

CORRECTION

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Correction: Identification of five hub genes as monitoring biomarkers for breast cancer metastasis in silico

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Correction: *Hereditas* 156, 20 (2019).

<https://doi.org/10.1186/s41065-019-0096-6>.

Following publication of the original article [1], the author reported that Fig. 6B and 6C were repeated. The correct Figure is included here and the original article has been updated.

[†]Yun Cai, Jie Mei and Zhuang Xiao contributed equally to this work and should be considered co-first authors.

The online version of the original article can be found at <https://doi.org/10.1186/s41065-019-0096-6>.

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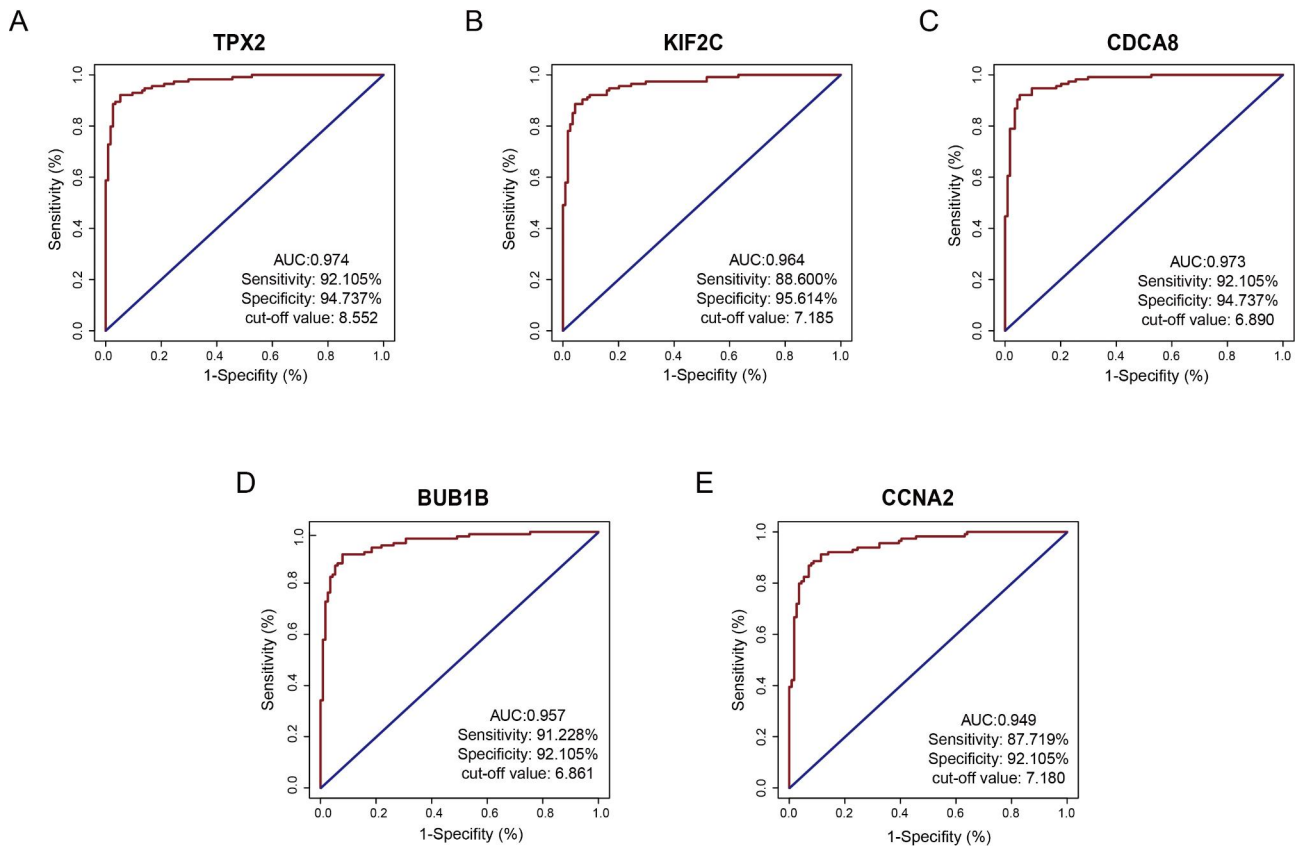


Fig. 6 Diagnostic value of the five hub genes in identifying normal and breast cancer tissues. The ROC curve revealed that the mRNA levels of these five genes exhibited excellent diagnostic efficiency for breast cancer and adjacent tissues. **(a)** TPX2, **(b)** KIF2C, **(c)** CDCA8, **(d)** BUB1B, **(e)** CCNA2

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References

1. Cai Y, Mei J, Xiao Z, et al. Identification of five hub genes as monitoring biomarkers for Breast cancer Metastasis in silico. *Hereditas*. 2019;156:20. <https://doi.org/10.1186/s41065-019-0096-6>.