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Erratum

Profiling the epigenetic interplay of lncRNA RUNXOR and oncogenic RUNX1 in breast cancer cells by gene in situ cis-activation: Am J Cancer Res. 2019; 9(8): 1635-1649

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In this article, we found an error in Figure S4B (A3 panel) due to a "mis-paste" of cropped images during figure preparation using the Illustrator software. We have now corrected it. Therefore, we would like to publish this Erratum to reflect this change. This correction does not affect the interpretation and conclusions of the study. We apologize for this error. The corrected Figure S4 is as follows.

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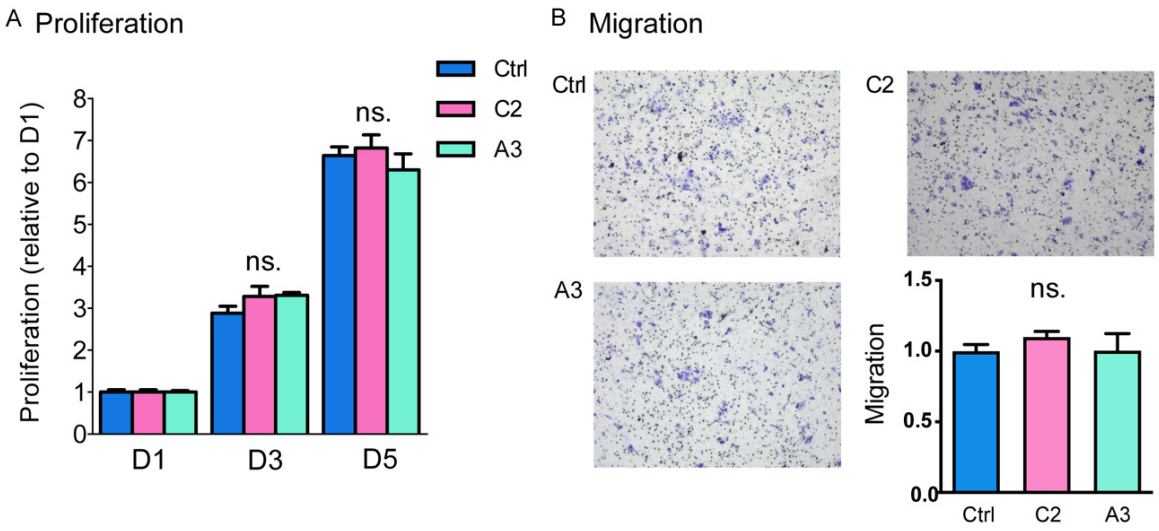

Figure S4. Phenotypic changes upon RUNXOR cis-overexpression. A. Cell proliferation. Cells were cultured for five consecutive days, and proliferation was assessed at D1, D3, and D5 using the CCK-8 assay. Cell growth was calculated based on absorbance values relative to day 1 (D1, set as 1). Experiments were performed in triplicate, and data are presented as mean ± SEM. B. Cell migration. Cell migration was assessed using a Transwell assay. Cells that migrated through the membrane were fixed, stained with crystal violet, and quantified using ImageJ. Migration ability was calculated based on the number of migrated cells relative to the Ctrl group (Ctrl, set as 1). Experiments were performed in triplicate. Representative images of migrated cells are shown.