

OPEN ACCESS

APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

*CORRESPONDENCE
Dan-dan Cao,
caodd@hku-szh.org
Yuan-Qing Yao,
yaoyuanqing@301hospital.com.cn

[†]These authors share first authorship

SPECIALTY SECTION

This article was submitted to RNA, a section of the journal Frontiers in Genetics

RECEIVED 28 June 2022 ACCEPTED 30 June 2022 PUBLISHED 10 August 2022

CITATION

Shang J, Cheng Y-F, Li M, Wang H, Zhang J-N, Guo X-M, Cao D-d and Yao Y-Q (2022), Corrigendum: Identification of key endometrial microRNAs and their target genes associated with pathogenesis of recurrent implantation failure by integrated bioinformatics analysis. *Front. Genet.* 13:979941. doi: 10.3389/fgene.2022.979941

COPYRIGHT

© 2022 Shang, Cheng, Li, Wang, Zhang, Guo, Cao and Yao. This is an openaccess article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Corrigendum: Identification of key endometrial microRNAs and their target genes associated with pathogenesis of recurrent implantation failure by integrated bioinformatics analysis

Jin Shang^{1†}, Yan-Fei Cheng^{2,3†}, Min Li^{4†}, Hui Wang⁵, Jin-Ning Zhang¹, Xin-Meng Guo⁶, Dan-dan Cao^{2*} and Yuan-Qing Yao^{2,4*}

¹Medical School of Chinese People's Liberation Army (PLA), Beijing, China, ²Shenzhen Key Laboratory of Fertility Regulation, Reproductive Medicine Center, The University of Hong Kong-Shenzhen Hospital, Shenzhen, China, ³Shenzhen Institute of Advanced Technology, Chinese Academy of Science, Shenzhen, China, ⁴Department of Obstetrics and Gynecology, The Seventh Medical Center, Chinese PLA General Hospital, Beijing, China, ⁵Department of Obstetrics and Gynecology, The First Medical Center, Chinese PLA General Hospital, Beijing, China, ⁶College of Medicine, Nankai University, Tianjin, China

KEYWORDS

recurrent implantation failure, differentially expressed genes, differentially expressed miRNAs, bioinformatics, endometrial transcriptomics

A Corrigendum on

Identification of key endometrial microRNAs and their target genes associated with pathogenesis of recurrent implantation failure by integrated bioinformatics analysis

by Shang J, Cheng Y-F, Li M, Wang H, Zhang J-N, Guo X-M, Cao D-d and Yao Y-Q (2022). Front. Genet. 13:919301. doi: 10.3389/fgene.2022.919301

In the published article, there was an error in author Dan-dan Cao's listed affiliation. Instead of "Dan-dan Cao^{3*}," it should be "Dan-dan Cao^{2*}."

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.