## Corrigendum

## BubR1 allelic effects drive phenotypic heterogeneity in mosaic-variegated aneuploidy progeria syndrome

Cynthia J. Sieben, Karthik B. Jeganathan, Grace G. Nelson, Ines Sturmlechner, Cheng Zhang, Willemijn H. van Deursen, Bjorn Bakker, Floris Foijer, Hu Li, Darren J. Baker, and Jan M. van Deursen

Original citation: J Clin Invest. 2020;130(1):171-188. https://doi.org/10.1172/JCI126863.

Citation for this corrigendum: J Clin Invest. 2020;130(11):6188. https://doi.org/10.1172/JCI144781.

During the preparation of this manuscript, the same sample was inadvertently included for the +/L1002P and +/- images in Figure 2E. The authors were able to use the original samples to prepare a corrected version. The correct figure is below.

The description in Methods for this panel has also been corrected, as below:

To assess the mitotic index of spontaneous lymphatic tumors, mitotic cells were visualized by IF labelling of at least 1 paraffin section for pHH3 (EMD Millipore, catalog 06-570), as described (6, 46).

The online version of the article has been updated with the corrected information.

The authors regret the errors.

