

Author Correction: Long non-coding RNA HEIH modulates CDK8 expression by inhibiting miR-193a-5p to accelerate nasopharyngeal carcinoma progression

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After publication, the authors found some mistakes in the article.

The Publisher apologizes for any inconvenience this may cause.

The correct sections are reproduced below.

- Figures 1, 2, and 4 report some wrong data. The authors have checked their raw data and realized that they had chosen the incorrect data for the analysis.
- In the paragraph entitled “CDK8 Was a Direct Target Gene of MiR-193a-5p in NPC”, “R2=0.6701, Figure 4G” was changed into “R2=0.6490, Figure 4G”.

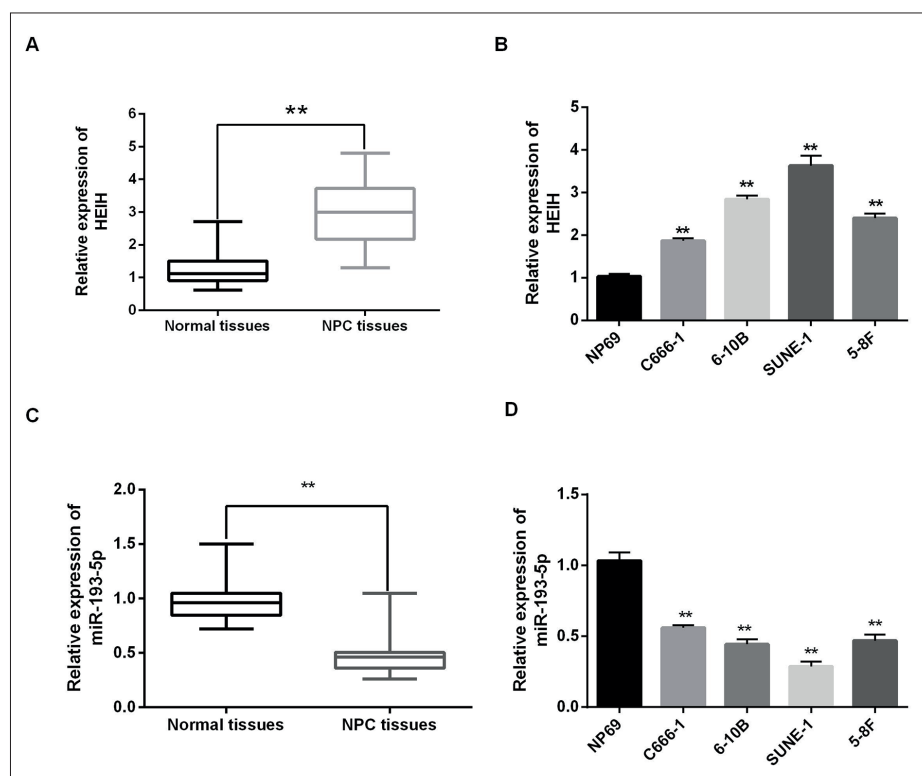


Figure 1. The expression of HEIH and miR-193a-5p were found in NPC. **A**, The expression of HEIH in NPC tissues (n=46). **B**, HEIH expression in NPC cell lines (SUNE-1, 6-10B, 5-8F and C666-1) and NP69 cells. **C**, MiR-193a-5p expression in NPC tissues. **D**, MiR-193a-5p expression in NPC cell lines (SUNE-1, 6-10B, 5-8F and C666-1) and NP69 cells. ** $p < 0.01$.

