

Publisher Correction: Profiling and promise of supermeres

James W. Clancy, Alex C. Boomgarden and Crislyn D'Souza-Schorey 

Correction to: *Nature Cell Biology* <https://doi.org/10.1038/s41556-021-00808-5>, published online 9 December 2021.










In the version of this article initially published, there were errors in Figs. 1 and 2 and the main text. In Fig. 1, “Extracellular particles” box, “Supermere” section, “TGF- β 1” replaces “AGFB1” under “Markers.” In the lower-right section of Fig. 1, the lower, second “LDL” header has been replaced by “VLDL” and “VLDL, very-low density lipoprotein” added to the caption. In Fig. 2 “Putative supermere biomarkers” box, “TGF- β 1” replaces “TGF β 1.” Further, in the last sentence of the fourth paragraph, “TGF- β 1” replaces “TGF- β 1.” The changes have been made to the online version of the article.

Published online: 10 January 2022

<https://doi.org/10.1038/s41556-022-00841-y>

© Springer Nature Limited 2022

Publisher Correction: Histone variant H3.3 maintains adult haematopoietic stem cell homeostasis by enforcing chromatin adaptability

Peipei Guo , Ying Liu, Fuqiang Geng, Andrew W. Daman, Xiaoyu Liu , Liangwen Zhong , Arjun Ravishankar , Raphael Lis , José Gabriel Barcia Durán , Tomer Itkin, Fanying Tang, Tuo Zhang , Jenny Xiang, Koji Shido , Bi-sen Ding, Duancheng Wen, Steven Z. Josefowicz and Shahin Rafii 

Correction to: *Nature Cell Biology* <https://doi.org/10.1038/s41556-021-00795-7>, published online 27 December 2021.

In the version of this article initially published, several edits were missing. In the legends of Fig. 1a and 2a, the word “view” has now been included, to read “Schematic view of non-competitive transplantation experiments” and “Schematic view of the in vitro HUVEC–HSPC co-culture experiment,” respectively. In the Fig. 3e *y*-axis label, “Percentage of LK cells” now replaces “Percentage of LKS cells,” while in Fig. 4i, second graph, the *y*-axis label “% CD16/32⁺ among LKS cells” replaces “% CD16/32⁺ among LK cells.” In the third sentence of the main text subsection “Derepressed ERV elicits antiviral responses,” “with H3.3 ... in HSPCs” has been removed from the sentence originally reading “There is a link between histone variants or histone modifying enzymes in suppressing ERVs and interferon signalling pathways^{37–39}; with H3.3 regulating ERV expression via H3K9me3 enrichment in HSPCs.” Further, in the last sentence of the penultimate paragraph of main text, “transcriptional” has been added to the sentence now reading “Representative genes with reduced H3K9me3 around promoters and reduced expression are Zfp84 and Zfp30, suggesting context-specific transcriptional regulations (Extended Data Fig. 10i and Supplementary Table 9).” The changes have been made to the HTML and PDF versions of the article.

Published online: 20 January 2022

<https://doi.org/10.1038/s41556-022-00851-w>

© The Author(s), under exclusive licence to Springer Nature Limited 2022