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

Frédéric Cadet, Nicolas Fontaine, Guangyue Li, Joaquin Sanchis, Matthieu Ng Fuk Chong, Rudy Pandjaitan, Iyanar Vetrivel , Bernard Offmann & Manfred T. Reetz

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.

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

now reads:

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

The original Article has been corrected.