

Correction

COVID-19 immune features revealed by a large-scale single-cell transcriptome atlas

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<https://doi.org/10.1016/j.cell.2021.10.023>

(Cell 184, 1895–1913.e1–e19; April 1, 2021)

This article focused on dissecting the immune landscape of COVID-19 by single-cell RNA sequencing. The raw data were deposited into Genome Sequence Archive for human prior to publication, but the access has only been approved recently and is now available with accession ID HRA001149 (<https://ngdc.cncb.ac.cn/gsa-human/browse/HRA001149>). Accordingly, we have updated the data and code availability section by adding the following sentence: “The raw data are available from Genome Sequence Archive for human with accession ID HRA001149 (<https://ngdc.cncb.ac.cn/gsa-human/browse/HRA001149>).” Additionally, we inadvertently omitted the information of the following funding sources of this study in the published version of the paper and would like to add them to the acknowledgements: the Beijing Municipal of Science and Technology Major Project (grant number: Z201100005320014) and the National Natural Science Foundation of China (grant number: 81988101). The text has now been corrected online, and the authors apologize for any confusion these omissions may have caused.