

Supplemental information

**Early detection and improved
genomic surveillance of SARS-CoV-2
variants from deep sequencing data**

Daniele Ramazzotti, Davide Maspero, Fabrizio Angaroni, Silvia Spinelli, Marco Antoniotti, Rocco Piazza, and Alex Graudenzi

Supplementary Information: Early detection and improved genomic surveillance of SARS-CoV-2 variants from deep sequencing data

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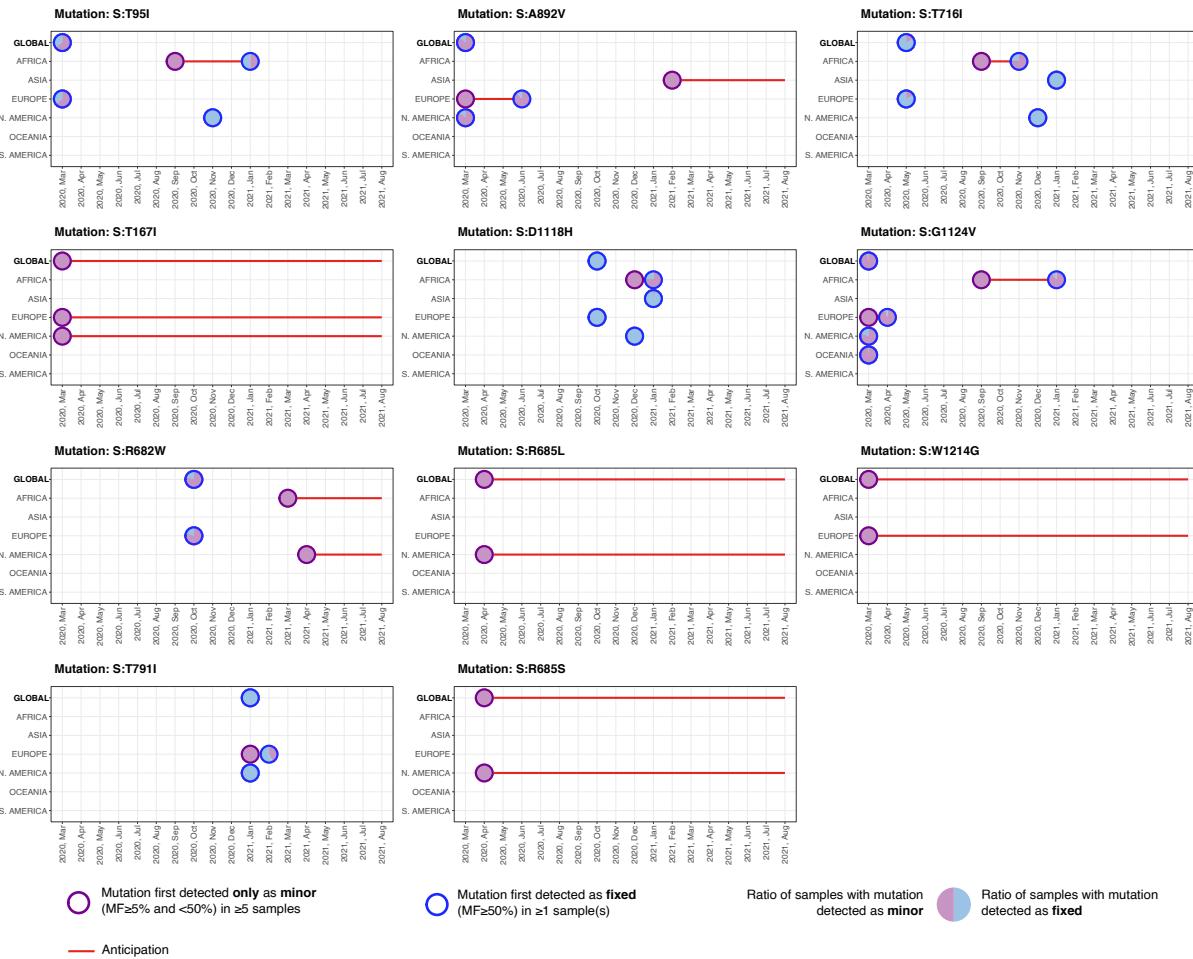


Figure S 1: Early detection of 11 highly diffused S mutations from deep sequencing data.

Related to Figures 2, 4, 5. Analysis of S mutations: S:T95I, S:T167I, S:R682W, S:R685L, S:R685S, S:T716I, S:T791I, S:A892V, S:D1118H, S:G1124V, S:W1214G. Circles with *purple* borders mark the first month in which the mutation was detected as minor ($MF \geq 5\%$ and $< 50\%$) in at least 5 samples, while been still undetected as fixed ($MF \geq 50\%$); circles with *blue* borders mark the month in which the mutation was first detected as fixed in at least 1 sample; red lines highlight the anticipation (when > 1 months). The analysis is performed by splitting the samples in the 6 distinct geographical regions and by reporting the corresponding results at the global scale. All circles contain a pie-chart that displays the ratio of samples showing that mutation either as minor or fixed in that month (further details are provided in the main text). For each SMoI the related variants are also reported.

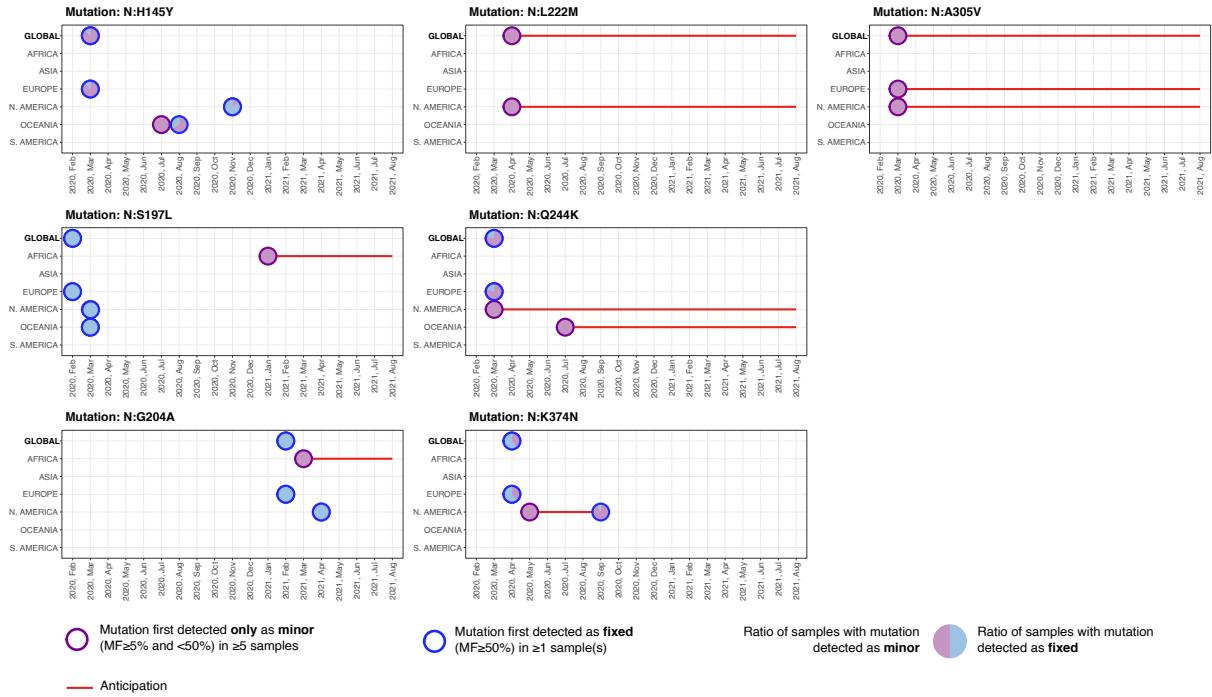


Figure S 2: Early detection of 7 highly diffused N mutations from deep sequencing data. Related to Figures 2, 4, 5. Analysis of N mutations: N:H145Y, N:S197L, N:G204A, N:L222M, N:Q244K, N:A305V, N:K374N. Circles with *purple* borders mark the first month in which the mutation was detected as minor ($MF \geq 5\%$ and $< 50\%$) in at least 5 samples, while been still undetected as fixed ($MF \geq 50\%$); circles with *blue* borders mark the month in which the mutation was first detected as fixed in at least 1 sample; red lines highlight the anticipation (when > 1 months). The analysis is performed by splitting the samples in the 6 distinct geographical regions and by reporting the corresponding results at the global scale. All circles contain a pie-chart that displays the ratio of samples showing that mutation either as minor or fixed in that month (further details are provided in the main text). For each SMoI the related variants are also reported.



Figure S 3: Mutant frequency and prevalence variation in time of mutation: S:A701V.

Related to Figure 3. The leftmost panels return the distribution of the mutation frequency (MF) of all samples with the mutation, grouped by month and geographical region. Each cell shows the proportion of samples showing the mutation with that specific MF. The rightmost panels show the number of samples showing the mutations either as minor ($MF \geq 5\%$) and $< 50\%$) or fixed ($MF \geq 50\%$).

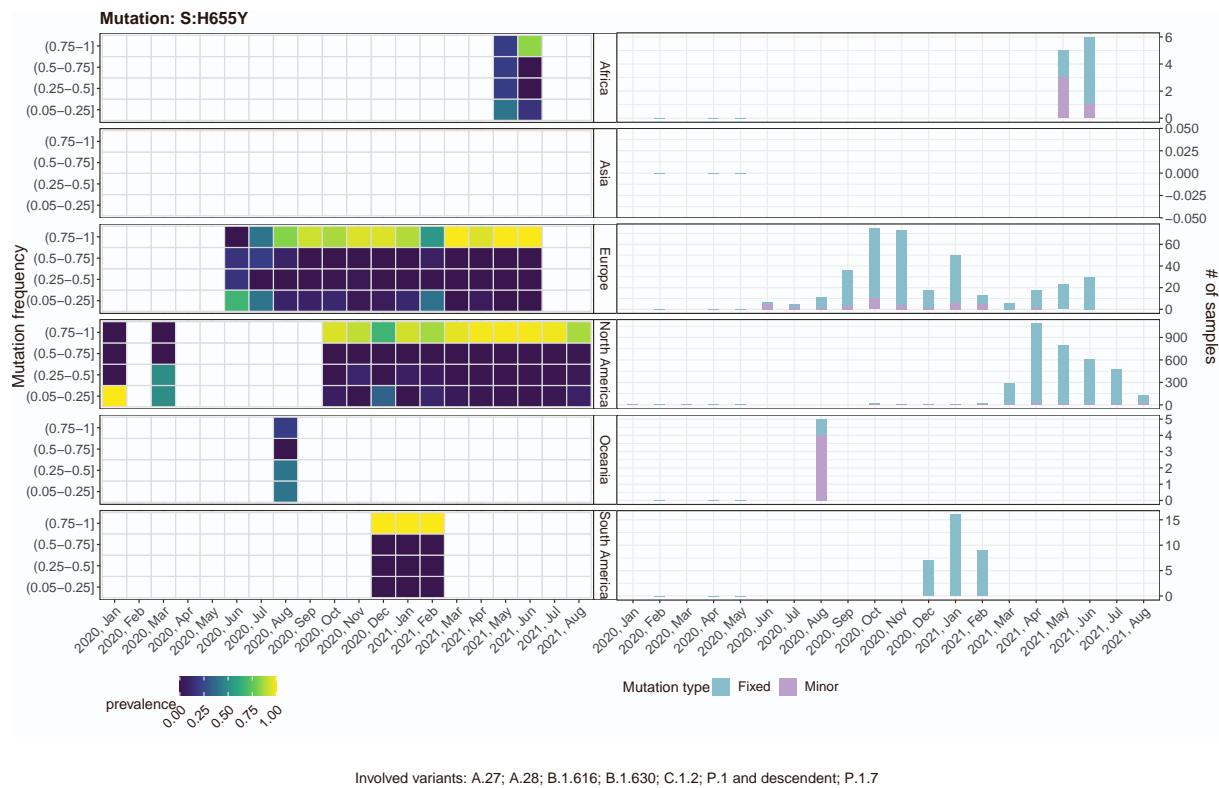


Figure S 4: Mutant frequency and prevalence variation in time of mutation: S:H655Y.
Related to Figure 3. Please refer to the caption of Figure S 3

