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Corrigendum: SNORA72 activates the Notch1/ c-Myc pathway to promote stemness transformation of ovarian cancer cells

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A Corrigendum on

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In the original article, there was a mistake in Figure 1 as published. The transwell picture for CA in Figure 1C was mistaken and misused. The corrected Figure 1 appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 1

SNORA72 is overexpressed in ovarian cancer stems cells (OCSCs). (A) Morphology of OVCAR-3 (OV), OVCAR-3 spheroid (OS), CAOV-3 (CA), and CAOV-3 spheroid (CS) cells shown under a microscope (x10). (B) Expression of CD133 detected by flow cytometry in OV vs. OS and in CA vs. CS cells. (C) Migration abilities of OV, OS, CA, and CS cells by Transwell assay. (D) Differentiation morphology of OS and CS cells at 0, 12, 24, 48, and 72 h. (E) Hierarchical clustering analysis of small nucleolar RNA (snoRNA) expression from non-coding RNA-ChIP data in OV and OS cells. Red, higher expression levels; green, lower expression levels. (F) Relative SNORA72 expression to U6, as an endogenous control, analyzed by qRT-PCR in OV and OS cells. The SNORA72 expression in OV cells was set as 1. Data are shown as the mean ± SD from three independent experiments. **p < 0.01, ****p < 0.0001.