SCIENTIFIC REPORTS

OPEN

Author Correction: Whole genome diversity of inherited chromosomally integrated HHV-6 derived from healthy individuals of diverse geographic origin

Marco Telford¹, Arcadi Navarro^{1,2,3,4} & Gabriel Santpere^{1,5}

Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-018-21645-x>, published online 22 February 2018

The Acknowledgements section in this Article is incomplete.

“This work was supported by Ministerio de Ciencia e Innovación, Spain (BFU2012-38236 and BFU2015-68649-P to A.N.), by Direcció General de Recerca, Generalitat de Catalunya (2014SGR1311 and 2014SGR866), by the Spanish National Institute of Bioinformatics (PT13/0001/0026) and the REEM (RD12/0032/0011, RD16/0015/0017) of the Instituto de Salud Carlos III and by FEDER (Fondo Europeo de Desarrollo Regional)/FSE (Fondo Social Europeo).”

should read:

“This work was supported by Ministerio de Ciencia e Innovación, Spain (BFU2012-38236 and BFU2015-68649-P (MINECO/FEDER, UE) to A.N.), by Direcció General de Recerca, Generalitat de Catalunya (2014SGR1311 and 2014SGR866), by the Spanish National Institute of Bioinformatics (PT13/0001/0026) and the REEM (RD12/0032/0011, RD16/0015/0017) of the Instituto de Salud Carlos III and by FEDER (Fondo Europeo de Desarrollo Regional)/FSE (Fondo Social Europeo).”

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2018

¹Institute of Evolutionary Biology (UPF-CSIC), Departament de Ciències Experimentals i la Salut, Universitat Pompeu Fabra, PRBB, Barcelona, Catalonia, Spain. ²National Institute for Bioinformatics (INB), PRBB, Barcelona, Catalonia, Spain. ³Institució Catalana de Recerca i Estudis Avançats (ICREA), PRBB, Barcelona, Catalonia, Spain. ⁴Center for Genomic Regulation (CRG), PRBB, Barcelona, Catalonia, Spain. ⁵Department of Neuroscience, Yale School of Medicine, New Haven, CT, 06510, USA. Correspondence and requests for materials should be addressed to A.N. (email: arcadi.navarro@upf.edu) or G.S. (email: gabrielsantperebaro@gmail.com)