



<https://doi.org/10.1038/s41467-021-21726-y>

OPEN

# Author Correction: An all-to-all approach to the identification of sequence-specific readers for epigenetic DNA modifications on cytosine

Guang Song , Guohua Wang , Ximei Luo , Ying Cheng, Qifeng Song, Jun Wan , Cedric Moore, Hongjun Song , Peng Jin , Jiang Qian  & Heng Zhu 

Correction to: *Nature Communications* <https://doi.org/10.1038/s41467-021-20950-w>, published online 04 February 2021.

In the original version of this Article, the “Methods” section “Genome-wide hmC profiling of human embryonic stem cell H1” incorrectly stated “Human embryonic stem cell H1 was purchased from WiCell Research Institute (WiCell) and the ethics approval was obtained from the Robert-Koch Institute, Berlin, Germany.”. Ethical approval was not required for the use of hESC H1 cells purchased from WiCell Research Institute. The statement has been corrected to “Human embryonic stem cell H1 was purchased from WiCell Research Institute (WiCell).” This has been corrected in the HTML and PDF version of this Article.

Published online: 23 February 2021



**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) 2021