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# Corrigendum: Comprehensive pan-cancer analysis of the prognostic and immunological roles of the METTL3/lncRNA-SNHG1/miRNA-140-3p/UBE2C axis

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## KEYWORDS

epigenetics, SNHG1, miRNA-140-3p, UBE2C, pan-cancer, prognosis, immunotherapy, drug sensitivity

## A Corrigendum on Comprehensive pan-cancer analysis of the prognostic and immunological roles of the METTL3/lncRNA-SNHG1/miRNA-140-3p/UBE2C axis

by Jiang X, Yuan Y, Tang L, Wang J, Liu Q, Zou X and Duan L (2021). *Front. Cell Dev. Biol.* 9:765772.  
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In the published article, there was an error in [Figure 9E](#) as published. Due to negligence and lack of careful inspection, and this part of the sequence is very similar. Therefore, some base mapping errors occurred, the inaccurate WT sequence for SNHG1 is: UCUUUAUCUUGAGCUGUGGUA (In [Figure 9E](#)). The corrected WT sequence for SNHG1 is: AUUUUUCUACUGCUCGUGGUA. The [Figure 9E](#) caption was correct. The corrected [Figure 9E](#) appear below.

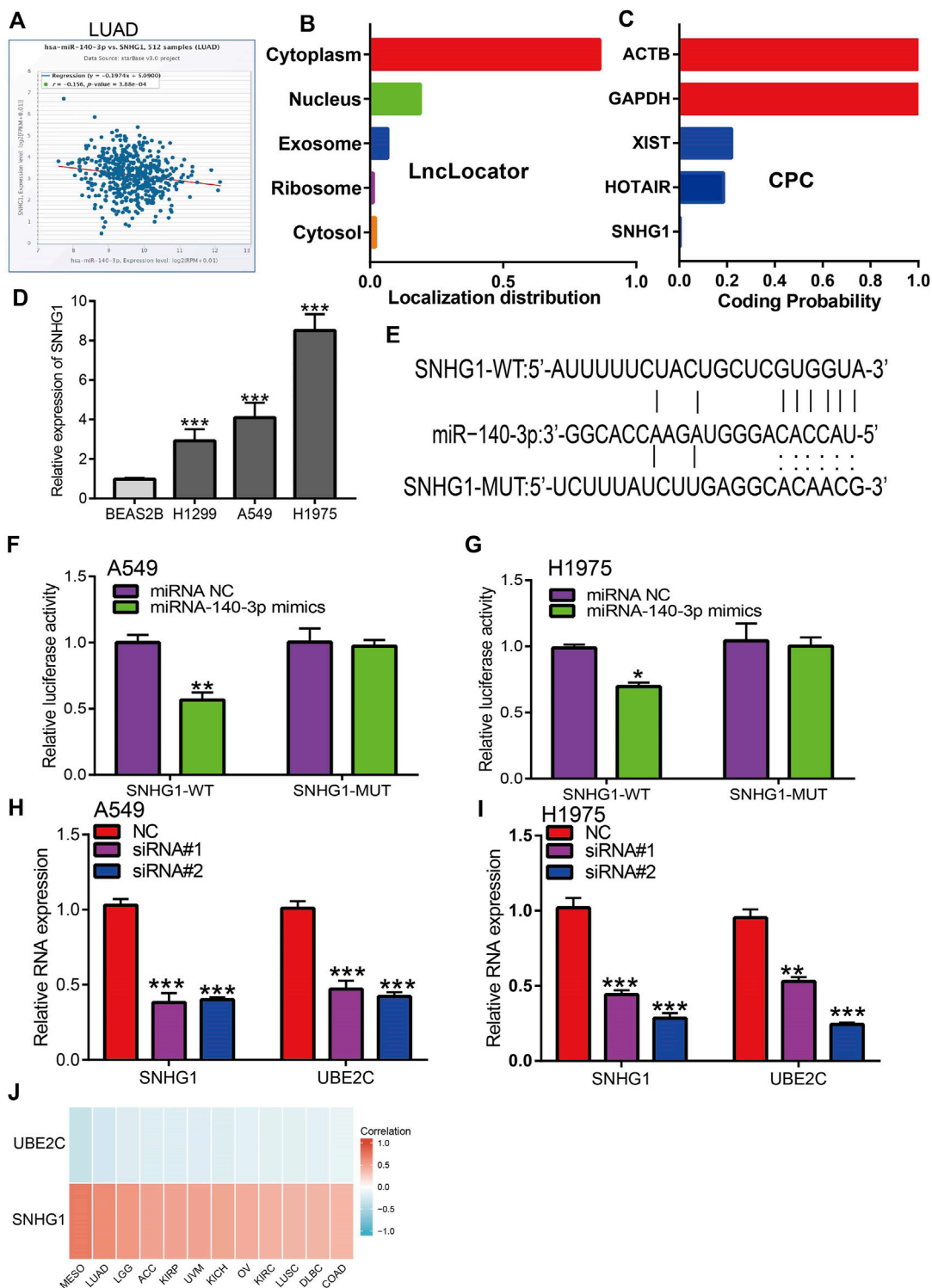
In the published article, there was an error in the **Materials and methods**, Quantitative Real-Time PCR section: primer sequences: SNHG1-F: AGCATCCACGAGCAAGAGAC, SNHG1-R: GATGCTACTAGTGTGCGGG.

A correction has been made to primer sequences: SNHG1-F: GCATCTCATAATCTA TCCTGG, SNHG1-R: CCTAGTTTCTCCTCAAACCTCCT. This sentence previously stated: “SNHG1-F: AGCATCCACGAGCAAGAGAC, SNHG1-R: GATGCTACTAGTGTG GCGGG”

The corrected sentence appears below:

“SNHG1-QPCR-F-GCATCTCATAATCTATCCTGG, SNHG1-QPCR-R-CCTAGTTTCTCCTCAAACCTCCT”

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



**FIGURE 9** Analysis of upstream lncRNA of miRNA-140-3p in pan-cancer. **(A)** The correlation between the SNHG1 and miRNA-140-3p in LUAD analysis by Starbase. **(B)** The subcellular localization of SNHG1 examined by LncLocator. **(C)** The coding potential of SNHG1 analysis by coding the potential calculator. **(D)** The expression of SNHG1 in NSCLC cell lines examined by qRT-PCR assay. **(E)** The target sites between miRNA-140-3p and SNHG1 was predicted by Starbase. **(F,G)** Relative luciferase activities of wild-type (WT) and mutated (MUT) SNHG1 reporter plasmid in A549 and H1975 cells co-transfected with miR-140-3p mimics examined by luciferase reporter assay. **(H,I)** The expression of UBE2C after depletion of SNHG1 in NSCLC cell lines examined by qRT-PCR assay. **(J)** The correlation between miRNA-140-3p and UBE2C, SNHG1 in NSCLC examined by Starbase. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

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