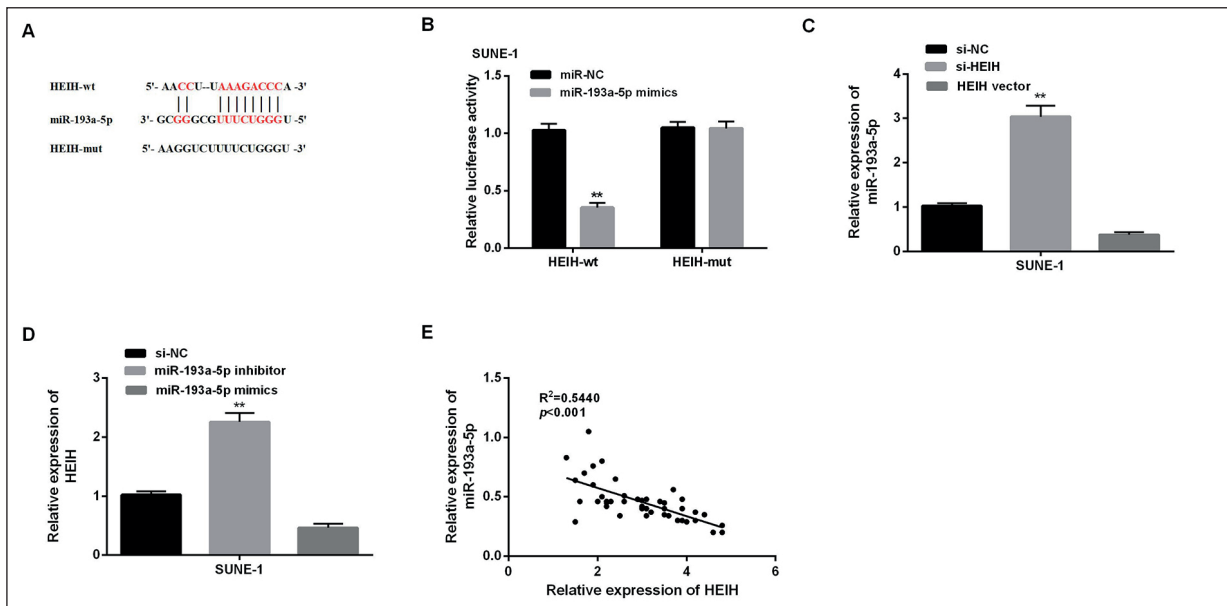


Figure 2. HEIH played as a sponge of miR-193a-5p in NPC. **A**, The binding sites between HEIH and miR-193a-5p. **B**, Dual-Luciferase assay. **C**, The expression of miR-193a-5p with si-HEIH and HEIH vector in SUNE-1 cells. **D**, HEIH expression regulated by miR-193a-5p mimic or inhibitor. **E**, HEIH negatively regulated miR-193a-5p expression in NPC tissues. ****** $p < 0.01$.

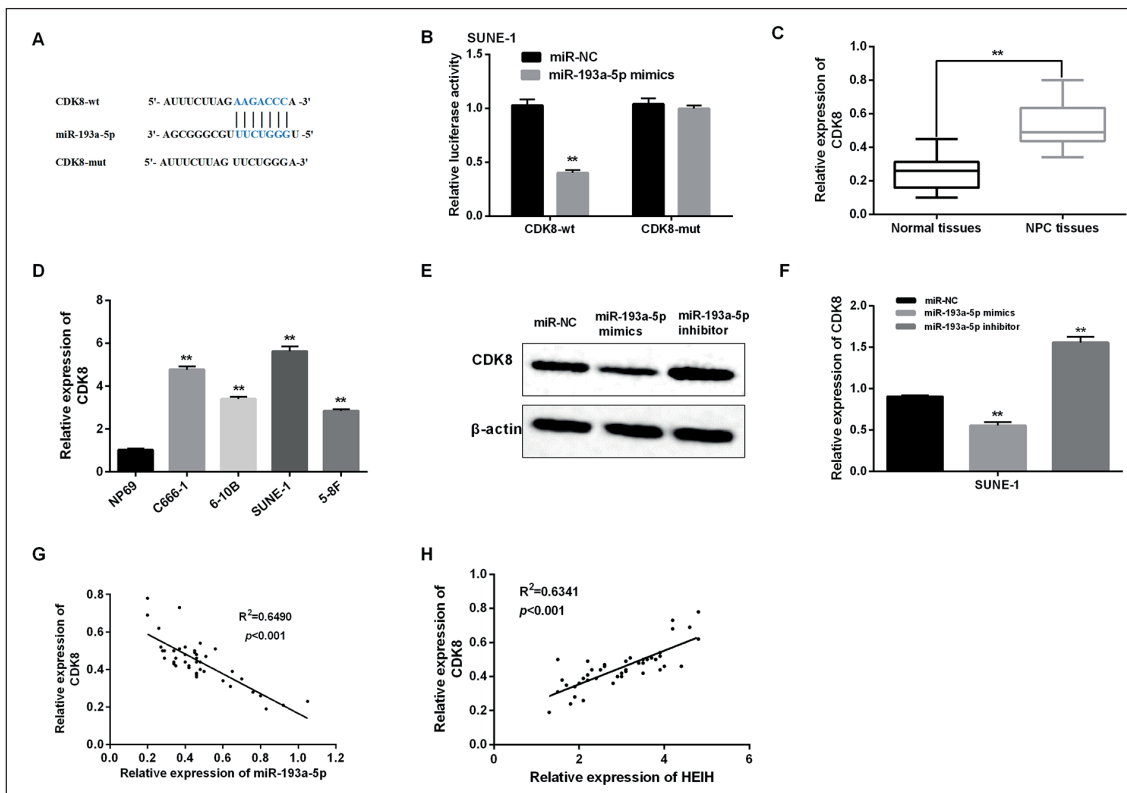


Figure 4. CDK8 was a target gene of miR-193a-5p. **A**, The binding sites between miR-193a-5p and CDK8. **B**, Dual-Luciferase reporter assay. **C**, The expression of CDK8 in NPC tissues. **D**, CDK8 expression in NPC cell lines and NP69 cells. **E-F**, CDK8 mRNA expression and protein expression in SUNE-1 cells with miR-193a-5p mimic or inhibitor. **G**, CDK8 was negatively correlated with miR-193a-5p in NPC tissues. **H**, CDK8 was positively correlated with HEIH in NPC tissues. ****** $p < 0.01$.