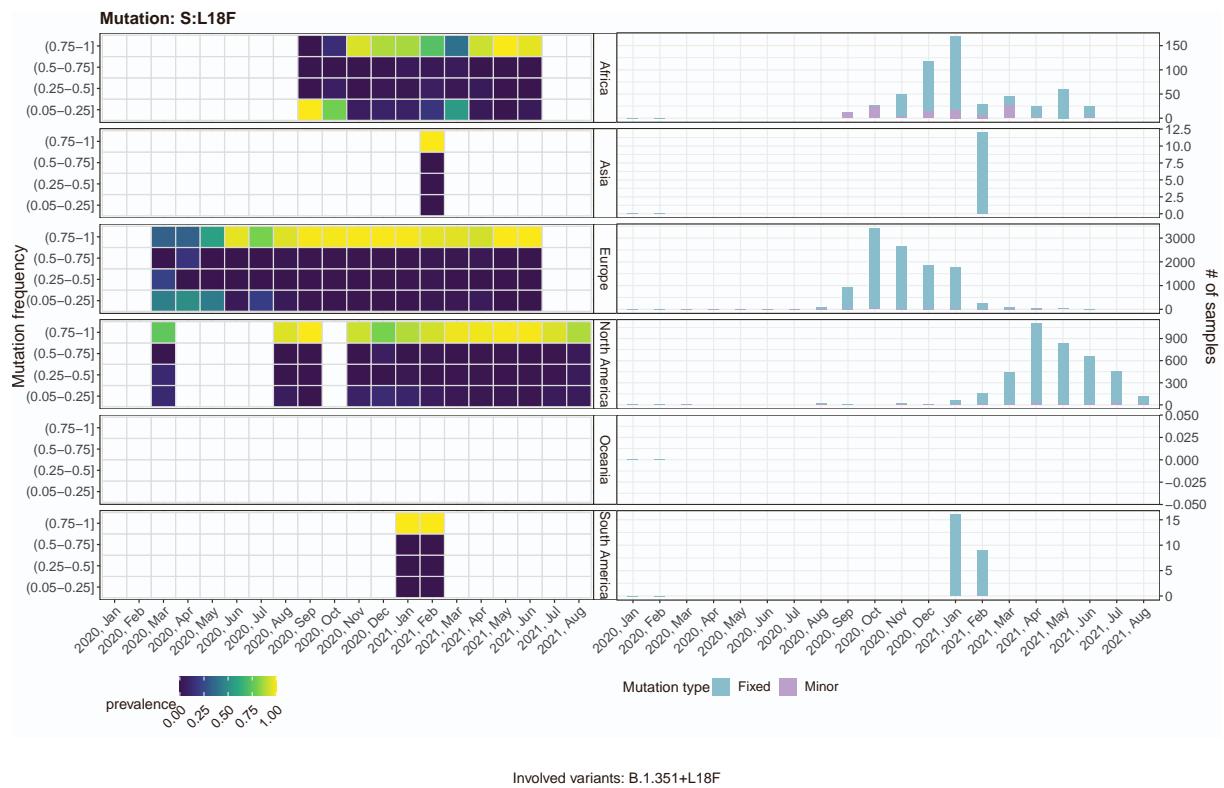


Figure S 5: **Mutant frequency and prevalence variation in time of mutation: S:L18F. Related to Figure 3.** Please refer to the caption of Figure S 3

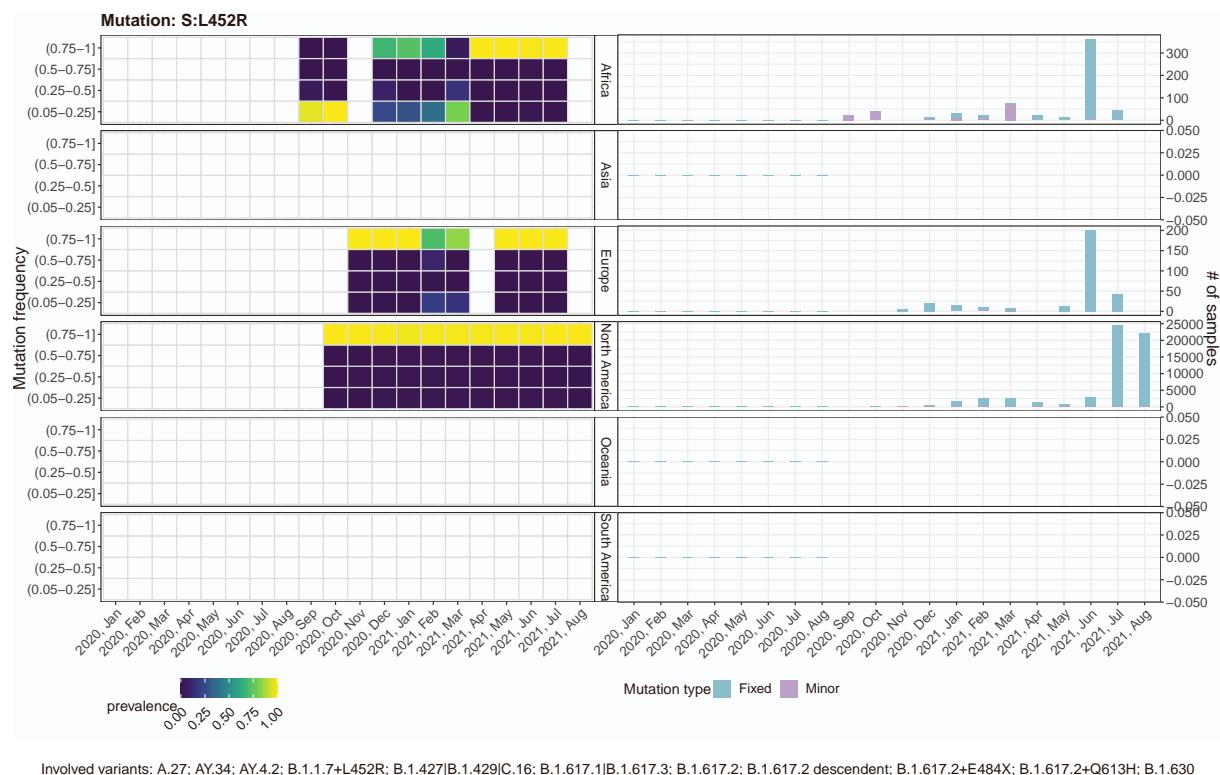


Figure S 6: Mutant frequency and prevalence variation in time of mutation: S:L452R.
Related to Figure 3. Please refer to the caption of Figure S 3

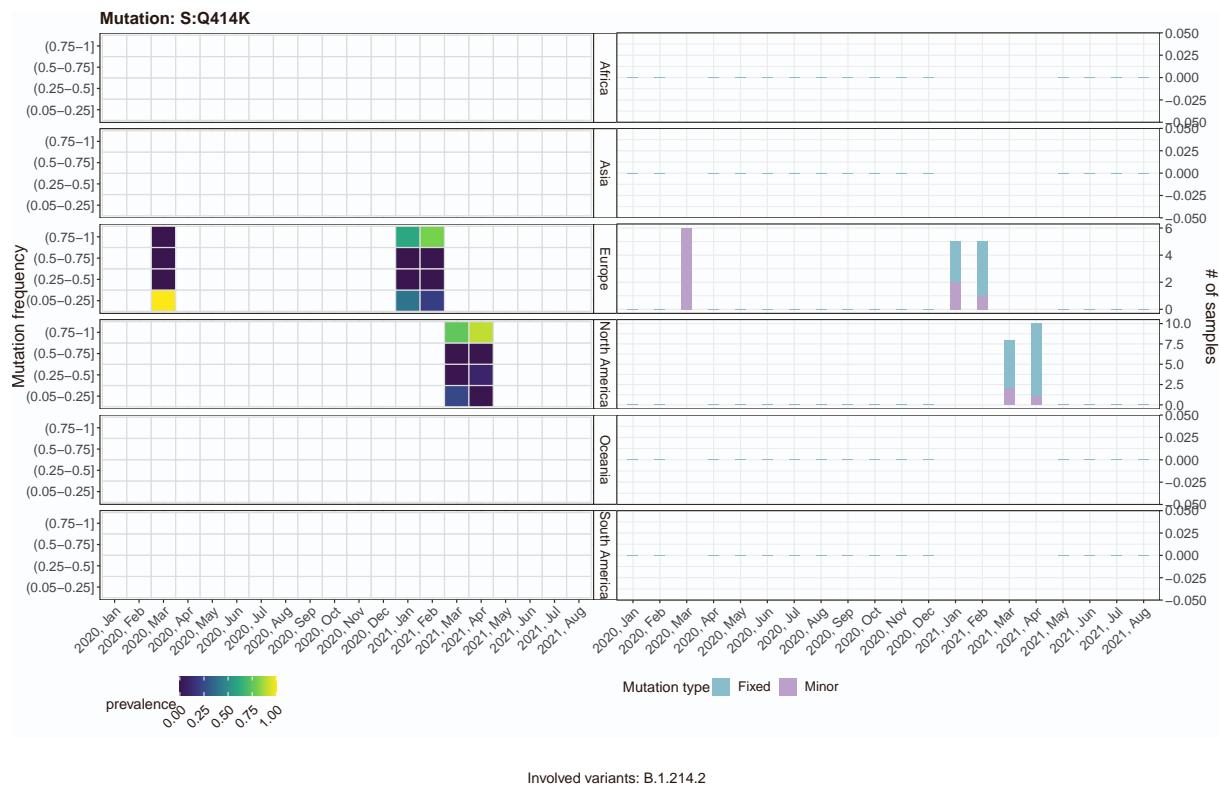


Figure S 7: Mutant frequency and prevalence variation in time of mutation: S:Q414K.
Related to Figure 3. Please refer to the caption of Figure S 3



Figure S 8: Mutant frequency and prevalence variation in time of mutation: S:T478K.
Related to Figure 3. Please refer to the caption of Figure S 3

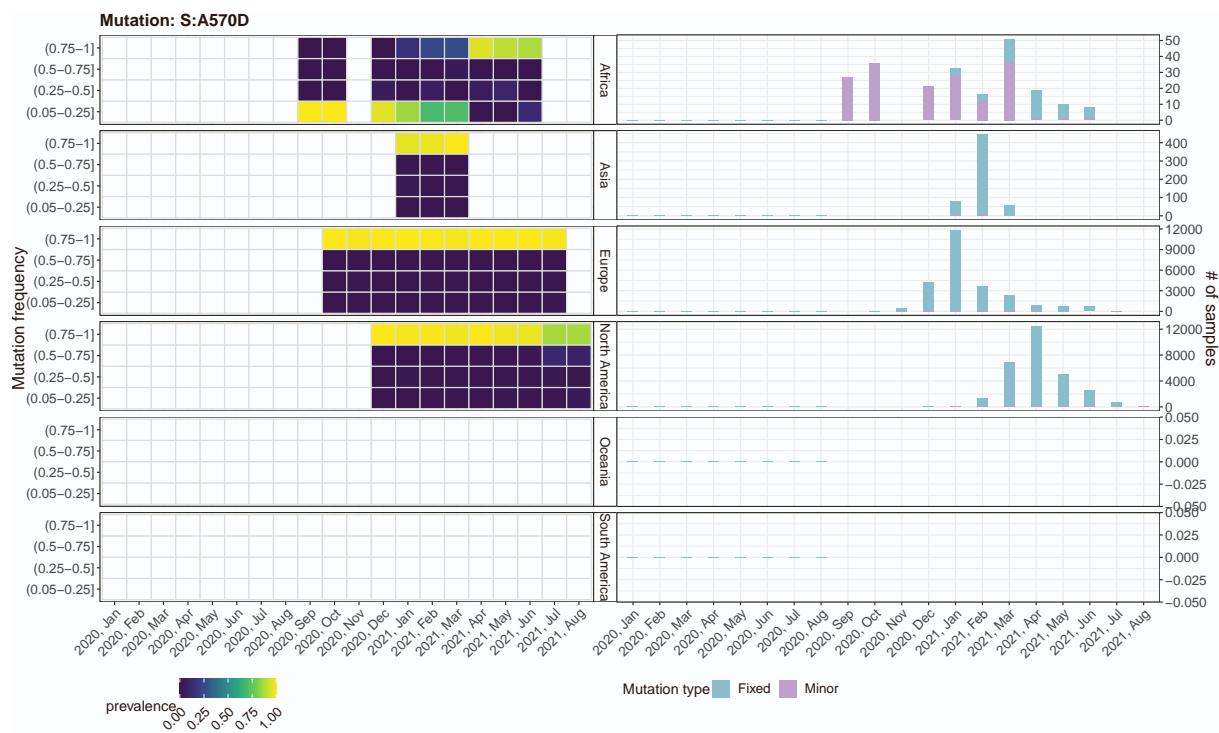


Figure S 9: Mutant frequency and prevalence variation in time of mutation: S:A570D. Related to Figure 3. Please refer to the caption of Figure S 3

