

Published online: 02 August 2018

OPEN Author Correction: Assembly and **Analysis of Unmapped Genome Sequence Reads Reveal Novel** Sequence and Variation in Dogs

Lindsay A. Holden 1, Meharji Arumilli 2,3,4, Marjo K. Hytönen K. Hytönen Hundi Hundi Lindsay A. Holden 1, Meharji Arumilli Jarkko Salojärvi 5,6, Kim H. Brown & Hannes Lohi^{2,3,4}

Correction to: Scientific Reports https://doi.org/10.1038/s41598-018-29190-3, published online 18 July 2018

In this Article, the Data availability statement contains an incorrect link to the Supplementary Data 1-3. The correct Data availability statement is shown below:

"Novel contigs and Supplementary Data 1-3 are available at, https://pdxscholar.library.pdx.edu/bio_data/3/. Raw whole genome sequence reads for Border Collie dogs used in this study are available upon request."

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

¹Department of Biology, Portland State University, Portland, Oregon, USA. ²Research Programs Unit, Molecular Neurology, University of Helsinki, Helsinki, Finland. ³Department of Veterinary Biosciences, University of Helsinki, Helsinki, Finland. ⁴Folkhälsan Institute of Genetics, Helsinki, Finland. ⁵Research Programme on Individuals and Populations, Faculty of Biological and Environmental Sciences, University of Helsinki, Helsinki, Finland. ⁶School of Biological Sciences, Nanyang Technological University, Singapore, Singapore. Lindsay A. Holden and Meharji Arumilli contributed equally to this work. Kim H. Brown and Hannes Lohi jointly supervised this work. Correspondence and requests for materials should be addressed to K.H.B. (email: kibr2@pdx.edu) or H.L. (email: hannes.lohi@helsinki.fi)