

Author Correction: LncRNA HOTAIRM1 inhibits the progression of hepatocellular carcinoma by inhibiting the Wnt signaling pathway

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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.

- In the Abstract section, the “progression-free survival (PFS)” is not correct, it should be “overall survival”.
- In the Patient and Methods section, “Tumor tissues and the corresponding paracancerous tissues of HCC patients with surgical resection in our hospital from July 2012 to February 2017 were collected” should be “Tumor tissues and the corresponding paracancerous tissues of HCC patients with surgical resection in our hospital from July 2002 to February 2007 were collected”.
- In the Results section, “Higher expressions of lncRNA HOTAIRM1 were observed in HCC tissues than those of the paracancerous tissues” should be “Lower expressions of lncRNA HOTAIRM1 were observed in HCC tissues than those of the paracancerous tissues”.

Abstract. – **PATIENTS AND METHODS:** LncRNA HOTAIRM1 expressions in 30 pairs of hepatocellular carcinoma tissues and paracancerous tissues were detected by quantitative Reverse Transcriptase-Polymerase Chain Reaction (qRT-PCR). The survival analysis and receiver operating characteristic (ROC) curve were introduced to explore the relationship between lncRNA HOTAIRM1 expressions and prognosis of HCC patients. The correlation between overall survival and clinical variables of HCC patients was estimated by single-factor and multiple-factor regression analysis, respectively.

Patients and Methods

Collection and Preservation of HCC Specimens

Tumor tissues and the corresponding paracancerous tissues of HCC patients with surgical resection in our hospital from July 2002 to February 2007 were collected. Enrolled patients were pathologically diagnosed as HCC and did not receive any preoperative treatment.

Results

LncRNA HOTAIRM1 Was Down-Regulated in HCC

We detected the lncRNA HOTAIRM1 expressions in 30 pairs of HCC and paracancerous tissues by qRT-PCR. Lower expressions of lncRNA HOTAIRM1 were observed in HCC tissues than those of the paracancerous tissues ($p < 0.001$, Figure 1A, 1B).