*Correction*

# Correction: Kurpas et al. Genomic Analysis of SARS-CoV-2 Alpha, Beta and Delta Variants of Concern Uncovers Signatures of Neutral and Non-Neutral Evolution. *Viruses* 2022, *14*, 2375

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## Missing Funding

In the original publication [1], the funder Polish National Science Center, 2021/41/B/NZ2/04134 to Marek Kimmel (MK) was not included. The correct funding should be as follows:

## Text Correction

There was an error in the original publication. Practical computations of the SFS based on Tung–Durrett model were carried out using notation $\alpha = 1 + \lambda_0/\lambda_1$. However, this is inconsistent with the original notation $\alpha = \lambda_0/\lambda_1$. The correction makes expressions (6) and (7) consistent with the original notation.

A correction has been made to Materials and Methods, SFS under Birth-and-Death Process Model with Mutation and Selection, paragraph 1:

Under these hypotheses, if the fitnesses of the two types are $\lambda_0 < \lambda_1$, then the site frequency spectrum approximately follows the power curve, in our notation

$$S(x) = Cx^{-(1+\alpha)}, x = 1, \ldots, n \tag{6}$$

hence the tail $T(x) = \sum_{\xi > x} S(\xi)$ follows the law

$$T(x) = \frac{C}{\alpha}\left(n^{-\alpha} - x^{-\alpha}\right), x = 1, \ldots, n \tag{7}$$

where $\alpha = \lambda_0/\lambda_1$.

The authors state that the scientific conclusions are unaffected. These corrections were approved by the Academic Editor. The original publication has also been updated.

## Reference

1.    Kurpas, M.K.; Jaksik, R.; Kuś, P.; Kimmel, M. Genomic Analysis of SARS-CoV-2 Alpha, Beta and Delta Variants of Concern Uncovers Signatures of Neutral and Non-Neutral Evolution. *Viruses* **2022**, *14*, 2375. [CrossRef] [PubMed]

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