	25	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jun 29	26	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jul 06	27	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jul 13	28	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 Jul	29	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 19 Jul	30	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
27 19 Aug	31	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
03 19 Aug	32	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
10 19 Aug	33	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
17 19 Aug	34	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
24 19 Aug	35	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
31 19 Sep	36	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
07 19 Sep	37	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
14 19 Sep	38	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 19 Sep	39	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
28 19 Oct	40	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
05 19 Oct	41	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
12 19 Oct	42	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
19 19 Oct	43	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
26 19 Nov	44	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
02 19 Nov	45	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
09 19 Nov	46	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
16 19 Nov	47	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
23 19 Nov 30	48	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$

Week		Estimated no. of		Importation lag	Detection lag
start-	Epi-	importations of	Lineage sizes (median	$(\text{mean} \pm \text{SD})$	$(\text{mean} \pm \text{SD})$
ing	week	Germany	and IQR) Germany	Germany	Germany
19 Dec 07	49	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
$19  \mathrm{Dec}$	50	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
14 19 Dec	51	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 19 Dec	0	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
28 20 Jan 04	1	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Jan 11	2	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Jan 18	3	3	NA [144-144]	$NA \pm NA$	$NA \pm NA$
20 Jan 25	4	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Feb 01	5	3	NA [229-229]	$NA \pm NA$	$NA \pm NA$
20 Feb 08	6	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Feb 15	7	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Feb 22	8	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Feb 29	9	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Mar 07	10	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Mar 14	11	3	NA [3-3]	$NA \pm NA$	$NA \pm NA$
20 Mar 21	12	3	NA [8-8]	$NA \pm NA$	$NA \pm NA$
20 Mar 28	13	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Apr 04	14	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Apr 11	15	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Apr 18	16	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Apr 25	17	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 May 02	18	3	NA [7-7]	$NA \pm NA$	$NA \pm NA$
20 May	19	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
09 20 May	20	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
16 20 May 23	21	3	NA [10-10]	$NA \pm NA$	$NA \pm NA$

Week start-	Epi-	Estimated no. of importations of	Lineage sizes (median	Importation lag (mean $\pm$ SD)	Detection lag (mean $\pm$ SD)
ing	week	Germany	and IQR) Germany	Germany	Germany
20 May 30	22	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Jun 06	23	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Jun 13	24	3	NA [153-153]	$NA \pm NA$	$NA \pm NA$
20 Jun 20	25	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
20 Jun 27	26	3	NA [7-7]	$NA \pm NA$	$NA \pm NA$
20  Jul	27	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
04 20 Jul	28	4	NA [178-290]	$NA \pm NA$	$NA \pm NA$
11 20 Jul	29	4	NA [2.5-3.5]	$NA \pm NA$	$NA \pm NA$
18 20 Jul	30	6	NA [81.5-198]	$NA \pm NA$	$NA \pm NA$
25 20 Aug	31	8	NA [4-209.25]	$NA \pm NA$	$NA \pm NA$
01 20 Aug	32	4	NA [728.25-2012.75]	$NA \pm NA$	$NA \pm NA$
08 20 Aug	33	7	NA [2-378]	$NA \pm NA$	$NA \pm NA$
15 20 Aug	34	11	NA [36-363]	$NA \pm NA$	$NA \pm NA$
22 20 Aug	35	8	NA [20.25-199]	$NA \pm NA$	$NA \pm NA$
29 20 Sep	36	9	NA [92-968.5]	$NA \pm NA$	$NA \pm NA$
05 20 Sep	37	11	NA [11-45]	$NA \pm NA$	$NA \pm NA$
12 20 Sep	38	6	NA [2.75-18]	$NA \pm NA$	$NA \pm NA$
19 20 Sep	39	14	NA [12.25-41.75]	$NA \pm NA$	$NA \pm NA$
26 20 Oct	40	11	NA [7-216]	$NA \pm NA$	$NA \pm NA$
03 20 Oct	41	14	NA [8.25-144.25]	$NA \pm NA$	$NA \pm NA$
10 20 Oct	42	13	NA [21.5-448.5]	$NA \pm NA$	$NA \pm NA$
17 20 Oct	43	8	NA [20.25-163.5]	$NA \pm NA$	$NA \pm NA$
24 20 Oct	44	10	NA [16-81.25]	$NA \pm NA$	$NA \pm NA$
31 20 Nov	45	12	NA [9.5-79.5]	$NA \pm NA$	$NA \pm NA$
07 20 Nov 14	46	10	NA [5-127]	$NA \pm NA$	$NA \pm NA$

