

AUTHOR CORRECTION

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Author Correction: Combining different CRISPR nucleases for simultaneous knock-in and base editing prevents translocations in multiplex-edited CAR T cells

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Following publication of the original article [1], the authors identified an error in one of the guide RNA spacer sequences disclosed in Supplementary Table S3. The correct sequence for base editing mediated silencing of the *CIITA* is 5'–3': CACTCACCTTAG CCTGAGCA, as originally described in Gaudelli et al. 2020 [2].

This error does not affect the main results and conclusions of the paper.

The Supplementary Table S3 of the original article [1] has been corrected.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s13059-025-03548-z>.

Additional file 2: Supplementary Tables

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- 2 Gaudelli NM, Lam DK, Rees H, et al. Directed evolution of adenine base editors with increased activity and therapeutic application. *Nat Biotechnol.* 2020;38:7. <https://doi.org/10.1038/s41587-020-0491-6>.