

CORRECTION

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Correction to: Genome-wide profiling of DNA methylation and gene expression identifies candidate genes for human diabetic neuropathy

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Correction to: Clin Epigenet 12, 123 (2020)

<https://doi.org/10.1186/s13148-020-00913-6>

Following publication of the original article [1], we were notified of a mistake in the way the author corrections have been implemented: a few random numbers had been incorrectly assigned to the second column in Fig. 5A. These have now been removed.

The original article has been corrected.

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Published online: 27 August 2020

Reference

1. Guo, et al. Genome-wide profiling of DNA methylation and gene expression identifies candidate genes for human diabetic neuropathy. *Clin Epigenet.* 2020;12:123. <https://doi.org/10.1186/s13148-020-00913-6>.

The original article can be found online at <https://doi.org/10.1186/s13148-020-00913-6>.

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