



Long noncoding RNA promotes oncogenic pathway to drive stem cell behaviors of hepatocellular carcinoma

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Abstract

Hepatocellular carcinoma (HCC) is a notoriously malignant cancer in the liver that causes high mortality and morbidity. The poor prognosis of HCC patients is largely attributed to recently recognized stem cell behaviors of HCC. Notably, a recent study revealed a novel mechanism by which long noncoding RNA promotes oncogenic pathway to drive stem cell behaviors of HCC. Kinesin Family Member 9-Antisense RNA 1 (KIF9-AS1) is a long noncoding RNA significantly upregulated in HCC patients and its level is correlated with poor prognosis. Furthermore, m6A writer METTL3 is upregulated in HCC patients and promotes the m6A methylation and stability of KIF9-AS1. The stabilized KIF9-AS1 then promotes ubiquitin-specific peptidase 1 mediated deubiquitination of short stature homeobox (SHOX2). The stabilized SHOX2 then drives the transcription of key genes involved in Wnt/ β -catenin pathway, TGF β signaling pathway and other unclear pathways to promote stem cell behaviors of HCC. These findings bring new hope to reduce HCC mortality and morbidity by revealing METTL3/KIF9-AS1/SHOX2 axis as prognostic markers and therapeutic targets.

Keywords: Hepatocellular carcinoma; KIF9-AS1; METTL3; ubiquitin-specific peptidase 1; SHOX2; stem cell

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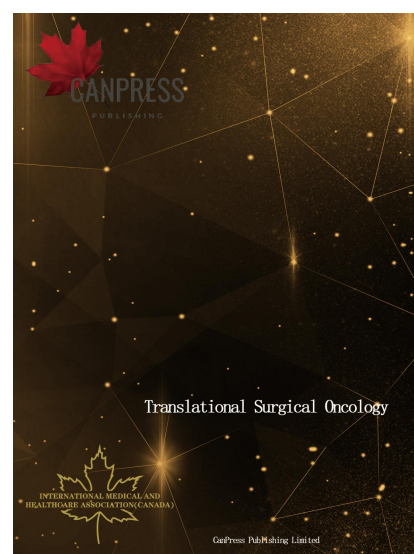
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Hepatocellular carcinoma (HCC) is a notoriously malignant cancer in the liver that causes high mortality and morbidity. At present, HCC is the fifth most common cancer and the fourth leading cause of cancer-related death worldwide. Hepatitis B virus (HBV) infection is one of the most important risk factors of HCC. Therefore, HCC has been widespread in China in the past decades, mainly due to high prevalence of HBV infection in China [1-3]. With the implement of HBV vaccination and the development of novel anti-HBV drugs, the incidence of HCC has reduced significantly in the past years. However, other risk factors such as obesity, diabetes, and metabolic syndrome cause increasing cases of HCC recently [4,5].

Currently, most common treatments for HCC include surgery and chemotherapy, and immunotherapy has recently been applied to improve the efficacy of HCC therapy [6-9]. In

particular, sorafenib is an effective first-line chemotherapy for HCC due to the ability of sorafenib to inhibit many kinases involved in the progression of HCC. Unfortunately, the prognosis of HCC patients remains poor due to recently recognized stem cell behaviors of HCC, including the development of drug resistance to sorafenib. The "stem cell behaviors" refer to a subset of cancer cells within HCC called "liver cancer stem cells" (LCSCs) that exhibit properties similar to normal stem cells, including self-renewal, the ability to differentiate into various cancer cell types, resistance to therapy, and the potential to initiate tumor growth, and act as a key driver of tumor progression, metastasis, and recurrence in HCC [10]. Indeed, genes that regulate stemness of HCC have been reported to be associated with the prognosis in HCC patients [11,12].



Interestingly, a recent study showed that a long noncoding RNA called Kinesin Family Member 9-Antisense RNA 1 (KIF9-AS1) was associated with the prognosis of HCC patients and it regulated stem cell behaviors of HCC via a novel mechanism [13]. In detail, the authors found that KIF9-AS1 level was significantly higher in HCC patients than controls, and KIF9-AS1 level was correlated with poor prognosis of HCC patients. Moreover, the expression of m6A writer METTL3 was higher in HCC cells, which could promote m6A modification of KIF9-AS1. Consequently, IGF2BP1 as an m6A reader stabilizes m6A-marked KIF9-AS1 to increase the stability of KIF9-AS1 in HCC cells.

Furthermore, the authors revealed that KIF9-AS1 upregulated ubiquitin-specific peptidase 1 (USP1) to enhance the stability of short stature homeobox (SHOX2). SHOX2 is a significant transcriptional factor that regulates development. Recently, SHOX2 has been proposed as a marker for the diagnosis and prognosis of various types of cancer, including HCC [14-16]. Mechanistically, SHOX2 promotes tumorigenesis via Wnt/ β -catenin and TGF β pathways [17,18]. Therefore, this study provides novel insight into the role of METTL3/KIF9-AS1/SHOX2 axis in the regulation of stem cell behaviors of HCC (Figure 1). METTL3 promotes the m6A methylation of KIF9-AS1 to stabilize it in IGF2BP1 dependent manner. The stabilized KIF9-AS1 then upregulates USP1 in an unclear mechanism. The upregulated USP1 then inhibits the ubiquitination and degradation of SHOX2. The stabilized SHOX2 then drives the transcription of key genes involved in Wnt/ β -catenin pathway, TGF β signaling pathway and other unclear pathways to promote stem cell behaviors of HCC, including but not limited to cell stemness and resistance to chemotherapy. Further studies are needed to explore the unclear mechanism or pathways highlighted in Figure 1 to get better understanding of the role of METTL3/KIF9-AS1/SHOX2 axis in HCC.

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Conflict of interest

The authors declare that they have no conflict of interest.

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