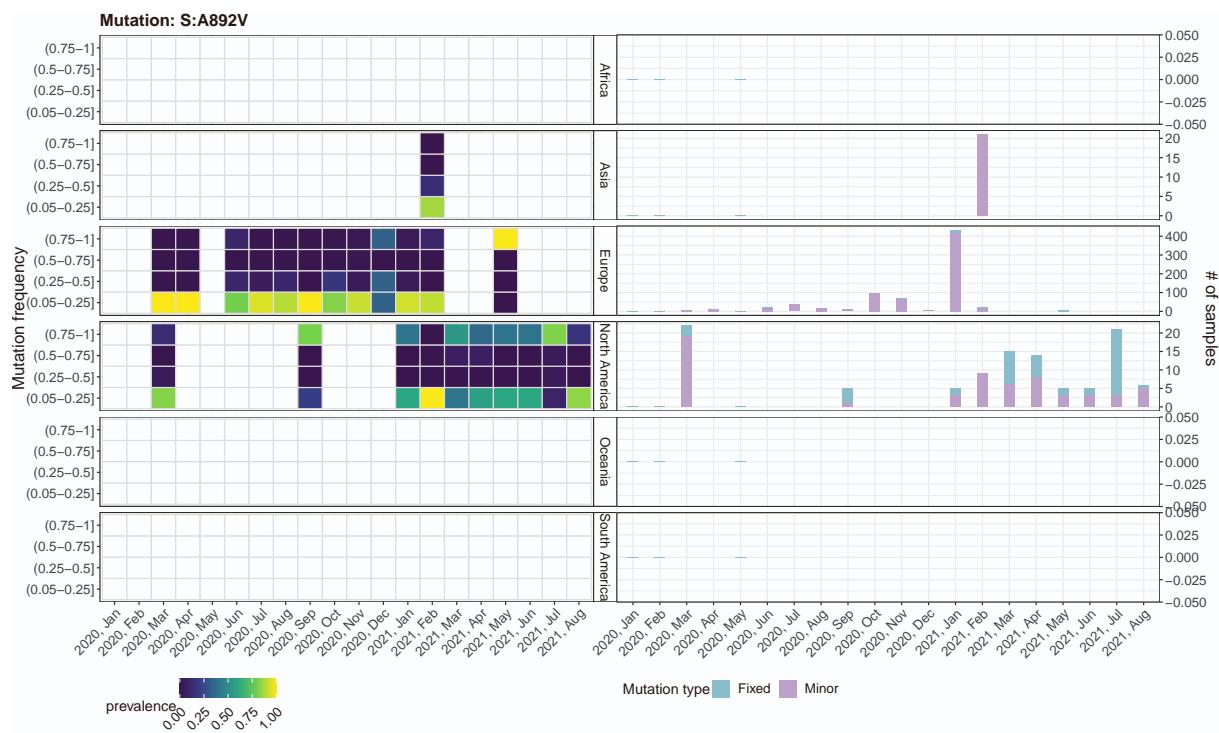


Figure S 10: Mutant frequency and prevalence variation in time of mutation: S:A892V.
Related to Figure 3. Please refer to the caption of Figure S 3

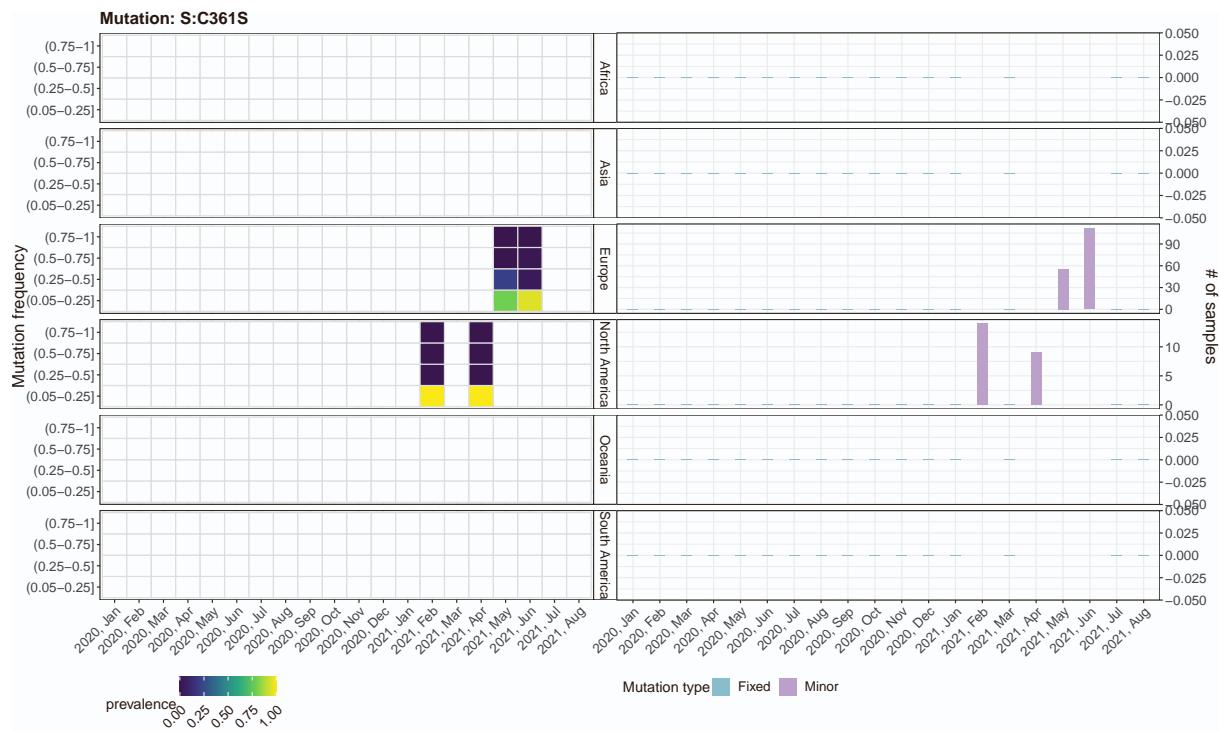


Figure S 11: Mutant frequency and prevalence variation in time of mutation: S:C361S.
Related to Figure 3. Please refer to the caption of Figure S 3

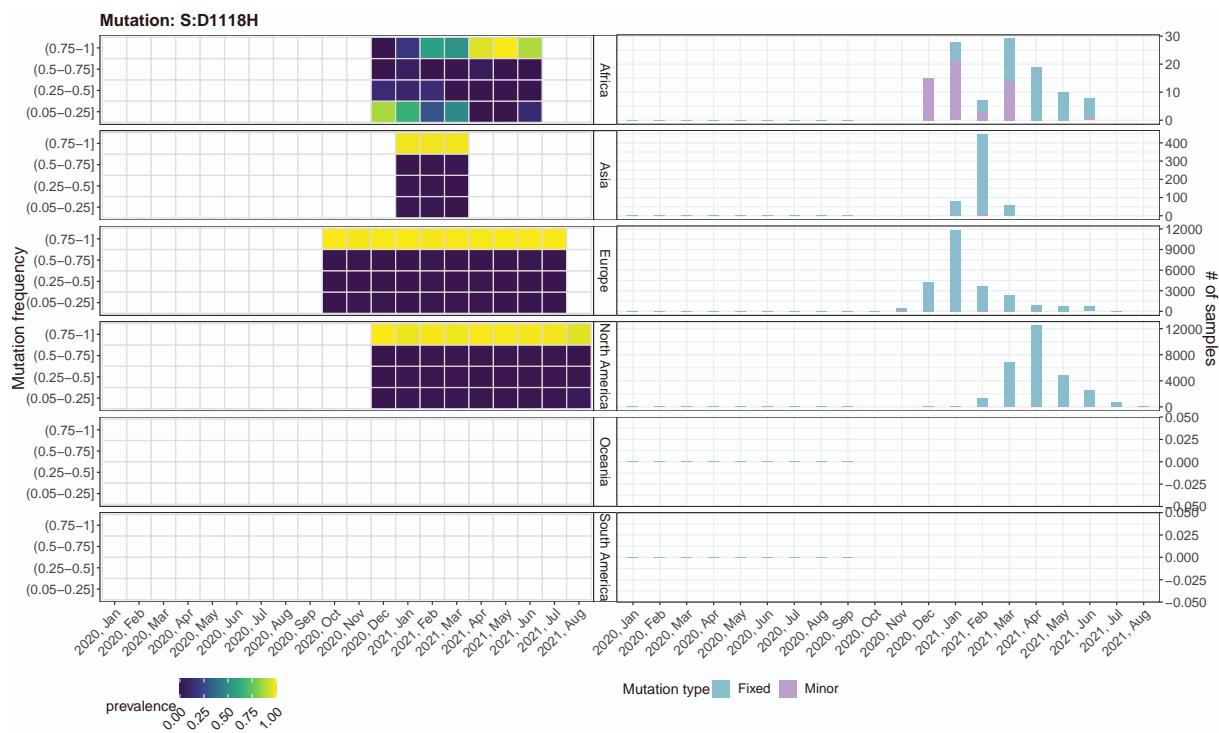


Figure S 12: Mutant frequency and prevalence variation in time of mutation: S:D1118H.
Related to Figure 3. Please refer to the caption of Figure S 3

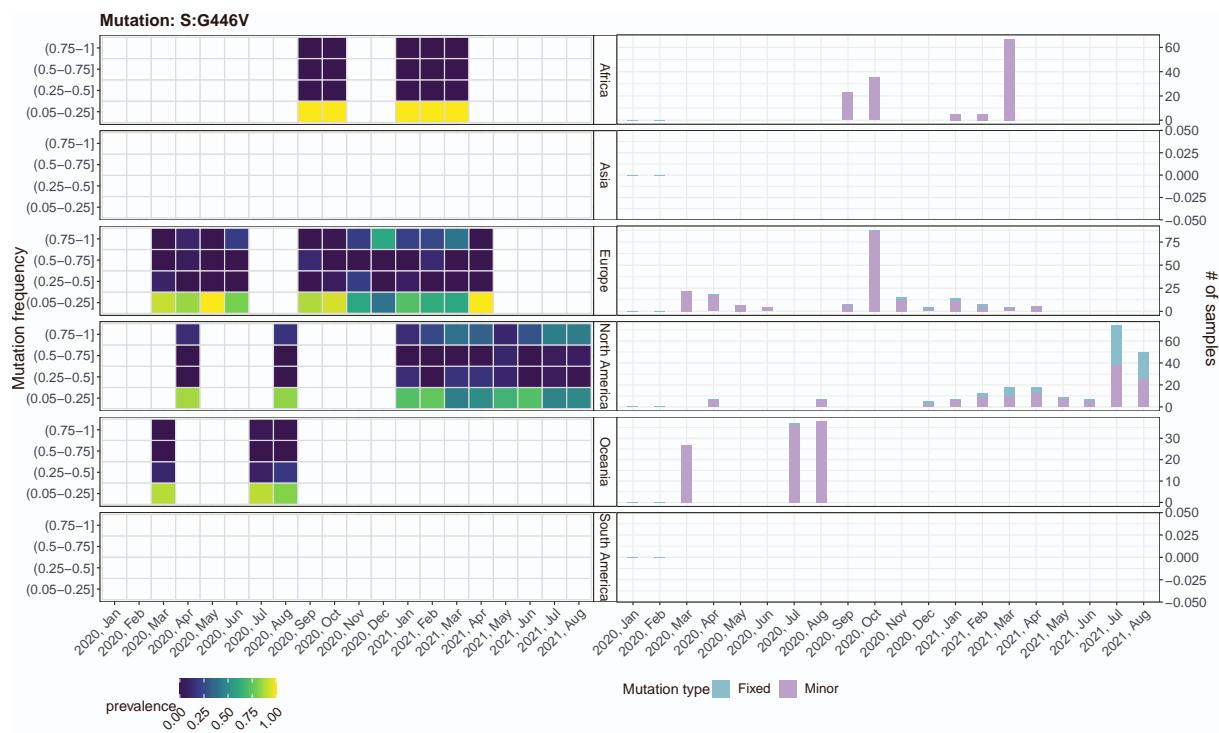


Figure S 13: Mutant frequency and prevalence variation in time of mutation: S:G446V.
Related to Figure 3. Please refer to the caption of Figure S 3

