

CORRECTION

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Correction: Lymphatic metastasis-associated circRNA–miRNA–mRNA network for exploring the pathogenesis and therapeutic target of triple negative breast cancer based on whole-transcriptome sequencing analysis: an experimental verification study

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Following publication of the original article [1], we have been notified that affiliation of the first author should be modified. They should be as follows:

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1. Luo J, Cao D, Hu C, Liang Z, Zhang Y, Lai J. Lymphatic metastasis-associated circRNA-miRNA-mRNA network for exploring the pathogenesis and therapeutic target of triple negative breast cancer based on whole-transcriptome sequencing analysis: an experimental verification study. *J Transl Med.* 2022;20(1):1.

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