








Author Correction: Phage-assisted evolution of an adenine base editor with improved Cas domain compatibility and activity

Michelle F. Richter, Kevin T. Zhao , Elliot Eton , Audrone Lapinaite , Gregory A. Newby , Benjamin W. Thuronyi, Christopher Wilson , Luke W. Koblan, Jing Zeng, Daniel E. Bauer, Jennifer A. Doudna  and David R. Liu 

Correction to: *Nature Biotechnology* <https://doi.org/10.1038/s41587-020-0453-z>, published online 16 March 2020.


















In the version of this article initially published online, the edited bases at site 18 in Fig. 2e were labeled A6 and A8; they are A9 and A11, respectively. In Supplementary Fig. 6, the x-axis labels for site 18 read, from left to right, A2, A3, A4, A6, A8, A16, A17, A19 and A20; the correct labels are A5, A6, A7, A9, A11, A19, A20, A22 and A23. The errors have been corrected in the print, PDF and HTML versions of the article.

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Publisher Correction: Base editors for simultaneous introduction of C-to-T and A-to-G mutations

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Publisher Correction: Engineered CRISPR–Cas12a variants with increased activities and improved targeting ranges for gene, epigenetic and base editing

Benjamin P. Kleinstiver , Alexander A. Sousa, Russell T. Walton, Y. Esther Tak, Jonathan Y. Hsu, Kendell Clement, Moira M. Welch, Joy E. Horng , Jose Malagon-Lopez, Irene Scarfò , Marcela V. Maus , Luca Pinello, Martin J. Aryee  and J. Keith Joung

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In the HTML version of this article initially published, Benjamin P. Kleinstiver was not listed as a corresponding author.

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