



Deposited via The University of Sheffield.

White Rose Research Online URL for this paper:

<https://eprints.whiterose.ac.uk/id/eprint/119426/>

Version: Supplemental Material

Article:

Elhaik, E., Tatarinova, T., Chebotarev, D. et al. (2014) Geographic population structure analysis of worldwide human populations infers their biogeographical origins. *Nature Communications*, 5 (3513). ISSN: 2041-1723

<https://doi.org/10.1038/ncomms4513>

Reuse

This article is distributed under the terms of the Creative Commons Attribution-NonCommercial-ShareAlike (CC BY-NC-SA) licence. This licence allows you to remix, tweak, and build upon this work non-commercially, as long as you credit the authors and license your new creations under the identical terms. More information and the full terms of the licence here: <https://creativecommons.org/licenses/>

Takedown

If you consider content in White Rose Research Online to be in breach of UK law, please notify us by emailing eprints@whiterose.ac.uk including the URL of the record and the reason for the withdrawal request.

Corrigendum: Geographic population structure analysis of worldwide human populations infers their biogeographical origins

Eran Elhaik, Tatiana Tatarinova, Dmitri Chebotarev, Ignazio S. Piras, Carla Maria Calò, Antonella De Montis, Manuela Atzori, Monica Marini, Sergio Tofanelli, Paolo Francalacci, Luca Pagani, Chris Tyler-Smith, Yali Xue, Francesco Cucca, Theodore G. Schurr, Jill B. Gaieski, Carlalynne Melendez, Miguel G. Vilar, Amanda C. Owings, Rocío Gómez, Ricardo Fujita, Fabrício R. Santos, David Comas, Oleg Balanovsky, Elena Balanovska, Pierre Zalloua, Himla Soodyall, Ramasamy Pitchappan, ArunKumar GaneshPrasad, Michael Hammer, Lisa Matisoo-Smith, R. Spencer Wells & The Genographic Consortium

Nature Communications 5:3513 doi: 10.1038/ncomms4513 (2014); Published 29 Apr 2014; Updated 31 Oct 2016

This article was published without any competing financial interests statement. A revised declaration that lists potentially competing financial interests is provided below: A.D.M., M.M. and M.A. are affiliated with Bcs Biotech S.r.l. J.B.G. is a consultant for the Genographic Project. M.H. is affiliated with FamilyTreeDNA. A.C., A.C.C., A.K.R., A.V.S., B.M.-C., C.F., C.J.A., C.S.I.D.S., D.E.P., D.F.S.H., D.R.L., H.L., J.B., J.R.S., J.S.Z., L.J., L.M.-S., L. Parida, L.Q.-M., M.C.D., M.E.K., M.G.V., M.H., O.A., N.C.M., P.P.V., P.S., S.A., S.L., S.R.W., R.J.M. and W.H. are part of the Genographic Consortium of the National Geographic. T.T. is developing a STTR grant application with Prosapia Genetics and IFXworks. The remaining authors declare no competing financial interests.

Furthermore, the affiliation details for Ajay K. Royyuru are incorrect in this Article. The correct affiliation details for this author are given below:

IBM, Somers, New York 10589, USA.



This work is licensed under a Creative Commons Attribution-NonCommercial-ShareAlike 4.0 International License. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder to reproduce the material. To view a copy of this license, visit <http://creativecommons.org/licenses/by-nc-sa/4.0/>

© The Author(s) 2016