

Author Correction: Capture-C: a modular and flexible approach for high-resolution chromosome conformation capture

Correction to: *Nature Protocols* <https://doi.org/10.1038/s41596-021-00651-w>, published online 4 February 2022.

<https://doi.org/10.1038/s41596-023-00860-5>

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 Check for updates

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In the version of this article initially published, the table in the “Fresh lysis buffer” section of “Reagent setup” showed incorrect information in the NaCl row: the stock concentration is 5 M, not 4 M, and the volume is 100 μ L, not 125 μ L. A Critical Step was missing after Step 13 in the Procedure: “Make sure that the cells are well re-suspended...”. Steps 21–23 omitted shaking instructions. In addition, Step 21 should have ended with “Keep Control 1 incubating along with digestion reactions without adding the enzyme”. In Step 43, the centrifuging time should have read 2 minutes instead of 30. Step 45 should have ended with “To increase the yield, leave the samples to dissolve at 4 °C overnight”. The second Critical Step in Step 52 was incomplete. Step 72 should have ended with “In indexed Tri-C samples, length should increase accordingly from -450 bp sonication product”. In Step 126, the two instances of “ μ g” should have been “ng”. Finally, Step 153 should not have ended with “and mix”. The errors have been corrected in the HTML and PDF versions of the article.

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Author Correction: Z-REX: shepherding reactive electrophiles to specific proteins expressed tissue specifically or ubiquitously, and recording the resultant functional electrophile-induced redox responses in larval fish

Correction to: *Nature Protocols* <https://doi.org/10.1038/s41596-023-00809-8>.
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