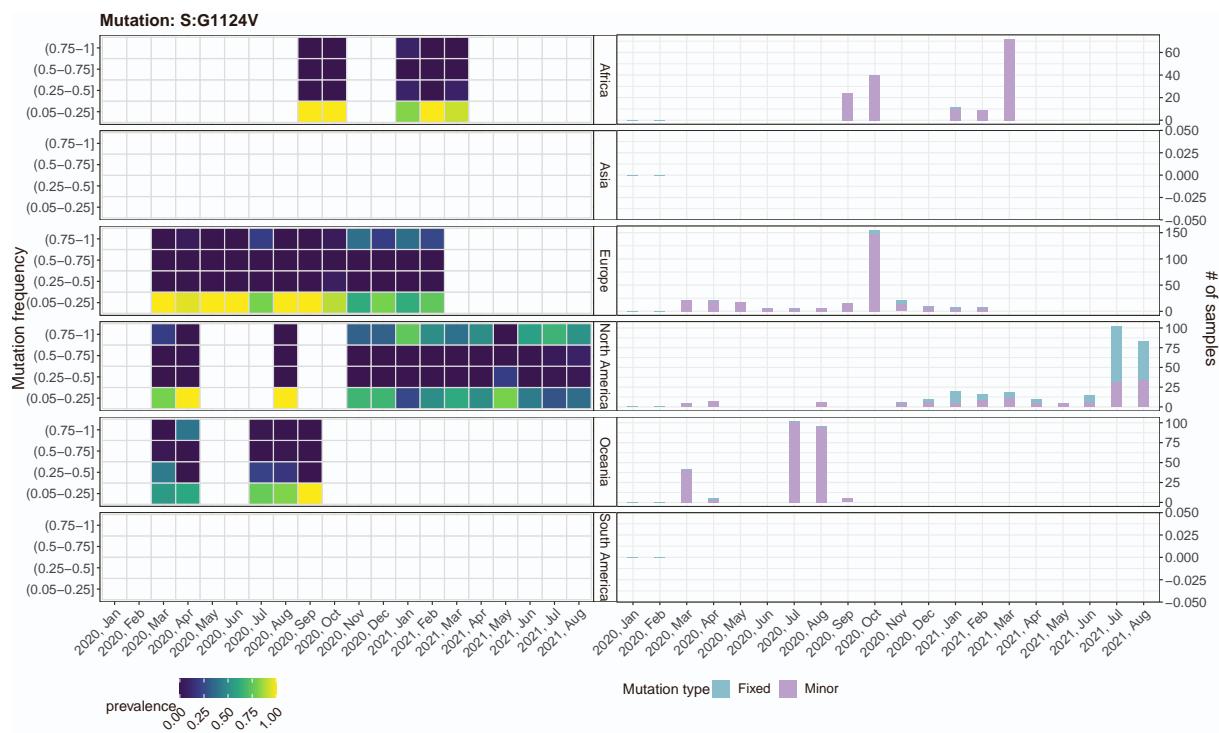


Figure S 14: Mutant frequency and prevalence variation in time of mutation: S:G1124V. Related to Figure 3. Please refer to the caption of Figure S 3

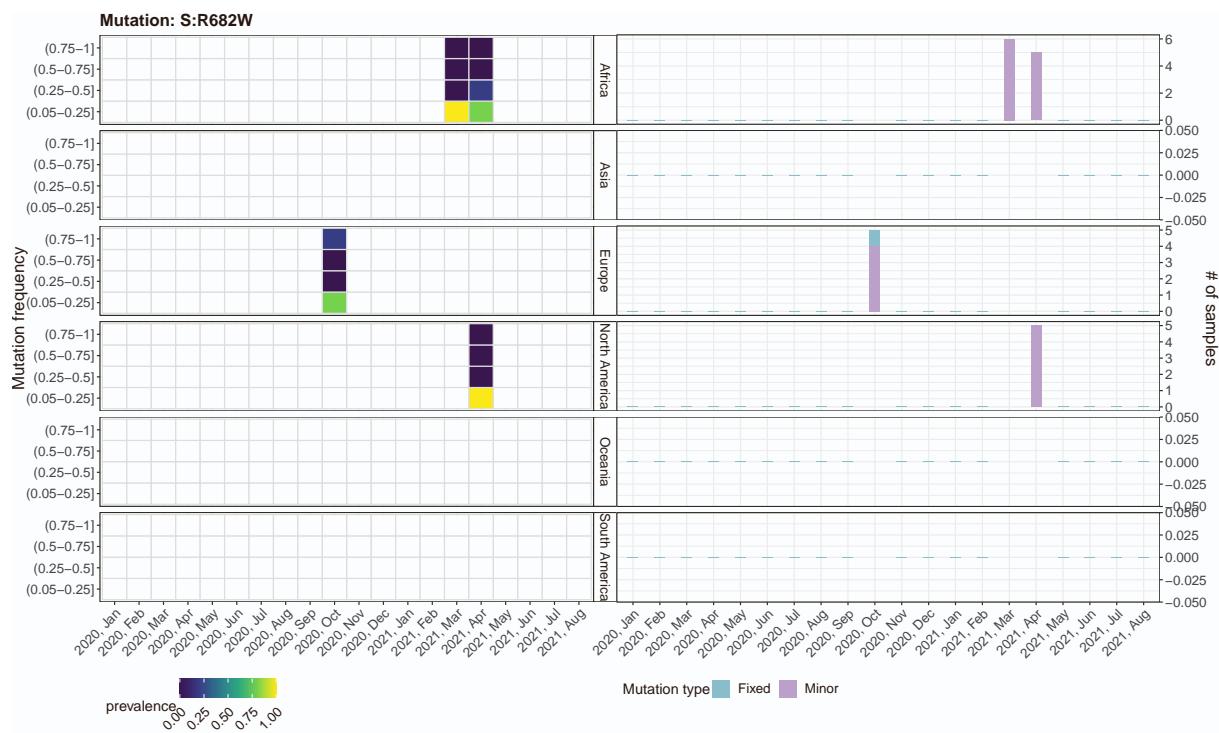


Figure S 15: Mutant frequency and prevalence variation in time of mutation: S:R682W.
Related to Figure 3. Please refer to the caption of Figure S 3

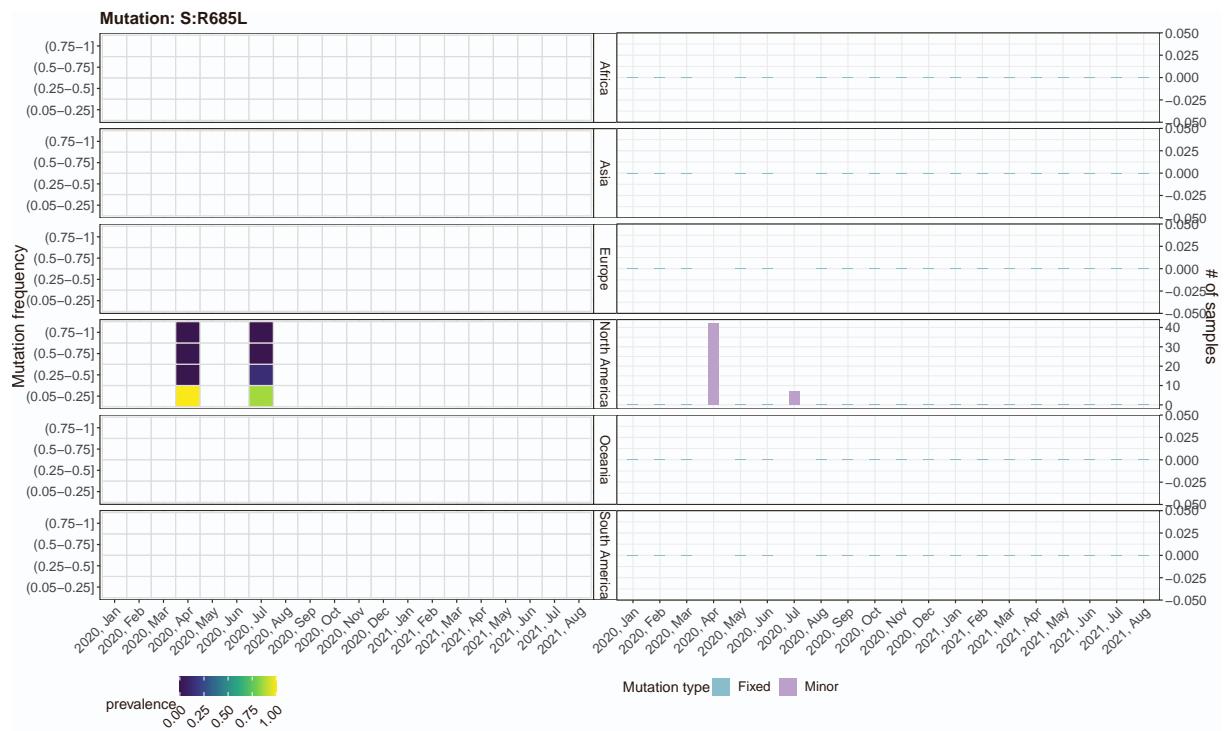


Figure S 16: Mutant frequency and prevalence variation in time of mutation: S:R685L.
Related to Figure 3. Please refer to the caption of Figure S 3

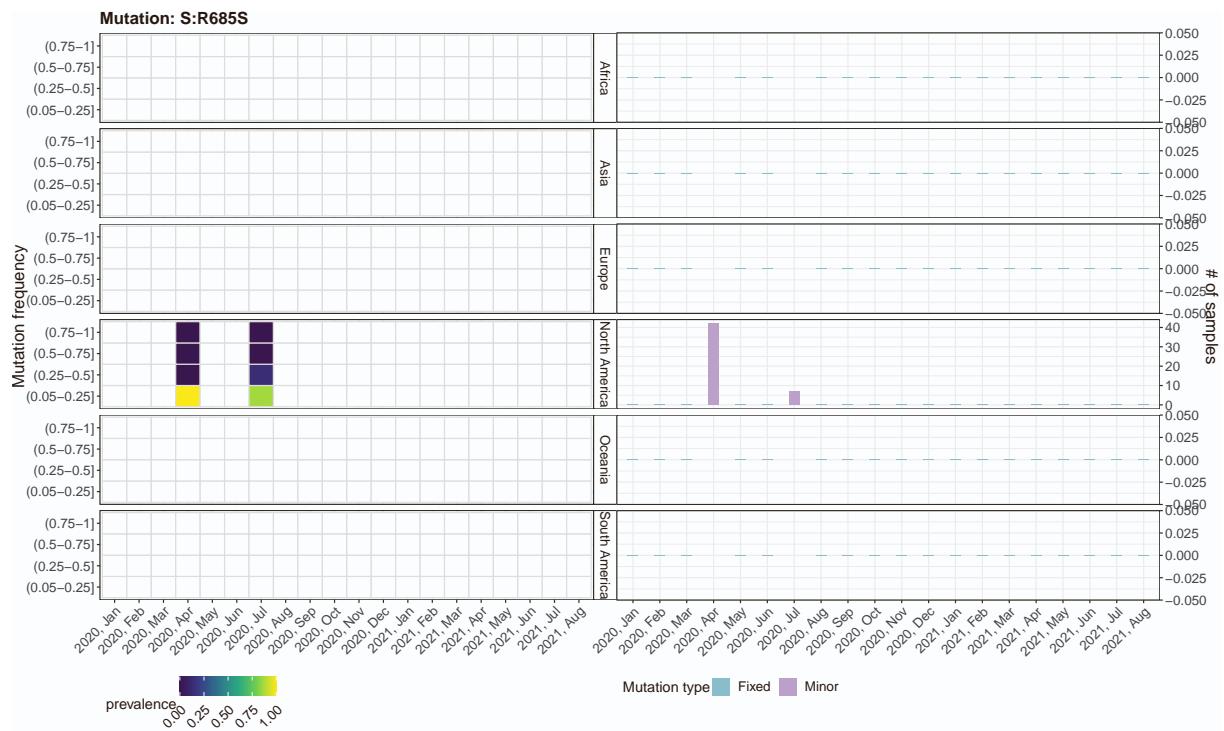


Figure S 17: Mutant frequency and prevalence variation in time of mutation: S:R685S.
Related to Figure 3. Please refer to the caption of Figure S 3



