






## Author Correction: Proteome-wide structural changes measured with limited proteolysis-mass spectrometry: an advanced protocol for high-throughput applications

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Correction to: *Nature Protocols* <https://doi.org/10.1038/s41596-022-00771-x>. Published online 16 December 2022.

In the version of this article initially published, the citation at the end of the Fig. 6b caption was incorrect. The correct reference—Murtagh, F., Legendre, P. Ward’s hierarchical agglomerative clustering method: which algorithms implement Ward’s criterion? *J. Classif.* **31**, 274–295 (2014)—has been added and the reference list renumbered. In the second-to-last paragraph of the Introduction, the callout to Fig. 6e originally cited Fig. 7a and has been updated. The text changes have been made to the HTML and PDF versions of the article. Further, the numbered citations to Extended Data figures in the Supplementary Information were incorrect and have been replaced in a revised file online.

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## Author Correction: Reliable detection of somatic mutations in solid tissues by laser-capture microdissection and low-input DNA sequencing

Peter Ellis, Luiza Moore , Mathijs A. Sanders, Timothy M. Butler , Simon F. Brunner, Henry Lee-Six, Robert Osborne, Ben Farr, Tim H. H. Coorens , Andrew R. J. Lawson , Alex Cagan, Mike R. Stratton, Inigo Martincorena and Peter J. Campbell 

Correction to: *Nature Protocols* <https://doi.org/10.1038/s41596-020-00437-6>, published online 14 December 2020.

In the version of this article initially published, Step 24 showed an incorrect amount of mixture to be added to each ligation reaction. The amount should have been 130 µl, not 35. The error has been corrected in the HTML and PDF versions of the article.

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