








<https://doi.org/10.1038/s41467-021-24441-w>

OPEN

# Publisher Correction: Integrating genomics and metabolomics for scalable non-ribosomal peptide discovery

Bahar Behsaz, Edna Bode, Alexey Gurevich, Yan-Ni Shi, Florian Grundmann, Deepa Acharya, Andrés Mauricio Caraballo-Rodríguez , Amina Bouslimani, Morgan Panitchpakdi, Annabell Linck, Changhui Guan, Julia Oh, Pieter C. Dorrestein , Helge B. Bode , Pavel A. Pevzner  & Hosein Mohimani 

Correction to: *Nature Communications* <https://doi.org/10.1038/s41467-021-23502-4>, published online 28 May 2021.

The original version of this Article omitted Bahar Behsaz and Edna Bode as equally contributing authors.

The original version of this Article contained an error in the author affiliations.

Pavel A. Pevzner was incorrectly associated with Computational Biology Department, School of Computer Science, Carnegie Mellon University, Pittsburgh, PA, USA.

This has now been corrected in both the PDF and HTML versions of the Article.

Published online: 08 July 2021



**Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2021