Figure S 18: Mutant frequency and prevalence variation in time of mutation: S:S297L. Related to Figure 3. Please refer to the caption of Figure S 3

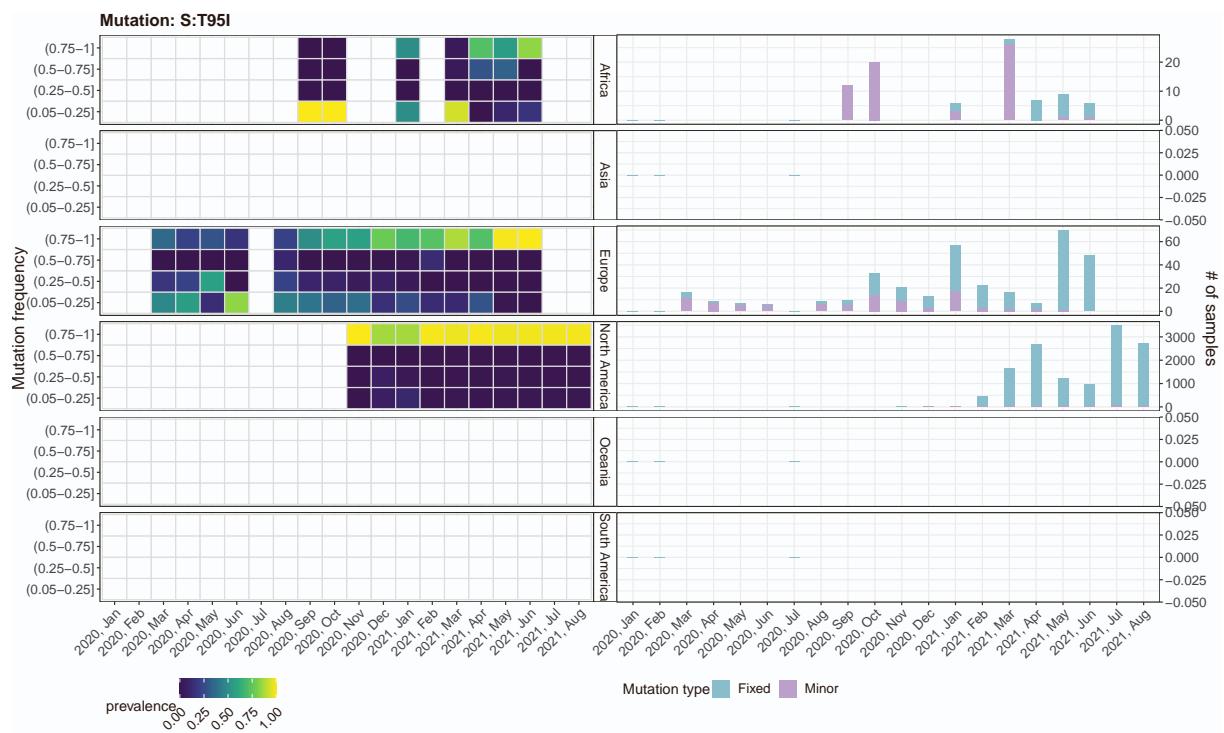


Figure S 19: Mutant frequency and prevalence variation in time of mutation: S:T95I.
Related to Figure 3. Please refer to the caption of Figure S 3

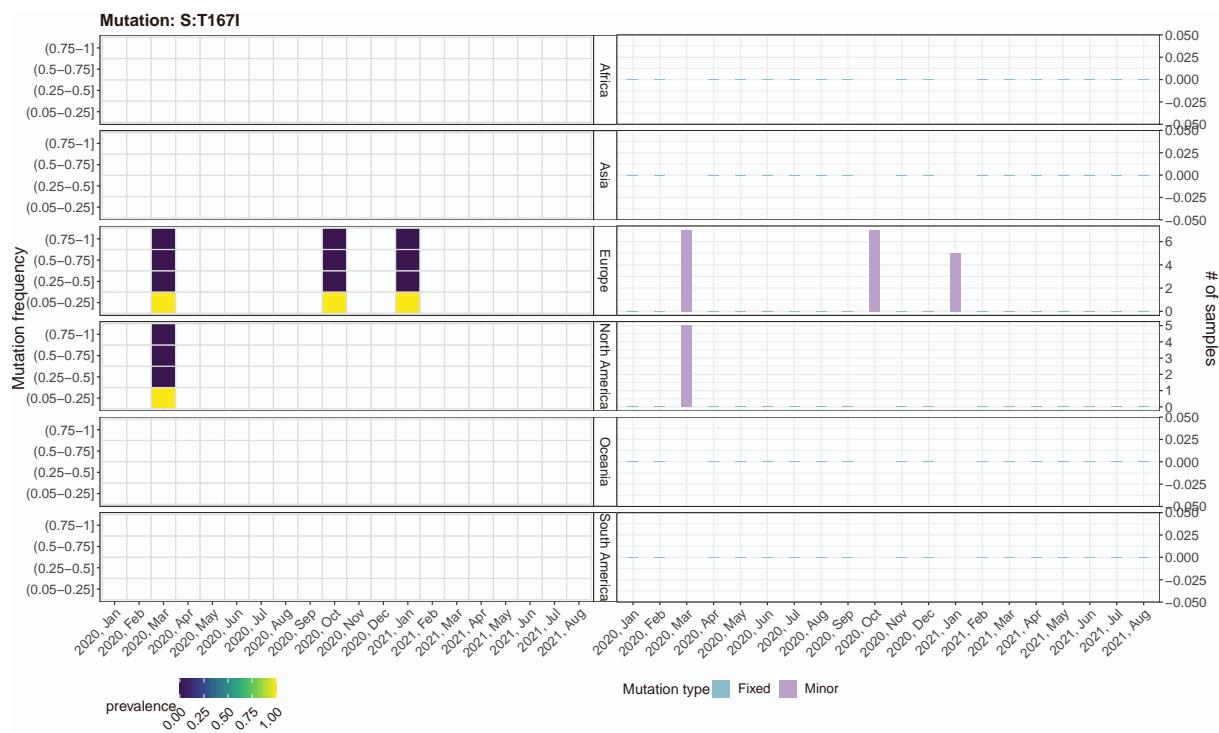


Figure S 20: Mutant frequency and prevalence variation in time of mutation: S:T167I.
Related to Figure 3. Please refer to the caption of Figure S 3

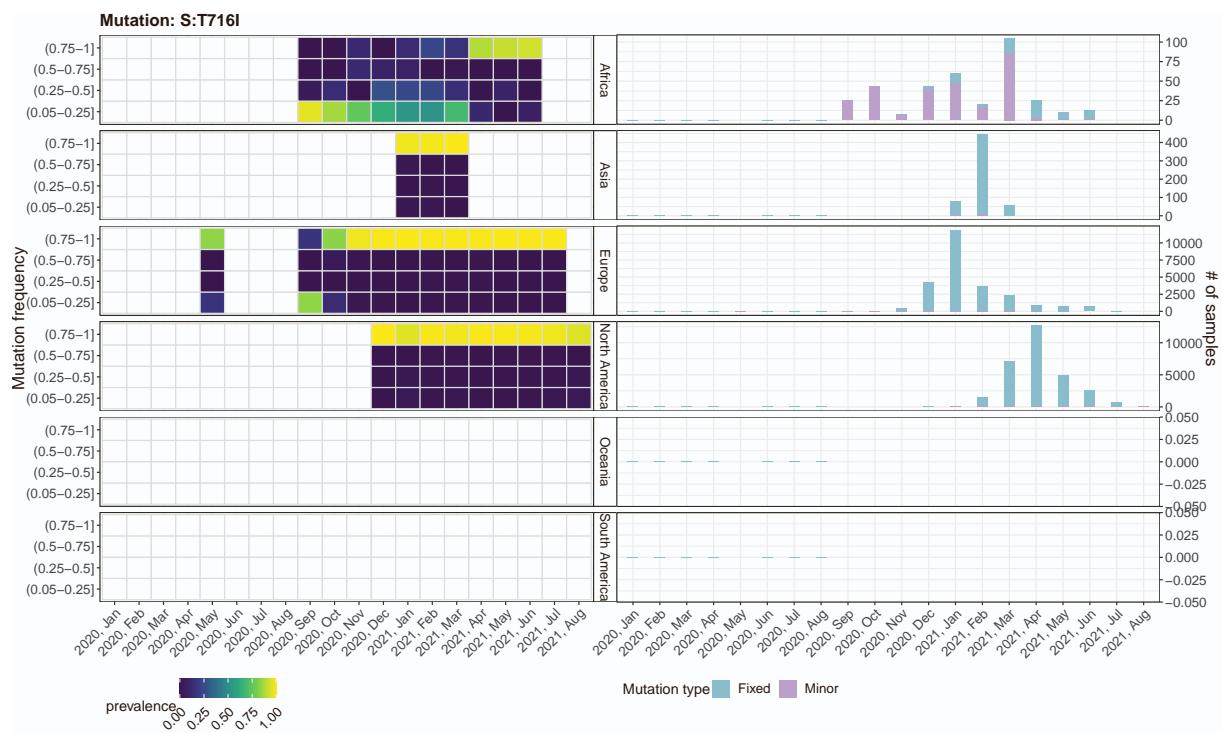


Figure S 21: Mutant frequency and prevalence variation in time of mutation: S:T716I.
Related to Figure 3. Please refer to the caption of Figure S 3

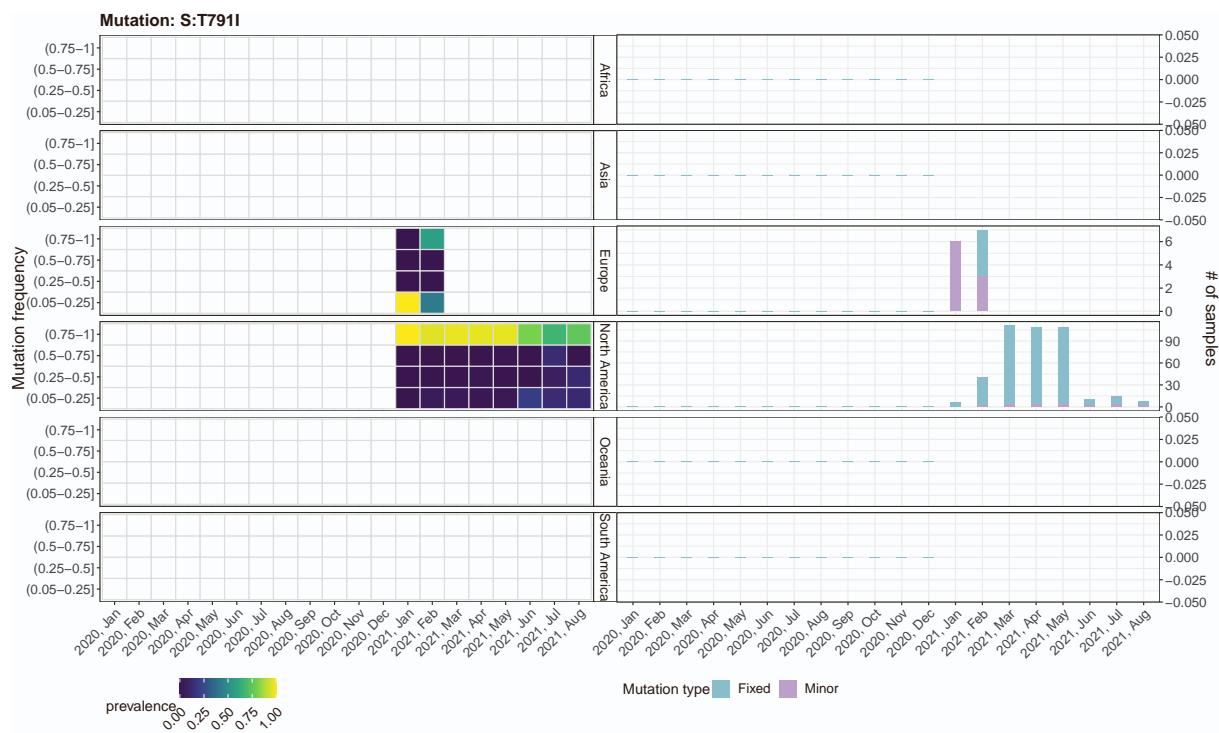


Figure S 22: Mutant frequency and prevalence variation in time of mutation: S:T791I. Related to Figure 3. Please refer to the caption of Figure S 3

