



## OPEN **Publisher Correction:** A machine learning approach for reliable prediction of amino acid interactions and its application in the directed evolution of enantioselective enzymes

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The original version of this Article contained a typographical error in Equation 1, where the division line in the fraction “ $\frac{n}{N}$ ” was omitted.

$$S(k) = \sum_{n=0}^{N-1} s(n)e^{-2i\pi k \frac{n}{N}}$$

now reads:

$$S(k) = \sum_{n=0}^{N-1} s(n)e^{-2i\pi k \frac{n}{N}}$$

The original Article has been corrected.



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