



## City Research Online

### City, University of London Institutional Repository

---

**Citation:** Cortés-Ciriano, I., Lee, J. J-K., Xi, R., Jain, D., Jung, Y. L., Yang, L., Gordenin, D., Klimczak, L. J., Zhang, C-Z., Pellman, D. S., et al (2020). Comprehensive analysis of chromothripsis in 2,658 human cancers using whole-genome sequencing. *Nature Genetics*, 52(3), pp. 331-341. doi: 10.1038/s41588-019-0576-7

This is the published version of the paper.

This version of the publication may differ from the final published version.

---

**Permanent repository link:** <https://openaccess.city.ac.uk/id/eprint/32290/>











**Link to published version:** <https://doi.org/10.1038/s41588-019-0576-7>

**Copyright:** City Research Online aims to make research outputs of City, University of London available to a wider audience. Copyright and Moral Rights remain with the author(s) and/or copyright holders. URLs from City Research Online may be freely distributed and linked to.

**Reuse:** Copies of full items can be used for personal research or study, educational, or not-for-profit purposes without prior permission or charge. Provided that the authors, title and full bibliographic details are credited, a hyperlink and/or URL is given for the original metadata page and the content is not changed in any way.



## Publisher Correction: Comprehensive molecular characterization of mitochondrial genomes in human cancers

Yuan Yuan , Young Seok Ju , Youngwook Kim , Jun Li , Yumeng Wang, Christopher J. Yoon, Yang Yang, Inigo Martincorena, Chad J. Creighton , John N. Weinstein , Yanxun Xu, Leng Han , Hyung-Lae Kim, Hidewaki Nakagawa, Keunchil Park , Peter J. Campbell , Han Liang  and PCAWG Consortium

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-019-0557-x>, published online 5 February 2020.

This paper was originally published without open access. As of the date of this correction, the paper is available online as an open-access paper under a Creative Commons Attribution 4.0 International License.










**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/>.

Published online: 11 February 2020

<https://doi.org/10.1038/s41588-020-0587-4>

© The Author(s) 2020

## Publisher Correction: Comprehensive analysis of chromothripsis in 2,658 human cancers using whole-genome sequencing

Isidro Cortés-Ciriano , Jake June-Koo Lee , Ruibin Xi , Dhawal Jain, Youngsook L. Jung, Lixing Yang, Dmitry Gordenin , Leszek J. Klimczak , Cheng-Zhong Zhang , David S. Pellman, PCAWG Structural Variation Working Group, Peter J. Park  and PCAWG Consortium

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-019-0576-7>, published online 5 February 2020.

In the version of this article initially published, author Peter J. Park had affiliation numbers 2 and 3; the correct affiliation numbers are 1 and 2. The error has been corrected in the HTML and PDF versions of the article.

Published online: 13 May 2020

<https://doi.org/10.1038/s41588-020-0634-1>

© The Author(s), under exclusive licence to Springer Nature America, Inc. 2020