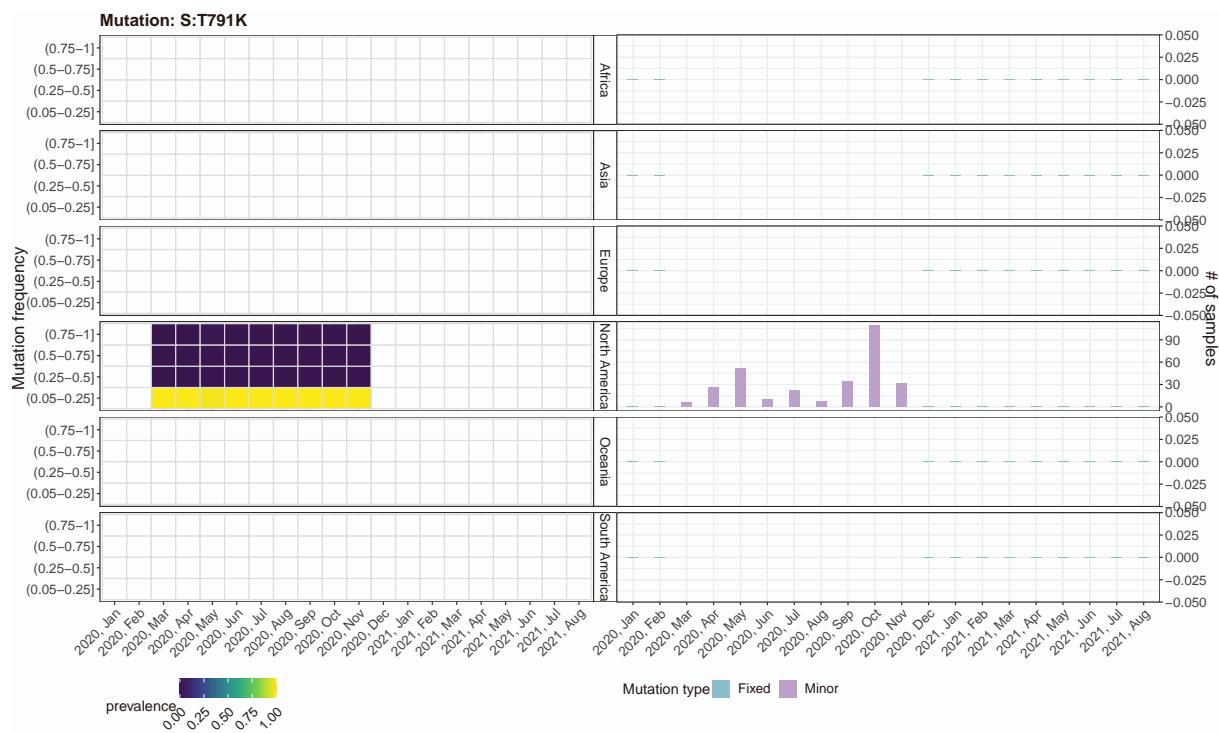


Figure S 23: Mutant frequency and prevalence variation in time of mutation: S:T791K. Related to Figure 3. Please refer to the caption of Figure S 3

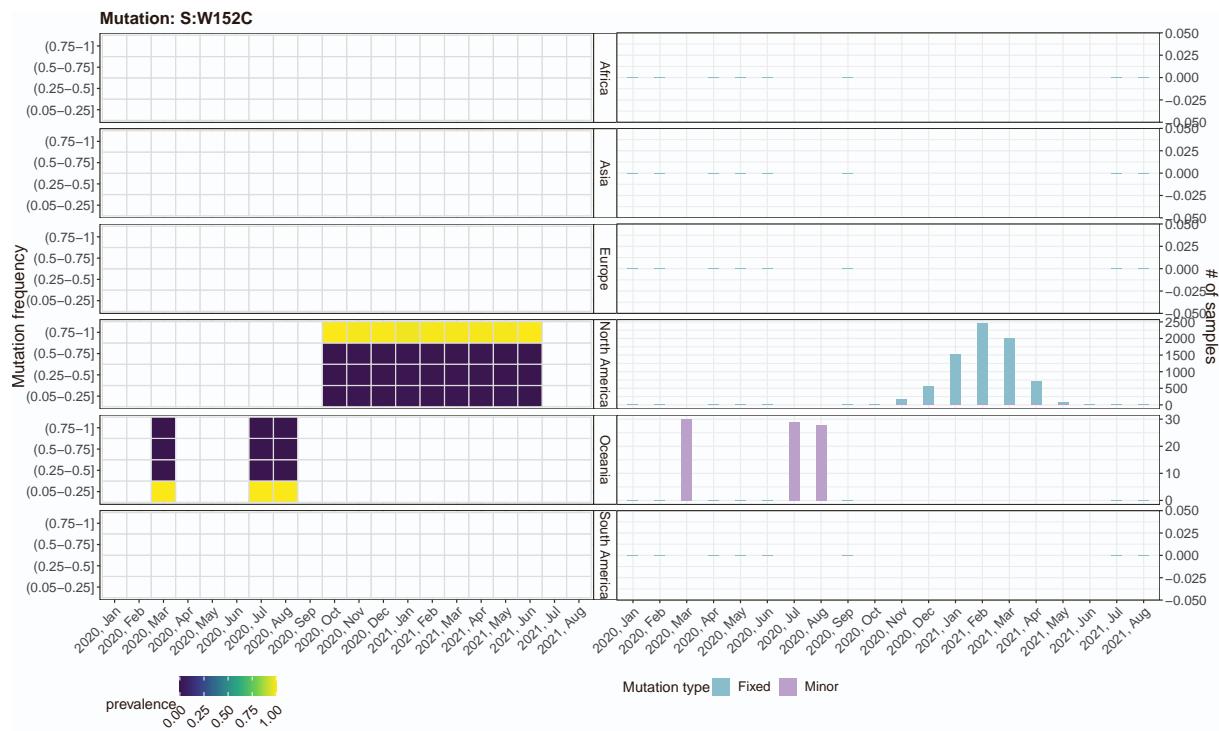


Figure S 24: Mutant frequency and prevalence variation in time of mutation: S:W152C.
Related to Figure 3. Please refer to the caption of Figure S 3

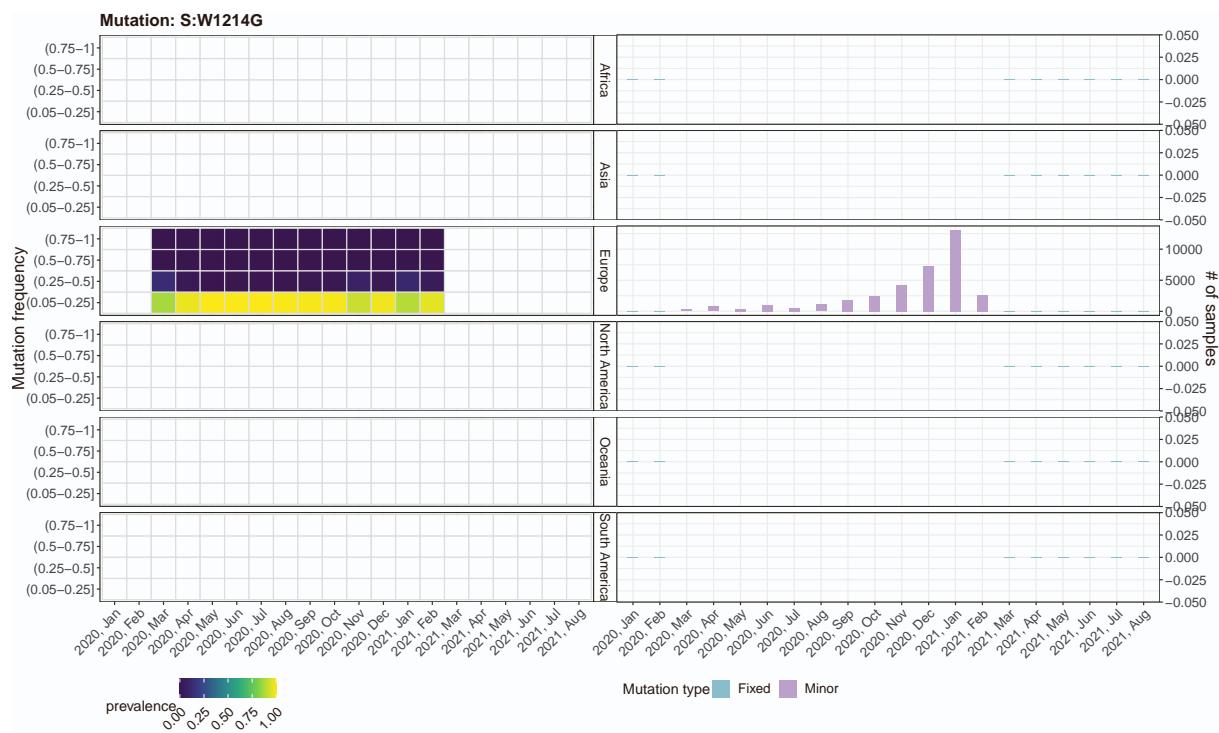


Figure S 25: Mutant frequency and prevalence variation in time of mutation: S:W1214G.
Related to Figure 3. Please refer to the caption of Figure S 3

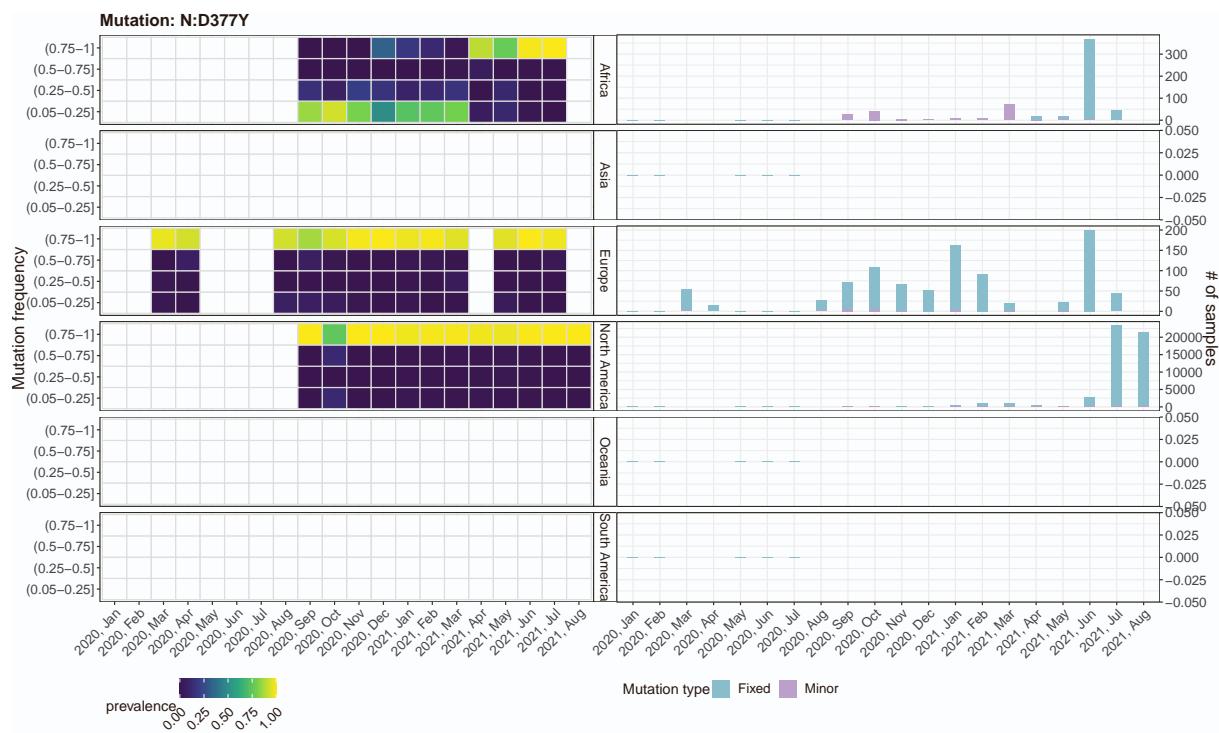


Figure S 26: Mutant frequency and prevalence variation in time of mutation: N:D377Y.
Related to Figure 3. Please refer to the caption of Figure S 3



