



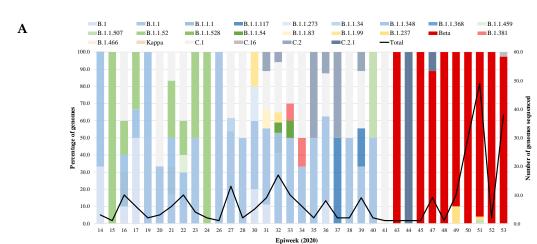
Correction

## Correction: Subramoney et al. Molecular Epidemiology of SARS-CoV-2 during Five COVID-19 Waves and the Significance of Low-Frequency Lineages. *Viruses* 2023, 15, 1194

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In the original publication [1], there was a mistake in Figure 1 as published. An error occurred occasionally, resulting in redundant horizontal lines in the bar graphs of 1A and 1C and the disappearance of the light yellow bars in 1C. The corrected Figure 1 appears below. The authors state that the scientific conclusions are unaffected. This correction was approved by the Academic Editor. The original publication has also been updated.



**Figure 1.** *Cont*.



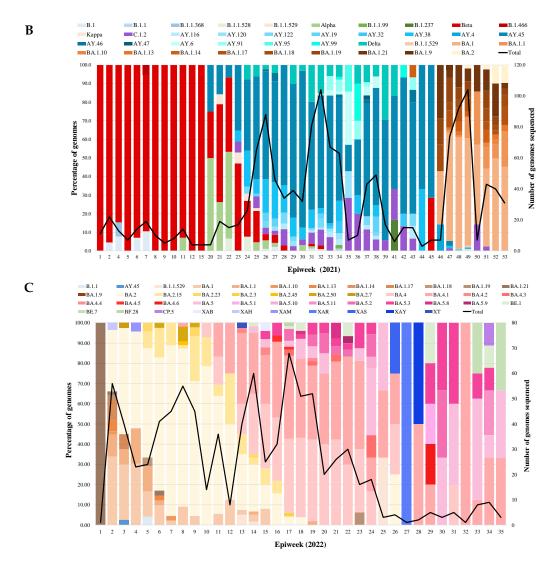
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Viruses **2023**, 15, 1502 2 of 2



**Figure 1.** Prevalence of SARS-CoV-2 lineages over time from 2020 to 2022 by epiweek. The bar graph represents the SAR-CoV-2 lineages and VOCs identified in our study cohort. The black line graph represents the total number of samples that were sequenced during each epiweek. **(A)** Distribution of SARS-CoV-2 lineages in 2020, **(B)** Distribution of SARS-CoV-2 lineages in 2021, and **(C)** Distribution of SARS-CoV-2 lineages in 2022.

## Reference

Subramoney, K.; Mtileni, N.; Giandhari, J.; Naidoo, Y.; Ramphal, Y.; Pillay, S.; Ramphal, U.; Maharaj, A.; Tshiabuila, D.; Tegally, H.; et al. Molecular Epidemiology of SARS-CoV-2 during Five COVID-19 Waves and the Significance of Low-Frequency Lineages. *Viruses* 2023, 15, 1194. [CrossRef] [PubMed]

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