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 PMID: 23667470

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