

CORRECTION

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Correction to: miR-205-5p inhibits human endometriosis progression by targeting ANGPT2 in endometrial stromal cells

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The original article [1] contains an error in Fig. 1 whereby in sub-panel A of Fig. 1, the blue and red thresholds were mistakenly reversed. The correct version of sub-panel A in Fig. 1 can be viewed ahead.

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1. Zhou C-F, et al. miR-205-5p inhibits human endometriosis progression by targeting ANGPT2 in endometrial stromal cells. *Stem Cell Res Ther.* 2019;10:287 <https://doi.org/10.1186/s13287-019-1388-5>.

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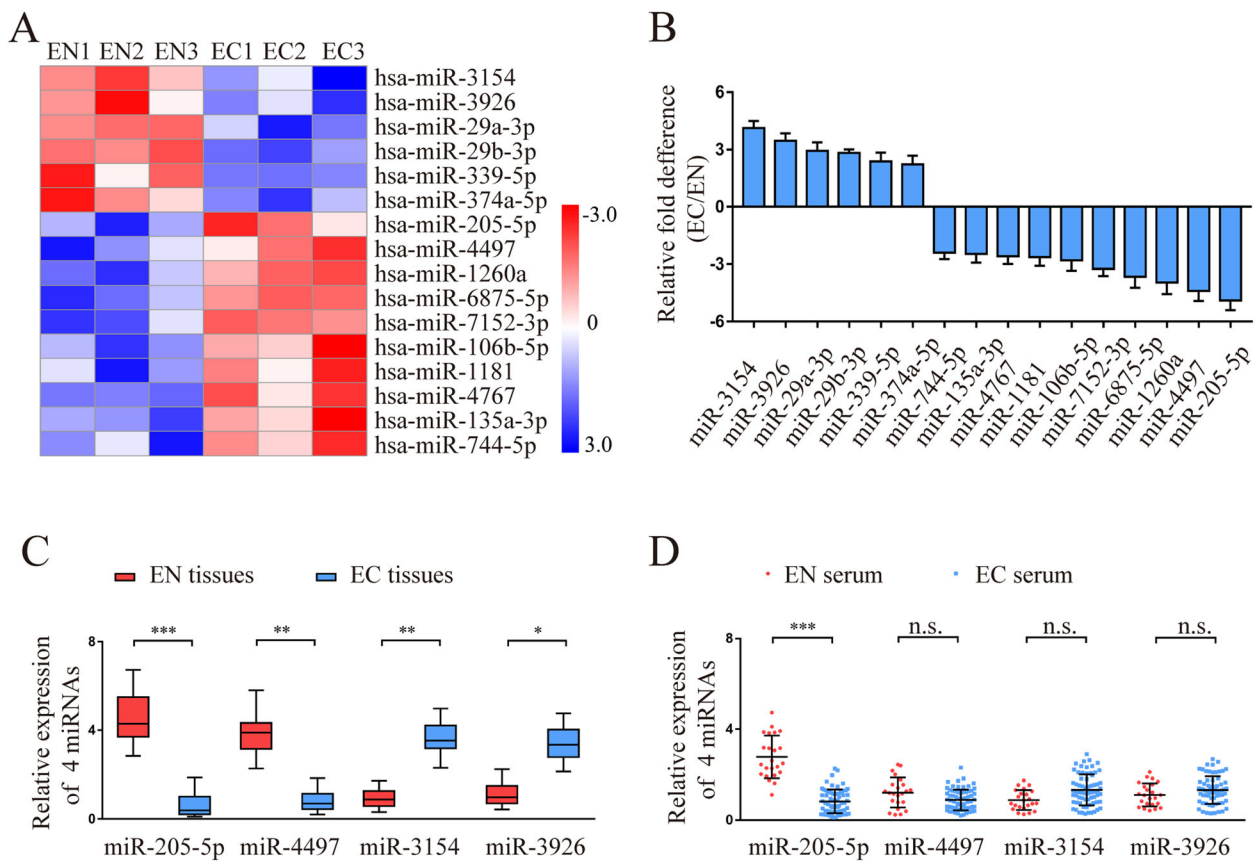


Fig. 1 Identification of miR-205-5p as a negatively pathologic miRNA in endometriosis. **a** The different miRNA expression profiles between the EN ($n = 3$) and EC ($n = 3$) groups were analysed by miRNA microarray. The heatmap diagram shows that the representative miRNAs were significantly associated with endometriosis. EN1, EN2, and EN3 indicate 3 normal endometria; EC1, EC2, and EC3 indicate 3 ectopic endometria. **b** The levels of 16 differentially expressed miRNAs in the endometrium used for microarray analysis were analysed by qRT-PCR. **c, d** The levels of miR-205-5p, miR-4497, miR-3154, and miR-3926 were validated by qRT-PCR in additional tissues and serum from the EN ($n = 23$) and EC ($n = 68$) groups. EN, normal endometrium; EC, ectopic endometrium. Error bars represent the mean \pm SD of 3 independent experiments. n.s., not significant; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$