

CORRECTION

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Correction: ScRNA-seq revealed an immunosuppression state and tumor microenvironment heterogeneity related to lymph node metastasis in prostate cancer

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and *EEF1A2*, resulting in an error in Fig. 3E, F in the published; the figure should have appeared as shown below.

In Fig. 3E, F of this article [1] the authors erroneously used the transwell picture of myc as the result of CCL5

[†]Shiyong Xin and Xiang Liu contributed equally to this work.

The original article can be found online at <https://doi.org/10.1186/s40164-023-00407-0>.

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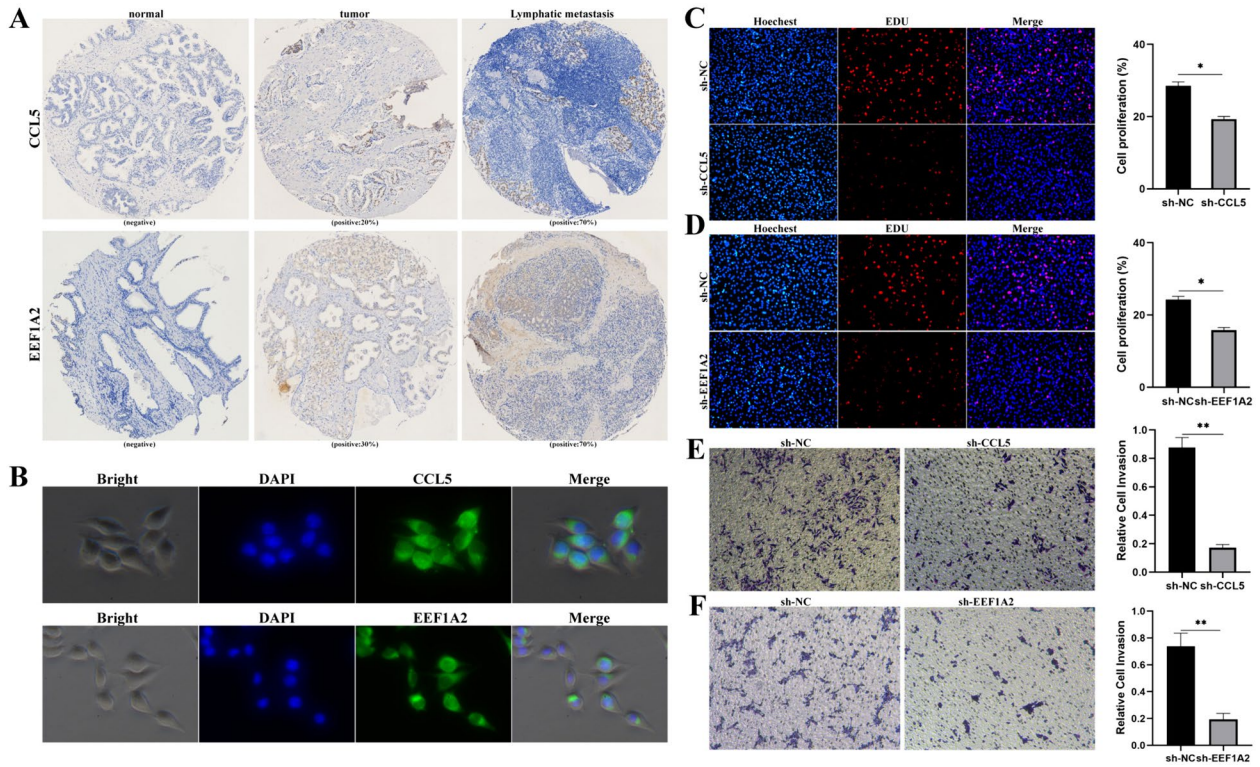


Fig. 3 EEF2+ and FOLH1+ luminal cells existed in PCa. **A** Immunohistochemistry analysis of CCL5 and EEF1A2 through PCa tissue microarray; **B** Immunofluorescence of CCL5 and EEF1A2 in LNCaP. **C, D** EDU showed the cell proliferation capacity of LNCaP after CCL5 and EEF1A2 down-regulation. **E, F** Metastatic ability of LNCaP cells was analyzed using transwell assay after down-regulation of CCL5 and EEF1A2

The original article has been corrected.

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- Xin S, Liu X, Li Z, Sun X, Wang R, Zhang Z, Feng X, Jin L, Li W, Tang C, Mei W, Cao Q, Wang H, Zhang J, Feng L, Ye L. ScRNA-seq revealed an immunosuppression state and tumor microenvironment heterogeneity related to lymph node metastasis in prostate cancer. *Exp Hematol Oncol*. 2023;12:49. <https://doi.org/10.1186/s40164-023-00407-0>.

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