

CORRECTION

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Correction to: Genome-wide DNA methylome analysis identifies methylation signatures associated with survival and drug resistance of ovarian cancers

David W. Chan^{1*} , Wai-Yip Lam², Fushun Chen¹, Mingo M. H. Yung¹, Yau-Sang Chan¹, Wai-Sun Chan¹, Fangfang He¹, Stephanie S. Liu¹, Karen K. L. Chan¹, Benjamin Li² and Hextan Y. S. Ngan^{1,3*}

Correction to: *Clin Epigenet* (2021) 13:142

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Following publication of the original article [1], the authors identified an error in the result section.

The sentence currently read:

Indeed, the hypermethylated gene promoter regions of CLDN16 [...].

The sentence should read:

Indeed, the hypomethylated gene promoter regions of CLDN16 [...].

The original article [1] has been corrected.

Reference

1. Chan, et al. *Clin Epigenet*. 2021;13:142. <https://doi.org/10.1186/s13148-021-01130-5>.

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Author details

¹Department of Obstetrics and Gynaecology, L747 Laboratory Block, LKS Faculty of Medicine, 21 Sassoon Road, Pokfulam, Hong Kong, SAR, People's Republic of China. ²Lee's Pharmaceutical (HK) Ltd, 1/F Building 20E, Phase 3, Hong Kong Science Park, Shatin, Hong Kong, People's Republic of China. ³Department of Obstetrics and Gynaecology, 6/F Professorial Block, Queen Mary Hospital, Pokfulam, Hong Kong, People's Republic of China.

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*Correspondence: dwchan@hku.hk; hysngan@hku.hk

¹ Department of Obstetrics and Gynaecology, L747 Laboratory Block, LKS Faculty of Medicine, 21 Sassoon Road, Pokfulam, Hong Kong, SAR, People's Republic of China

Full list of author information is available at the end of the article



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