Figure S 27: Mutant frequency and prevalence variation in time of mutation: N:A254S.
Related to Figure 3. Please refer to the caption of Figure S 3



Figure S 28: Mutant frequency and prevalence variation in time of mutation: N:A254V.
Related to Figure 3. Please refer to the caption of Figure S 3

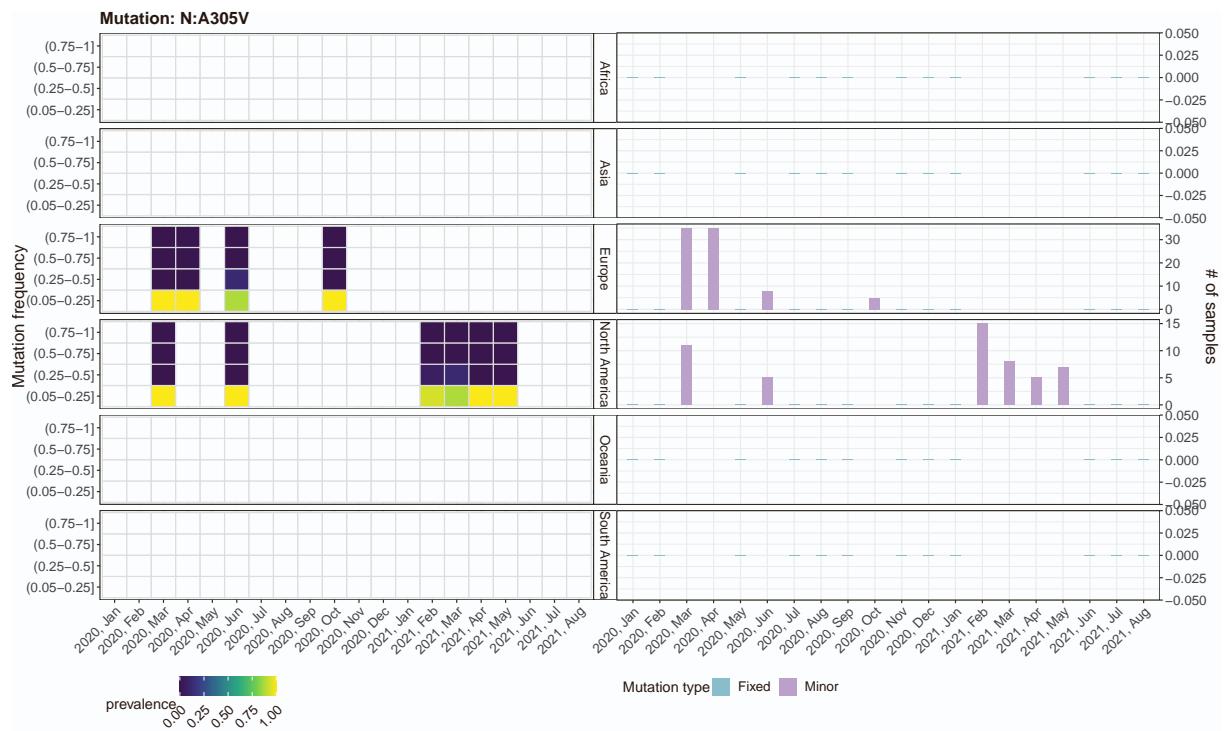


Figure S 29: Mutant frequency and prevalence variation in time of mutation: N:A305V.
Related to Figure 3. Please refer to the caption of Figure S 3

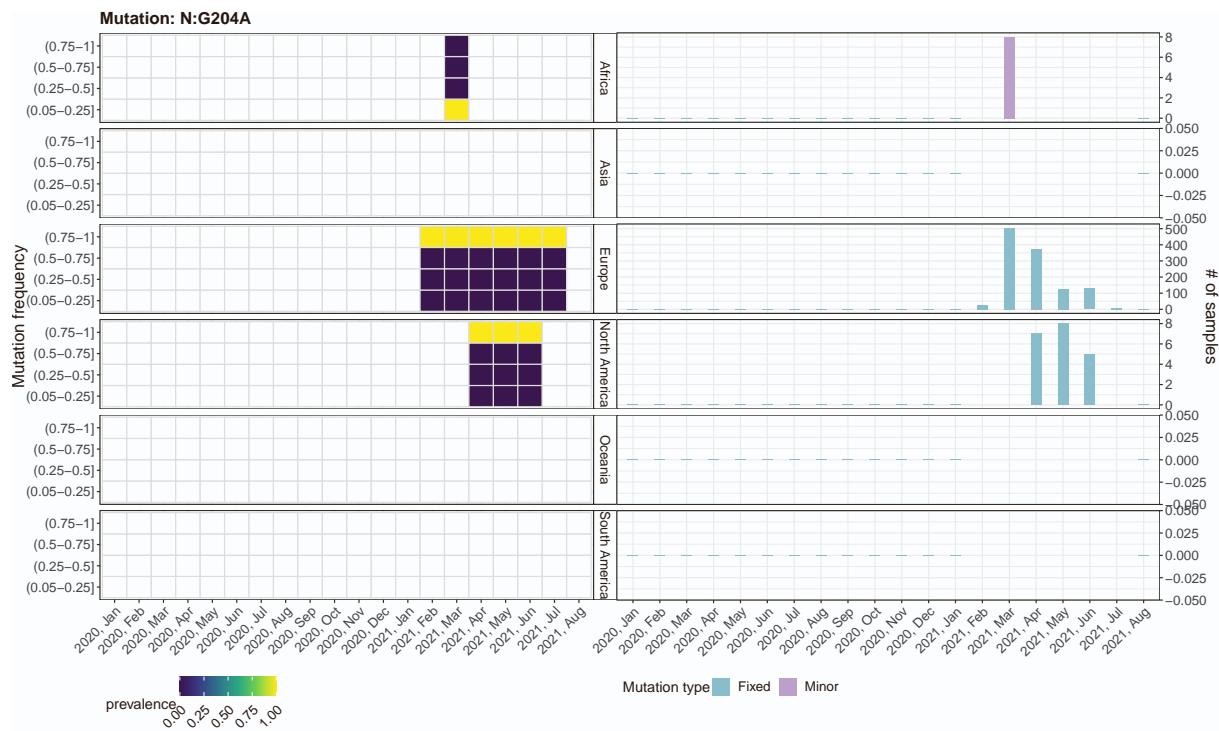


Figure S 30: Mutant frequency and prevalence variation in time of mutation: N:G204A.
Related to Figure 3. Please refer to the caption of Figure S 3

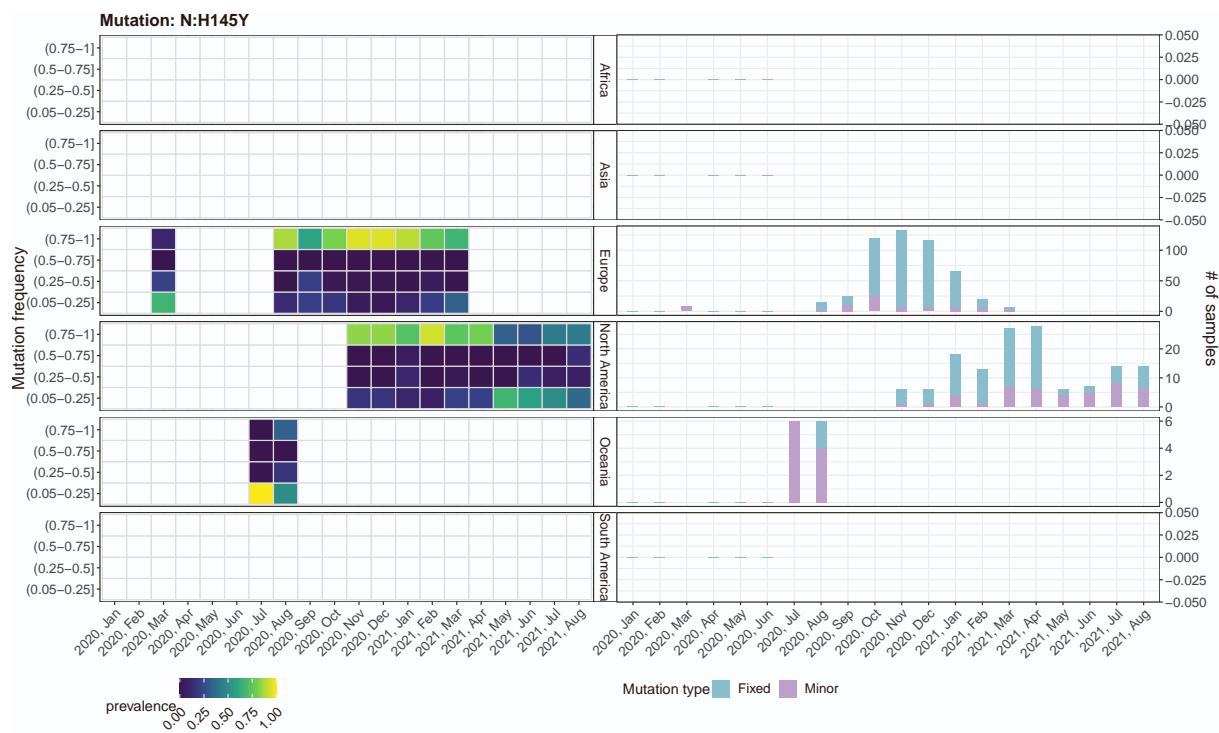


Figure S 31: Mutant frequency and prevalence variation in time of mutation: N:H145Y.
Related to Figure 3. Please refer to the caption of Figure S 3

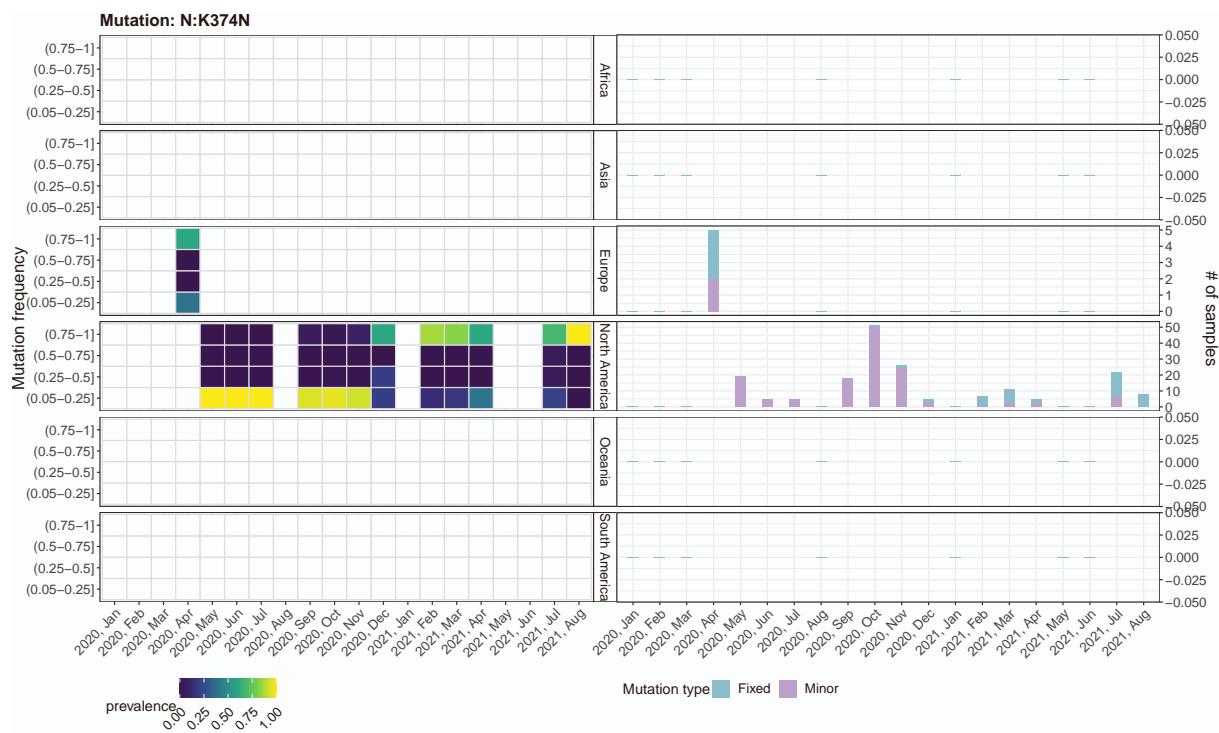


Figure S 32: Mutant frequency and prevalence variation in time of mutation: N:K374N.
Related to Figure 3. Please refer to the caption of Figure S 3

