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

Shiyong Xin^{1,2†}, Xiang Liu^{1,3†}, Ziyao Li¹, Xianchao Sun¹, Rong Wang⁴, Zhenhua Zhang⁵, Xinwei Feng⁶, Liang Jin¹, Weiyi Li¹, Chaozhi Tang¹, Wangli Mei¹, Qiong Cao⁷, Haojie Wang⁸, Jianguo Zhang², Lijin Feng^{9*} and Lin Ye^{1*}

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

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

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

*Correspondence: Lijin Feng fenglijin_2049@tongji.edu.cn

doctyelin@126.com

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

¹ Department of Urology, Shanghai East Hospital, School of Medicine, Tongji University, No.150, Ji-mo Rd, Pu-Dong New District, Shanghai 200120, China

² Department of Urology, The First Affiliated Hospital and College of Clinical Medicine of Henan University of Science and Technology, Luoyang 471003, China

³ Department of Urology, Putuo People's Hospital, School of Medicine, Shanghai, China

⁴ School of Pharmacy, Inner Mongolia Medical University, Hohhot 010000, China

⁵ School of Pharmaceutical Sciences, Sun Yat-Sen University, Guangzhou 510006, China

⁶ School of Pharmaceutical Sciences (Shenzhen), Shenzhen Campus of Sun Yat-Sen University, Shenzhen 518107, China

⁷ Department of Pathology, The Third Affiliated Hospital of Henan University of Science and Technology, Henan 471003, China

⁸ Department of Central Laboratory, Zhengzhou University, Luoyang Central Hospital, Luoyang 471003, China

⁹ Department of Pathology, Jing'an District Zhabei Central Hospital, No.619, Zhonghuaxin Road, Shanghai 200070, China

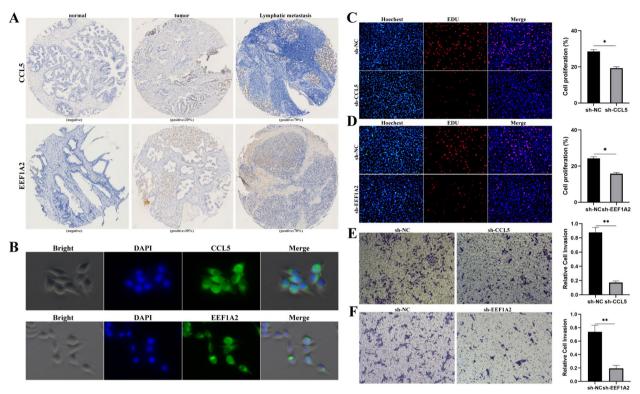


Fig. 3 EEF2+and FOLH1+luminal cells existed in PCa. A Immunohistochemistry analysis of CCL5 and EEF1A2 through PCa tissue microarray; B Immunofluorescence of CCL5 and EEF1A2 in LNCap. C, D EDU showed the cell proliferation capacity of LNCaP after CCL5 and EEF1A2 down-regulation. E, F Metastatic ability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