



## Correction: Integrative network biology analysis identifies miR-508-3p as the determinant for the mesenchymal identity and a strong prognostic biomarker of ovarian cancer

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Published online: 3 September 2019

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### Correction to: Oncogene

<https://doi.org/10.1038/s41388-018-0577-5>  
published online 26 November 2018

Following publication of this article, the authors noted that they accidentally misplaced the figure in supplementary

Fig. 7b when re-editing the figures to meet the format and the image resolution of the journal. The amended version of supplementary Fig. 7 is provided below. This change does not affect the results or conclusions reported in the paper.

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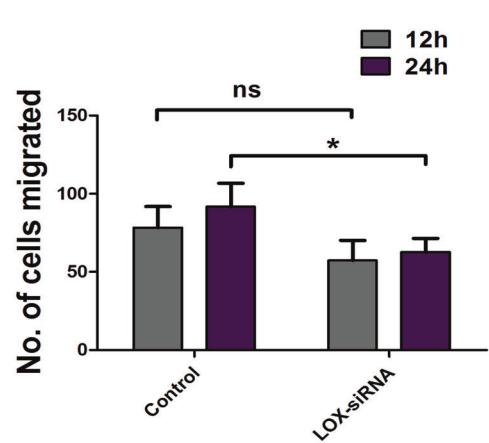
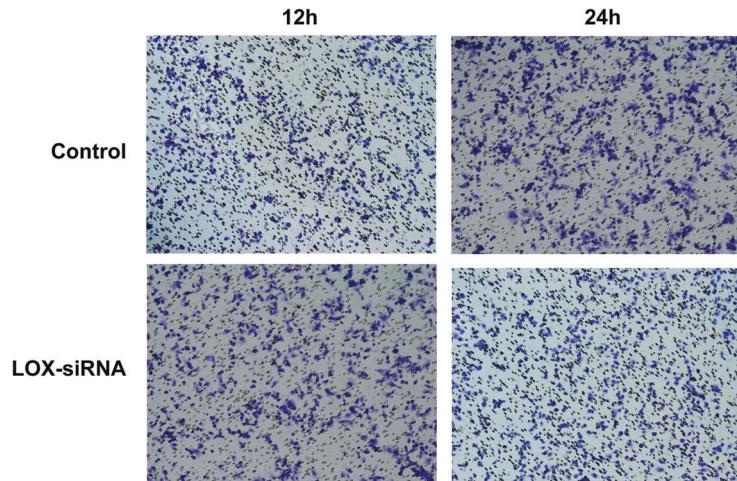
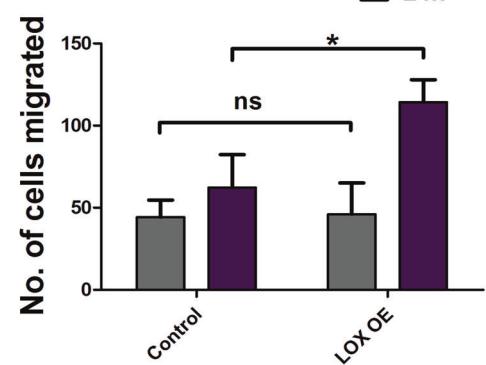
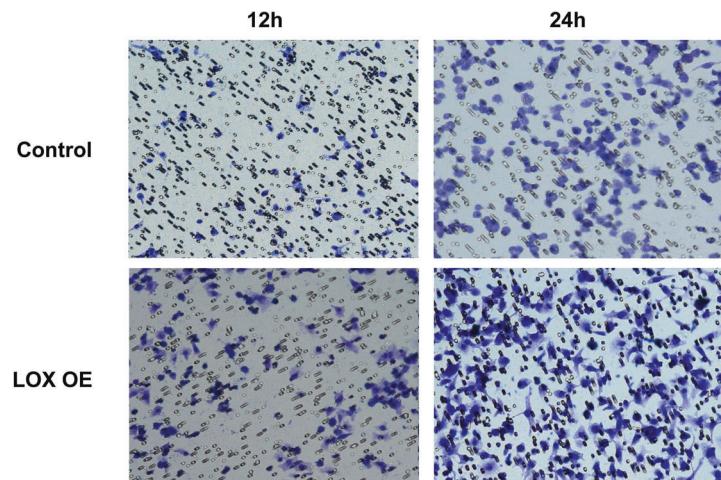
The original article can be found online at <https://doi.org/10.1038/s41388-018-0577-5>.

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**a****miR-508-3p inhibitor-transfected OV56 cells****b****miR-508-3p mimic-transfected COV504 cells**

**Fig. S7 a** Transwell chamber analysis of control or LOX silenced OV56 cells transfected with miR-508-3p inhibitor. **b** Transwell chamber analysis of control or LOX-overexpressing COV504 cells

transfected with miR-508-3p mimic. In all bar plots, p-values are based on two-tailed Student's t-tests ( $*P < 0.05$ ,  $**P < 0.01$ ,  $***P < 0.001$ )