



OPEN

Publisher Correction: Whole genome sequencing resolves 10 years diagnostic odyssey in familiar myxoma

Sára Pála, Judit Tőke, Anikó Bozsik, Henriett Butz, János Papp, István Likó, Enikő Kuroli, András Bánvölgyi, Mátyás Hamar, Jerome Bertherat, Márta Medvecz & Attila Patócs

Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-023-41878-9>, published online 05 September 2023

The Data availability section in the original version of this Article was incorrect.

“The datasets generated during and/or analysed during the current study are available from the corresponding author on reasonable request.”

now reads:

“We deposited the variant into the LOVD locus specific database, which was already revised and accepted by the curator. The reference number: <https://databases.lovd.nl/shared/variants/0000932922>”

The original Article has been corrected.



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 licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence 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 licence, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2023