



## Author Correction: The copy number and mutational landscape of recurrent ovarian high-grade serous carcinoma


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The original version of this Article mistakenly misspelled “Carolin M. Sauer” from the author list as “Carolyn Sauer,” who is from the ‘CRUK Cambridge Institute, University of Cambridge, Cambridge, UK.’ This has been corrected in both the PDF and HTML versions of the Article.

The original version of this Article mistakenly misspelled “Ionut-Gabriel Funingana” from the author list as “Ionat-Gabriel Funingana,” who is from the ‘CRUK Cambridge Institute, University of Cambridge, Cambridge, UK’ and ‘Cambridge University Hospitals NHS Foundation Trust, Cambridge, UK.’ This has been corrected in both the PDF and HTML versions of the Article.

The original version of this Article contained an error in Fig. 1a, in which a panel indicating copy number event analysis was missing from the study workflow diagram. This has been corrected in both the PDF and HTML versions of the Article.

The original version of this Article contained an error in the Methods section which incorrectly read ‘Fastq sequencing reads were aligned to GRCh37 (g1kp2) using bwa samse.’ The correct version states ‘hs37d5’ in place of ‘g1kp2.’ This has been corrected in both the PDF and HTML versions of the Article.

The original version of this Article contained an error in the Data Availability Statement stating ‘The pre-processed single nucleotide variant and copy number data are available through a Zenodo data repository and can be downloaded here’ which linked to an incorrect version of the Zenodo repository. The correct link is as follows: <https://zenodo.org/record/7573784>. This has been corrected in both the PDF and HTML versions of the Article.

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