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MicroRNA-10a通过靶向作用E2F3抑制肝癌细胞的增殖 [点此下载全文](#)

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摘要:

目的: 探讨微小RNA-10a (microRNA-10a, miR-10a) 对肝癌细胞增殖的影响及其作用机制。方法: 收集广西医科大学附属肿瘤医院肿瘤科2001年10月至2005年7月144例肝癌患者手术切除的肝癌组织和癌旁组织(距癌灶组织边缘2~5 cm)标本, Real-time PCR法分析144例肝癌组织及癌旁组织中miR-10a的表达量。在肝癌细胞(QGY-7701、Huh7、PCL/PRF/5)中转染miR-10a模拟物, Real-time PCR法检测转染后细胞miR-10a的表达水平; CCK-8法检测过表达miR-10a的肝癌细胞的增殖水平, 流式细胞术检测过表达miR-10a的肝癌细胞的凋亡和细胞周期; 生物信息学预测并以Western blotting检测过表达miR-10a的肝癌细胞中转录因子E2F3的表达量。结果: 与癌旁组织相比, 肝癌组织中的miR-10a显著低表达 $[-9.89 \pm 1.68]$  vs  $[-7.84 \pm 1.97]$ ,  $P = 0.000$ 。转染miR-10a模拟物后肝癌细胞系中miR-10a的表达量是转染对照小RNA组或空白组细胞的16倍左右。过表达miR-10a可显著抑制7种肝癌细胞(QGY-7701、QGY-7703、Huh7、PCL/PRF/5、HepG2、Bel-7402、SMMC-7721)的增殖(均  $P < 0.05$ ), 并引起肝癌细胞细胞周期G<sub>1</sub>/S期阻滞, 但并不能诱导肝癌细胞发生凋亡。生物信息学预测显示E2F3是miR-10a可能的靶分子, Western blotting检测显示过表达miR-10a可明显抑制肝癌细胞中E2F3的表达 $[0.50 \pm 0.12]$  vs  $[0.79 \pm 0.21]$ ,  $P < 0.05$ 。结论: 人肝癌组织中低表达miR-10a, 转染miR-10a模拟物后多种肝癌细胞的增殖均受到明显抑制, 其机制可能与miR-10a靶向作用转录因子E2F3并阻滞肝癌细胞细胞周期于G<sub>1</sub>/S期有关。

关键词: [肝癌](#) [微小RNA-10a](#) [增殖](#) [转录因子](#) [E2F3](#) [G<sub>1</sub>/S期阻滞](#)

MicroRNA-10a inhibits hepatocellular carcinoma cell proliferation through targeting E2F3 [Download Fulltext](#)

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Abstract:

Objective: To investigate the role of microRNA-10a (miR-10a) in hepatocellular carcinoma (HCC) growth. Methods: Paired HCC and adjacent non-tumor tissue specimens were surgically collected from 144 patients who were diagnosed with primary HCC in Guangxi Medical University-Affiliated Tumor Hospital between October 2001 and July 2005. HCC QGY-7701, Huh7, and PCL/PRF/5 cells were transfected with miR-10a mimics or scramble control miRNA. The abundance of miR-10a in both tissue specimens and transfected cells was quantified by real-time PCR and E2F3 protein in transfected cells was assessed by Western blotting. Proliferation of the transfectants was assessed by a colorimetric cell counting assay. Cell cycle progression and apoptosis of the transfectants were assessed by FACS. Results: The abundance of miR-10a mRNA was significantly lower in HCC tissue specimens than in normal tissue specimens  $(-9.89 \pm 1.68$  vs  $-7.84 \pm 1.97$ ,  $P = 0.0001$ ). HCC cells transfected with miR-10a mimics had miR-10a abundance 16 times higher than both wild-type HCC cells and HCC cells transfected with the control miRNA with scrambled sequences. Overexpression of miR-10a resulted in significant increases in suppression of HCC cell proliferation ( $P < 0.05$ ) and G<sub>1</sub> phase arrest. In contrast, overexpression of miR-10a had no influence on apoptosis of HCC cells. Bioinformatics suggested that transcription factor E2F3 might be a downstream target of miR-10a and the expression of E2F3 in HCC cells transfected with miR-10a was significantly lower than in wild-type HCC cells and HCC cells transfected with the control miRNA  $(0.50 \pm 0.12$  vs  $0.79 \pm 0.21$ ,  $P < 0.05$ ). Conclusion: MiR-10a may suppress HCC cell proliferation through G<sub>1</sub> phase arrest in an E2F3-dependent mechanism.

Keywords: [hepatocellular carcinoma](#) [microRNA-10a \(miR-10a\)](#) [transcription factor](#) [E2F3](#) [proliferation](#) [G<sub>1</sub>/S phase arrest](#)

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