

CORRECTION

Correction: A Next-Generation Sequencing Method for Genotyping-by-Sequencing of Highly Heterozygous Autotetraploid Potato

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There is an error in the second sentence of the sub-section Nucleotide Diversity within the Results. The correct sentence is: Mean π of the covered genome was 1.07×10^{-2} (Table 4).

Reference

1. Uitdewilligen JGAML, Wolters A-MA, D'hoop BB, Borm TJA, Visser RGF, van Eck HJ (2013) A Next-Generation Sequencing Method for Genotyping-by-Sequencing of Highly Heterozygous Autotetraploid Potato. PLoS ONE 8(5): e62355. doi: [10.1371/journal.pone.0062355](https://doi.org/10.1371/journal.pone.0062355) PMID: [23667470](https://pubmed.ncbi.nlm.nih.gov/23667470/)



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