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Correction

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

Xianwen Ren, Wen Wen, Xiaoying Fan, Wenhong Hou, Bin Su, Pengfei Cai, Jiasheng Li, Yang Liu, Fei Tang, Fan Zhang, Yu Yang, Jiangping He, Wenji Ma, Jingjing He, Pingping Wang, Qiqi Cao, Fangjin Chen, Yuqing Chen, Xuelian Cheng, Guohong Deng, Xilong Deng, Wenyu Ding, Yingmei Feng, Rui Gan, Chuang Guo, Weiqiang Guo, Shuai He, Chen Jiang, Juanran Liang, Yi-min Li, Jun Lin, Yun Ling, Haofei Liu, Jianwei Liu, Nianping Liu, Shu-Qiang Liu, Meng Luo, Qiang Ma, Qibing Song, Wujianan Sun, GaoXiang Wang, Feng Wang, Ying Wang, Xiaofeng Wen, Qian Wu, Gang Xu, Xiaowei Xie, Xinxin Xiong, Xudong Xing, Hao Xu, Chonghai Yin, Dongdong Yu, Kezhuo Yu, Jin Yuan, Biao Zhang, Peipei Zhang, Tong Zhang, Jincun Zhao, Peidong Zhao, Jianfeng Zhou, Wei Zhou, Sujuan Zhong, Xiaosong Zhong, Shuye Zhang, Lin Zhu, Ping Zhu, Bin Zou, Jiahua Zou, Zengtao Zuo, Fan Bai, Xi Huang, Penghui Zhou,* Qinghua Jiang,* Zhiwei Huang,* Jin-Xin Bei,* Lai Wei,* Xiu-Wu Bian,* Xindong Liu,* Tao Cheng,* Xiangpan Li,* Pingsen Zhao,* Fu-Sheng Wang,* Hongyang Wang,* Bing Su,* Zheng Zhang,* Kun Qu,* Xiaoqun Wang,* Jiekai Chen,* Ronghua Jin,* and Zemin Zhang*

*Correspondence: zhouph@sysucc.org.cn (P.Z.), qhjiang@hit.edu.cn (Q.J.), huangzhiwei2009@gmail.com (Z.H.), beijx@sysucc.org.cn (J.-X.B.), weilai@gzzoc.com (L.W.), bianxiuwu@263.net (X.-W.B.), xindongliu@hotmail.com (X.L.), chengtao@ihcams.ac.cn (T.C.), rm001227@whu.edu.cn (X.L.), zhaopingsen01@163.com (P.Z.), fswang302@163.com (F.-S.W.), hywangk@vip.sina.com (H.W.), bingsu@sjtu.edu.cn (B.S.), zhangzheng1975@aliyun.com (Z.Z.), qukun@ustc.edu.cn (K.Q.), xiaoqunwang@ibp.ac.cn (X.W.), chen_jiekai@gibh.ac.cn (J.C.), ronghuajin_youan@126.com (R.J.), zemin@pku.edu.cn (Z.Z.)
<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.