https://doi.org/10.1038/s41594-022-00776-w

OPEN

Check for updates

Author Correction: SETDB1-like MET-2 promotes transcriptional silencing and development independently of its H3K9me-associated catalytic activity

Colin E. Delaney[®], Stephen P. Methot[®], Veronique Kalck, Jan Seebacher[®], Daniel Hess[®], Susan M. Gasser[®] and Jan Padeken

Correction to: Nature Structural & Molecular Biology https://doi.org/10.1038/s41594-021-00712-4, published online 31 January 2022.

In the version of this article initially published, there was a typographical error in the abstract. The cofactor LIN-65 originally read "LIN-61." The corrected sentence now reads "Our study suggests that the noncatalytic, focus-forming function of this SETDB1-like protein and its intrinsically disordered cofactor LIN-65 is physiologically relevant." The change has been made to the HTML and PDF versions the article.



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 were inside in other to the partial of the article's Creative Commons license.

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/.

Published online: 15 April 2022

https://doi.org/10.1038/s41594-022-00776-w

© The Author(s) 2022