Genome Biology

Author Correction: Community-wide hackathons to identify central themes in single-cell multi-omics

Kim-Anh Lê Cao[1*], Al J. Abadi[1], Emily F. Davis-Marcisak[2], Lauren Hsu[3,4], Arshi Arora[5], Alexis Coullomb[6], Atul Deshpande[7], Yuzhou Feng[1], Pratheepa Jeganathan[8], Melanie Loth[7], Chen Meng[9], Wancen Mu[10], Vera Pancaldi[6,11], Kris Sankaran[12], Dario Righelli[13], Amrit Singh[14,15], Joshua S. Sodicoff[16,17], Genevieve L. Stein-O'Brien[2,7,18,19], Ayshwarya Subramanian[20], Joshua D. Welch[16,21], Yue You[22,23], Ricard Argelaguet[24], Vincent J. Carey[25], Ruben Dries[26,27,28], Casey S. Greene[29], Susan Holmes[30], Michael I. Love[10,31], Matthew E. Ritchie[22,23,32], Guo-Cheng Yuan[33], Aedin C. Culhane[3,4] and Elana Fertig[7,34,35]

The original article can be found online at https://doi.org/10.1186/s13059-021-02433-9.

* Correspondence: kimanh.lecao@nimelb.edu.au

[1]Melbourne Integrative Genomics, School of Mathematics and Statistics, University of Melbourne, Melbourne, Australia

Full list of author information is available at the end of the article

Correction to: Genome Biol 22, 220 (2021)

https://doi.org/10.1186/s13059-021-02433-9

Following publication of the original article [1], the authors identified that author Dario Righelli had erroneously been omitted from the author panel. The author group above has been updated and the original article [1] has been corrected.

Author details

[1]Melbourne Integrative Genomics, School of Mathematics and Statistics, University of Melbourne, Melbourne, Australia. [2]McKusick-Nathans Institute of the Department of Genetic Medicine, Johns Hopkins School of Medicine, Baltimore, MD, USA. [3]Data Science, Dana-Farber Cancer Institute, Boston, MA, USA. [4]Biostatistics, Harvard TH Chan School of Public Health, Boston, MA, USA. [5]Department of Epidemiology and Biostatistics, Memorial Sloan Kettering Cancer Center, New York, NY, USA. [6]Centre de Recherches en Cancérologie de Toulouse (INSERM), Université Paul Sabatier III, Toulouse, France. [7]Cancer Convergence Institute and Division of Quantitative Sciences, Department of Oncology, Sidney Kimmel Comprehensive Cancer Center, Johns Hopkins University School of Medicine, Baltimore, MD, USA. [8]Department of Mathematics and Statistics, McMaster University, Hamilton, Canada. [9]Bavarian Center for Biomolecular Mass Spectrometry (BayBioMS), School of Life Sciences, Technical University of Munich, Munich, Germany. [10]Department of Biostatistics, UNC, Chapel Hill, NC, USA. [11]Barcelona Supercomputing Center, Barcelona, Spain. [12]Department of Statistics, University of Wisconsin, Madison, WI, USA. [13]Department of Statistical Sciences, University of Padova, Padova, PD, Italy. [14]Department of Pathology and Laboratory Medicine, University of British Columbia, Vancouver, BC, Canada. [15]PROOF Centre of Excellence, Vancouver, BC, Canada. [16]Department of Computational Medicine and Bioinformatics, University of Michigan, Ann Arbor, MI, USA. [17]Department of Biomedical Engineering, University of Michigan, Ann Arbor, MI, USA. [18]Department of Neuroscience, Johns Hopkins University, Baltimore, MD, USA. [19]Kavli Neuroscience Discovery Institute, Johns Hopkins University, Baltimore, MD, USA. [20]Klarman Cell Observatory, Broad Institute of MIT and Harvard, Cambridge, MA, USA. [21]Department of Computer Science and Engineering, University of Michigan, Ann Arbor, MI, USA. [22]Epigenetics and Development Division, The Walter and Eliza Hall Institute of Medical Research, University of Melbourne, Melbourne, Australia. [23]Department of Medical Biology, University of Melbourne, Melbourne, Australia. [24]Epigenetics Programme, Babraham Institute, Cambridge CB22 3AT, UK. [25]Channing Division of Network Medicine, Brigham and Women's Hospital, Harvard Medical School, Boston, MA, USA. [26]Department of Hematology and Oncology, Boston Medical Center, Boston, MA, USA. [27]Department of Computational Biomedicine, Boston University School of Medicine, Boston, MA, USA. [28]Center for Regenerative Medicine (CReM), Boston University, Boston, MA, USA. [29]Center for Health AI and Department of Biochemistry and Molecular Genetics, University of Colorado School of Medicine, Aurora, CO, USA. [30]Department of Statistics, Stanford University, Stanford, CA, USA. [31]Department of Genetics, UNC, Chapel Hill, NC, USA. [32]School of Mathematics and Statistics, University of Melbourne, Melbourne, Australia. [33]Department of Genetics and Genomic

Cao *et al. Genome Biology*   (2021) 22:246

Page 2 of 2

Sciences, Charles Bronfman Institute for Personalized Medicine, Icahn School of Medicine at Mount Sinai, New York, NY, USA. [34]Department of Biomedical Engineering, Johns Hopkins University School of Medicine, Baltimore, MD, USA. [35]Department of Applied Mathematics and Statistics, Johns Hopkins University Whiting School of Engineering, Baltimore, MD, USA.

Reference

1.   Lê Cao KA, Abadi AJ, Davis-Marcisak EF, et al. Community-wide hackathons to identify central themes in single-cell multi-omics. Genome Biol. 2021;22:220 https://doi.org/10.1186/s13059-021-02433-9.