

## Publisher Correction: Single-cell RNA-seq analysis software providers scramble to offer solutions

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Correction to: *Nature Biotechnology* <https://doi.org/10.1038/s41587-020-0449-8>, published online 9 March 2020.

In the version of this article initially published, the development of dropSeqPipe was credited to the lab of high-throughput scRNA-seq pioneer Steve McCarroll. It should have been credited to Patrick Roelli and colleagues at the Swiss Institute of Bioinformatics. The error has been corrected in the HTML and PDF versions of the article.

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