

CORRECTION

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Correction to: Spatiotemporal expression profile of novel and known small RNAs throughout rice plant development focussing on seed tissues

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Following publication of the original article [1], a typographical error was reported in the ‘Background’ section of the article. The incorrect and corrected sentences are given below, with the correction highlighted in bold:

The incorrect sentence read:

Twenty-one nt siRNAs, on the other hand, mainly cause transcriptional repression through the plant-specific RNA-directed DNA methylation (RdDM) pathway, inducing de novo DNA methylation [8].

The corrected sentence is:

Twenty-four nt siRNAs, on the other hand, mainly cause transcriptional repression through the plant-specific RNA-directed DNA methylation (RdDM) pathway, inducing de novo DNA methylation [8].

The original article [1] has been corrected.

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Reference

1. Meijer A, De Meyer T, Vandepoele K, et al. Spatiotemporal expression profile of novel and known small RNAs throughout rice plant development focussing on seed tissues. *BMC Genomics*. 2022;23:44. <https://doi.org/10.1186/s12864-021-08264-z>.

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