

# Publisher Correction: Signatures of TOP1 transcription-associated mutagenesis in cancer and germline


<https://doi.org/10.1038/s41586-022-04812-z>

Published online: 3 May 2022

Correction to: *Nature* <https://doi.org/10.1038/s41586-022-04403-y>

Published online 9 February 2022

Open access

 Check for updates

**Martin A. M. Reijns, David A. Parry, Thomas C. Williams,  
Ferran Nadeu, Rebecca L. Hindshaw, Diana O. Rios Szwed,  
Michael D. Nicholson, Paula Carroll, Shelagh Boyle, Romina Royo,**

**Alex J. Cornish, Hang Xiang, Kate Ridout, The Genomics England  
Research Consortium\*, Colorectal Cancer Domain UK100,000  
Genomes Project\*, Anna Schuh, Konrad Aden, Claire Palles,  
Elias Campo, Tatjana Stankovic, Martin S. Taylor & Andrew P. Jackson**

In the version of this article initially published, there were errors in the affiliations shown for Colorectal Cancer Domain UK100,000 Genomes Project members Steve Thorn and Henry Wood. Steve Thorn is with the Cancer Research UK Edinburgh Centre, Institute of Genetics and Cancer, The University of Edinburgh, Edinburgh, UK (not University of Leeds). Henry Wood is with Pathology and Data Analytics, Leeds Institute of Medical Research, St James's University Hospital, University of Leeds, Leeds, UK (not The University of Edinburgh). The errors have been corrected in the HTML and PDF versions of the article.

\*A list of authors and their affiliations appears online.



**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) 2022