Week start-	Epi-	Estimated no. of importations of	Lineage sizes (median	Importation lag (mean ± SD)	Detection lag (mean ± SD)
ing	week	Germany	and IQR) Germany	Germany	Germany
20 Nov 21	47	11	NA [3-17]	$NA \pm NA$	$NA \pm NA$
20 Nov 28	48	10	NA [5.5-239.75]	$NA \pm NA$	$NA \pm NA$
20 Dec 05	49	11	NA [14-75]	$NA \pm NA$	$NA \pm NA$
20 Dec 12	50	12	NA [26.5-219.75]	$NA \pm NA$	$NA \pm NA$
20 Dec 19	51	6	NA [33.75-67]	$NA \pm NA$	$NA \pm NA$
20 Dec 26	0	4	NA [9.5-12.5]	$NA \pm NA$	$NA \pm NA$
21 Jan 02	1	14	NA [6.25-55.75]	$NA \pm NA$	$NA \pm NA$
21 Jan 09	2	7	NA [8-51]	$NA \pm NA$	$NA \pm NA$
21 Jan 16	3	11	NA [15-50]	$NA \pm NA$	$NA \pm NA$
21 Jan 23	4	7	NA [17-31]	$NA \pm NA$	$NA \pm NA$
21 Jan 30	5	7	NA [39-122]	$NA \pm NA$	$NA \pm NA$
21 Feb 06	6	5	NA [19.5-32]	$NA \pm NA$	$NA \pm NA$
21 Feb 13	7	6	NA [5.25-29]	$NA \pm NA$	$NA \pm NA$
21 Feb 20	8	9	NA [10-31.5]	$NA \pm NA$	$NA \pm NA$
21 Feb 27	9	4	NA [17.25-47.75]	$NA \pm NA$	$NA \pm NA$
21 Mar 06	10	4	NA [11.25-17.75]	$NA \pm NA$	$NA \pm NA$
21 Mar 13	11	6	NA [10.5-42]	$NA \pm NA$	$NA \pm NA$
21 Mar 20	12	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 Mar 27	13	3	NA [9-9]	$NA \pm NA$	$NA \pm NA$
21 Apr 03	14	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
$21~\mathrm{Apr}$	15	4	NA [9.25-17.75]	$NA \pm NA$	$NA \pm NA$
10 21 Apr	16	4	NA [17.5-22.5]	$NA \pm NA$	$NA \pm NA$
17 21 Apr	17	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
24 21 May	18	3	NA [2-2]	$NA \pm NA$	$NA \pm NA$
01 21 May 08	19	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$

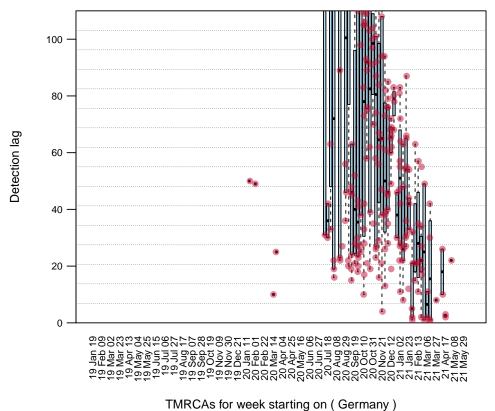
Week start-	Epi-	Estimated no. of importations of	Lineage sizes (median	Importation lag (mean $\pm$ SD)	Detection lag (mean $\pm$ SD)
ing	week	Germany	and IQR) Germany	Germany	Germany
21 May 15	20	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 May 22	21	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 May 29	22	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 Jun 05	23	2	NA [NA-NA]	$NA \pm NA$	$NA \pm NA$
21 Jun 12	24	2	NA [NA-NA]	NA ± NA	NA ± NA



TMRCAs for week starting on ( Germany )

Figure 3: Boxplots of the estimated importation lags for  ${\tt r}$  states transmission lineages ordered by importation date and aggregated by epi-week.

```
## Error in seq.int(0, to0 - from, by): 'to' must be a finite number
## Error in seq.int(0, to0 - from, by): 'to' must be a finite number
## Error in seq.int(0, to0 - from, by): 'to' must be a finite number
```



TWINGAS TO WEEK Starting On ( Germany )

Figure 4: Boxplots of the estimated detection lags for  ${\tt r}$  states transmission lineages ordered by importation date and aggregated by epi-week.

