







<https://doi.org/10.1038/s42003-022-03822-x>

OPEN

## Author Correction: GFI1 tethers the NuRD complex to open and transcriptionally active chromatin in myeloid progenitors

Anne Helness, Jennifer Fraszczak, Charles Joly-Beauparlant, Halil Bagci, Christian Trahan, Kaifée Arman , Peiman Shooshtarizadeh, Riyan Chen, Marina Ayoub, Jean-François Côté , Marlene Oeffinger, Arnaud Droit  & Tarik Möröy 

Correction to: *Communications Biology* <https://doi.org/10.1038/s42003-021-02889-2>, published online 02 December 2021.

The original version of this Article mistakenly referred to “anti-H3K9me2 (ab1220; AbCam)” when describing the Chromatin immunoprecipitation (ChIP) protocol in the Methods. The correct version of this text has now been corrected to reference “anti-H3K9me3 (ab8898; AbCam)”.

The original version of this Article was also missing the accession information for H3K4me1 ChIP-seq datasets in the Data Availability statement:

“Previously published ChIP-seq and ATAC-Seq data, which are presented in Fig. 4 and Suppl. Figure 4, are available under the following accession numbers: H3K27ac (GSM1441273), H3K27me3 (DRR023959), H3K9me3 (DRR023962), and ATAC-Seq (DRR023962).”

The correct statement now reads:

“Previously published ChIP-seq and ATAC-Seq data, which are presented in Fig. 4 and Suppl. Figure 4, are available under the following accession numbers: H3K27ac (GSM1441273), H3K27me3 (DRR023959), H3K9me3 (DRR023962), H3K4me1 (GSM1441289), and ATAC-Seq (DRR023962).”

Both errors have now been corrected in the PDF and HTML versions of the Article.

Published online: 07 September 2022



**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) 2022