CORRECTION



Correction to: Integrative proteomics highlight presynaptic alterations and c-Jun misactivation as convergent pathomechanisms in ALS

Amr Aly 1 · Zsofia I. Laszlo 2 · Sandeep Rajkumar 1 · Tugba Demir 1 · Nicole Hindley 2 · Douglas J. Lamont 3 · Johannes Lehmann 1 · Mira Seidel 1 · Daniel Sommer 1 · Mirita Franz-Wachtel 4 · Francesca Barletta 5 · Simon Heumos 5,6 · Stefan Czemmel 5 · Edor Kabashi 7 · Albert Ludolph 8,9 · Tobias M. Boeckers 1,9 · Christopher M. Henstridge 2 · Alberto Catanese 1,9

Published online: 12 September 2023

© The Author(s) 2023

Correction to:

Acta Neuropathologica (2023) 146:451–475 https://doi.org/10.1007/s00401-023-02611-y

The wrong Supplementary files (Figure and Table 2) were originally published with this article; it has now been replaced with the correct file.

The original article has been corrected.

Supplementary Information The online version contains supplementary material available at https://doi.org/10.1007/s00401-023-02630-9.

Data availability The MS proteomics data have been deposited to the ProteomeXchange Consortium (http://proteomecentral.proteomexchange.org) via the PRIDE partner repository and are available with the dataset identifiers PXD041543 (hiPSC-derivewd MNs) and PXD042617 (post mortem samples).

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/.

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

The original article can be found online at https://doi.org/10.1007/s00401-023-02611-y.

- Christopher M. Henstridge c.henstridge@dundee.ac.uk
- Alberto Catanese alberto.catanese@uni-ulm.de
- ¹ Institute of Anatomy and Cell Biology, Ulm University, Ulm, Germany
- Division of Cellular and Systems Medicine, School of Medicine, University of Dundee, Dundee, Scotland, UK
- FingerPrints Proteomics Facility, Discovery Centre, School of Life Sciences, University of Dundee, Dundee, Scotland, UK
- Proteome Center Tübingen, University of Tübingen, 72076 Tübingen, Germany

- Quantitative Biology Center (QBiC), University of Tübingen, 72076 Tübingen, Germany
- Biomedical Data Science, Department of Computer Science, University of Tübingen, 72076 Tübingen, Germany
- ⁷ Laboratory of Translational Research for Neurological Disorders, Imagine Institute, Université de Paris, INSERM, UMR 1163, 75015 Paris, France
- Department of Neurology, Ulm University School of Medicine, Ulm, Germany
- ⁹ German Center for Neurodegenerative Diseases (DZNE), Ulm Site, Germany