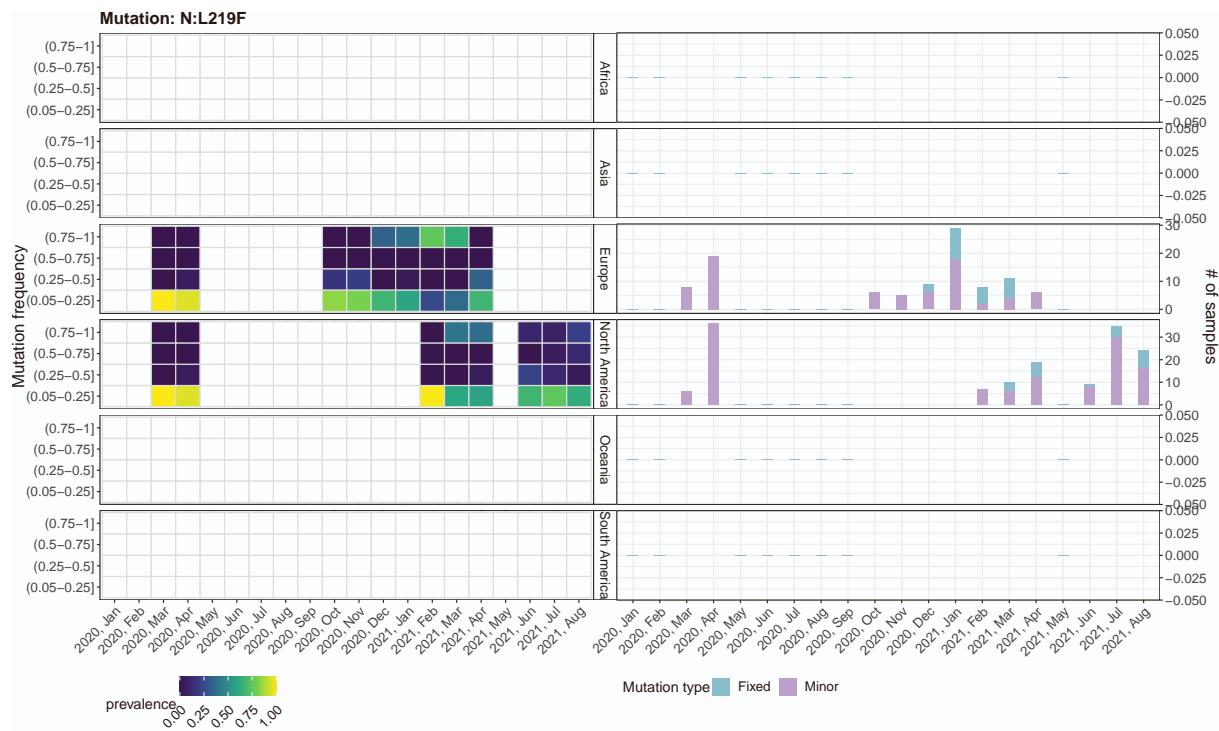


Figure S 33: Mutant frequency and prevalence variation in time of mutation: N:L219F.
Related to Figure 3. Please refer to the caption of Figure S 3

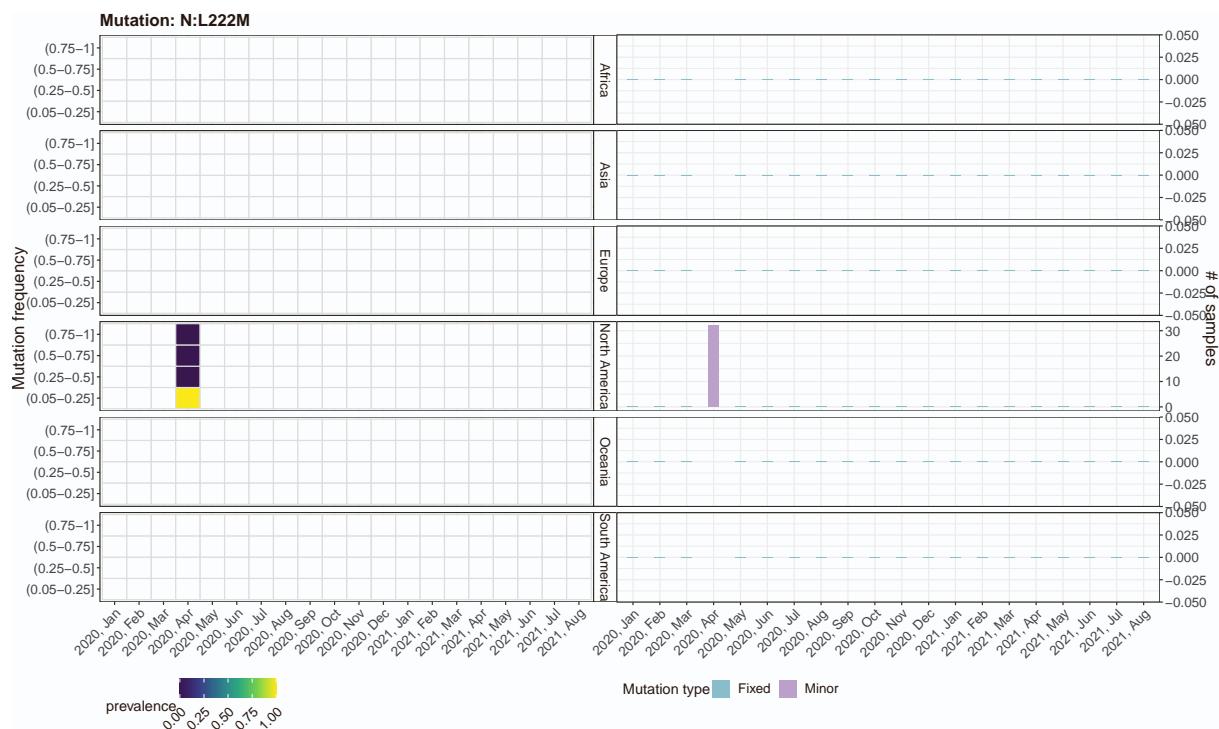


Figure S 34: Mutant frequency and prevalence variation in time of mutation: N:L222M.
Related to Figure 3. Please refer to the caption of Figure S 3

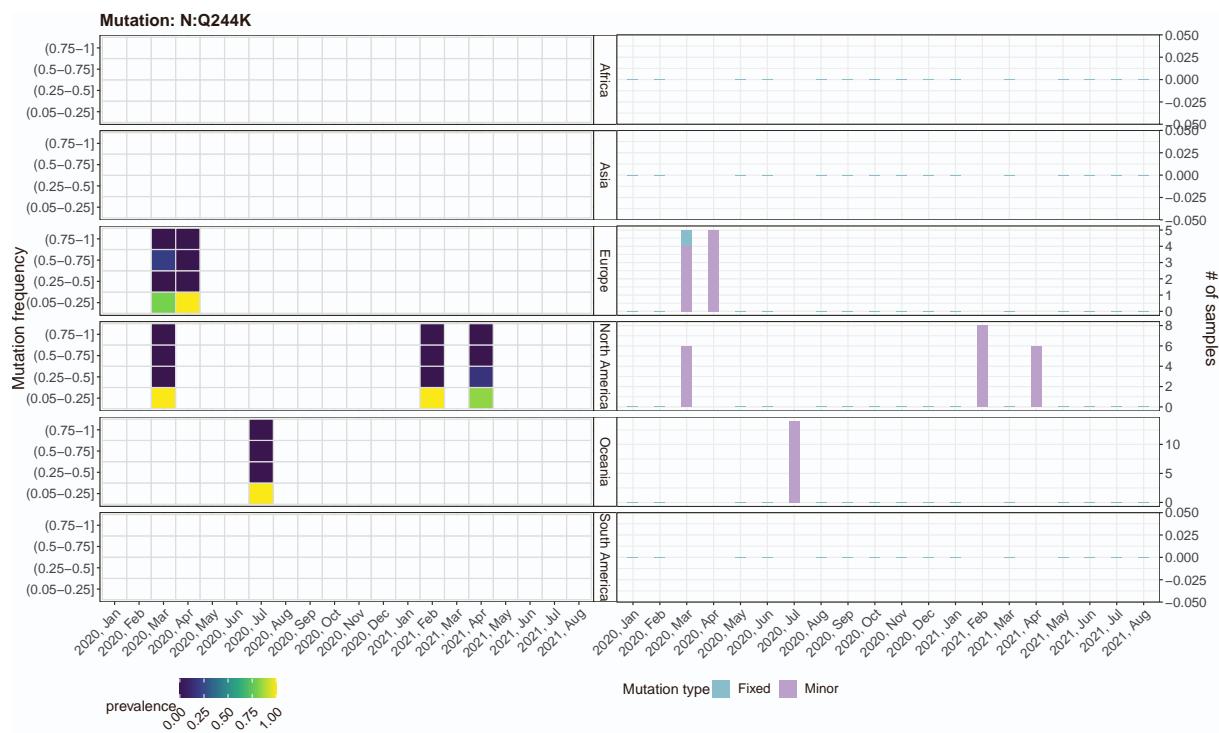


Figure S 35: Mutant frequency and prevalence variation in time of mutation: N:Q244K.
Related to Figure 3. Please refer to the caption of Figure S 3

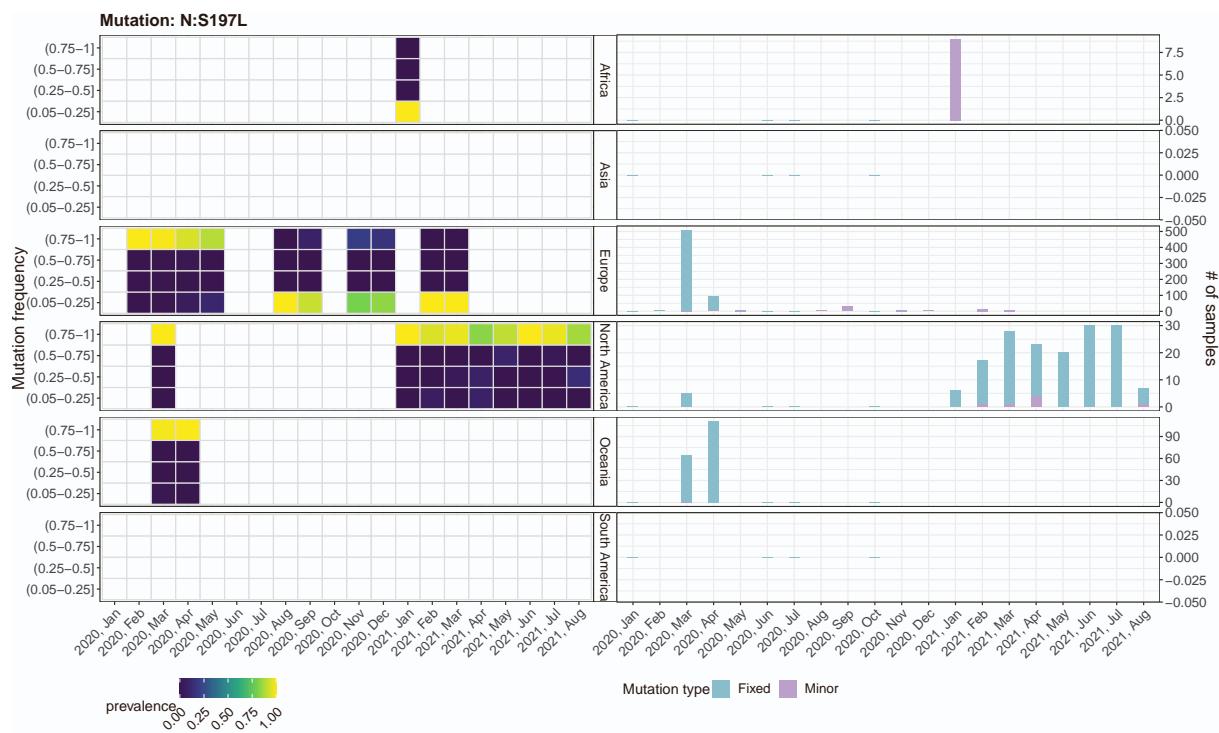


Figure S 36: Mutant frequency and prevalence variation in time of mutation: N:S197L.
Related to Figure 3. Please refer to the caption of Figure S 3