



## Correction to: Large-scale pathway specific polygenic risk and transcriptomic community network analysis identifies novel functional pathways in Parkinson disease

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The article Large-scale pathway specific polygenic risk and transcriptomic community network analysis identifies novel

functional pathways in Parkinson disease, written by S. Bandres-Ciga, S. Saez-Atienzar, J. J. Kim, M. B. Makarious, F. Faghri, M. Diez-Fairen, H. Iwaki, H. Leonard, J. Botia, M. Ryten, D. Hernandez, J. R. Gibbs, J. Ding, Z. Gan-Or,

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The original article has been corrected.

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