














## Author Correction: LAMP-Seq enables sensitive, multiplexed COVID-19 diagnostics using molecular barcoding

Kerstin U. Ludwig, Ricarda M. Schmithausen , David Li, Max L. Jacobs , Ronja Hollstein, Katja Blumenstock, Jana Liebing, Mikołaj Stabicki , Amir Ben-Shmuel, Ofir Israeli, Shay Weiss, Thomas S. Ebert, Nir Paran , Wibke Rüdiger, Gero Wilbring , David Feldman, Bärbel Lippke, Nina Ishorst, Lara M. Hochfeld , Eva C. Beins , Ines H. Kaltheuner, Maximilian Schmitz, Aliona Wöhler, Manuel Döhla , Esther Sib, Marius Jentzsch, Eva-Maria C. Moench, Jacob D. Borrajo, Jonathan Strecker, Julia Reinhardt, Brian Cleary , Matthias Geyer, Michael Hölzel, Rhiannon Macrae , Markus M. Nöthen, Per Hoffmann , Martin Exner, Aviv Regev , Feng Zhang  and Jonathan L. Schmid-Burgk 

Correction to: *Nature Biotechnology* <https://doi.org/10.1038/s41587-021-00966-9>, published online 29 June 2021.

In the version of this article initially published, Eva-Maria C. Moench (MVZ Laboratory, Dr. Quade and Associates, Cologne, Germany) was missing from the author list. The omission has been corrected in the HTML and PDF versions of the article.

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## Publisher Correction: Integrative spatial analysis of cell morphologies and transcriptional states with MUSE

Feng Bao , Yue Deng , Sen Wan, Susan Q. Shen, Bo Wang, Qionghai Dai, Steven J. Altschuler  and Lani F. Wu

Correction to: *Nature Biotechnology* <https://doi.org/10.1038/s41587-022-01251-z>, published online 28 March 2022.

In the version of this article initially published, Extended Data Fig. 1 was truncated at panel **j**. The full image, running to panel **u**, has been amended in the HTML and PDF versions of the article.

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