Clinical Epigenetics

CORRECTION   **Open Access**

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

Kai Guo[1†], Stephanie A. Eid[2†], Sarah E. Elzinga[2†], Crystal Pacut[2], Eva L. Feldman[2] and Junguk Hur[1*]

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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.

Author details
[1]Department of Biomedical Sciences, School of Medicine and Health Sciences, University of North Dakota, 1301 North Columbia Rd. Stop 9037, Grand Forks, ND 58202-9037, USA. [2]Department of Neurology, School of Medicine, University of Michigan, Ann Arbor, MI 48109, USA.

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.

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* Correspondence: junguk.hur@med.und.edu
[†]Kai Guo, Stephanie A. Eid and Sarah E. Elzinga contributed equally to this work.
[1]Department of Biomedical Sciences, School of Medicine and Health Sciences, University of North Dakota, 1301 North Columbia Rd. Stop 9037, Grand Forks, ND 58202-9037, USA
Full list of author information is available at the end of the article

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