

CORRECTION

Correction: All²: A tool for selecting mosaic mutations from comprehensive multi-cell comparisons

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There is an error in reference 11. The correct reference is:

Tu K, Lu K, Zhang Q, Huang W, Xie D. Accurate single-cell genotyping utilizing information from the local genome territory. *Nucleic Acids Res.* 2021 Jun 4;49(10):e57. doi: 10.1093/nar/gkab106. PMID: 33619552.

Reference

1. Sarangi V, Jang Y, Suvakov M, Bae T, Fasching L, Sekar S, et al. (2022) All²: A tool for selecting mosaic mutations from comprehensive multi-cell comparisons. *PLoS Comput Biol* 18(4): e1009487. <https://doi.org/10.1371/journal.pcbi.1009487> PMID: 35442945



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