## 4 Session info

```
## R version 4.1.0 (2021-05-18)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Ubuntu 20.04.2 LTS
## Matrix products: default
## BLAS:
           /usr/lib/x86_64-linux-gnu/atlas/libblas.so.3.10.3
## LAPACK: /usr/lib/x86 64-linux-gnu/atlas/liblapack.so.3.10.3
##
## locale:
  [1] LC CTYPE=en US.UTF-8
                                   LC NUMERIC=C
  [3] LC_TIME=en_US.UTF-8
                                   LC_COLLATE=en_US.UTF-8
   [5] LC_MONETARY=en_US.UTF-8
                                   LC_MESSAGES=en_US.UTF-8
##
  [7] LC_PAPER=en_US.UTF-8
                                   LC NAME=C
## [9] LC ADDRESS=C
                                   LC TELEPHONE=C
## [11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
## attached base packages:
## [1] stats
                 graphics grDevices utils
                                                datasets methods
                                                                    base
## other attached packages:
  [1] phytools_0.7-70 maps_3.3.0
                                           ape_5.5
                                                            ggtree_3.0.1
  [5] ggsci_2.9
                         ggplot2_3.3.3
                                           stringr_1.4.0
                                                            tictoc_1.0.1
## [9] knitr_1.33
                                           gplots_3.1.1
                         beastio_0.3.3
                                                            plyr_1.8.6
## [13] lubridate_1.7.10
##
## loaded via a namespace (and not attached):
## [1] Rcpp 1.0.6
                                lattice 0.20-44
                                                         tidyr_1.1.3
## [4] gtools_3.8.2
                                digest_0.6.27
                                                         utf8_1.2.1
## [7] R6_2.5.0
                                evaluate_0.14
                                                         coda_0.19-4
## [10] highr_0.9
                                                         rlang_0.4.11
                                pillar_1.6.1
## [13] lazyeval_0.2.2
                                phangorn_2.7.0
                                                         Matrix_1.3-4
## [16] combinat_0.0-8
                                rmarkdown_2.8
                                                         igraph_1.2.6
## [19] munsell_0.5.0
                                numDeriv_2016.8-1.1
                                                         compiler_4.1.0
                                pkgconfig_2.0.3
                                                         mnormt_2.0.2
## [22] xfun_0.23
                                                         tidyselect_1.1.1
## [25] tmvnsim_1.0-2
                                htmltools_0.5.1.1
## [28] tibble_3.1.2
                                expm_0.999-6
                                                         codetools_0.2-18
## [31] quadprog_1.5-8
                                fansi_0.5.0
                                                         crayon_1.4.1
                                                         MASS_7.3-54
## [34] dplyr_1.0.6
                                withr_2.4.2
## [37] bitops_1.0-7
                                grid_4.1.0
                                                         nlme_3.1-152
## [40] jsonlite_1.7.2
                                gtable_0.3.0
                                                         lifecycle_1.0.0
## [43] magrittr_2.0.1
                                scales_1.1.1
                                                         KernSmooth 2.23-20
## [46] tidytree 0.3.4
                                stringi 1.6.2
                                                         scatterplot3d 0.3-41
## [49] ellipsis_0.3.2
                                rvcheck_0.1.8
                                                         generics_0.1.0
## [52] vctrs 0.3.8
                                fastmatch 1.1-0
                                                         RColorBrewer 1.1-2
## [55] tools_4.1.0
                                treeio_1.16.1
                                                         glue_1.4.2
## [58] purrr_0.3.4
                                plotrix_3.8-1
                                                         parallel_4.1.0
## [61] yaml_2.2.1
                                colorspace_2.0-1
                                                         BiocManager_1.30.15
## [64] caTools_1.18.2
                                aplot_0.0.6
                                                         clusterGeneration_1.3.7
## [67] patchwork_1.1.1
```