

CORRECTION

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

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## 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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