

miR-106a负调控PTEN促进骨肉瘤细胞增殖并抑制凋亡

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Title: miR-106a promoted osteosarcoma cell proliferation and inhibited apoptosis via negatively regulating the expression of PTEN

作者: 曾桂平¹; 文昌明¹; 王聪²; 方忠³

1湖北省阳新县人民医院骨二科; 2手术室, 湖北 阳新 435200; 3华中科技大学同济医学院附属同济医院骨科, 湖北 武汉 430030

Author(s): Zeng Guiping¹; Wen Changming¹; Wang Cong²; Fang Zhong³

1.Second Department of Orthopaedics; 2.Operation room, Yangxin People's Hospital, Hubei Yangxin 435200, China; 3.Department of Orthopaedics, Tongji Hospital Affiliated to Tongji Medical College, Huazhong University of Science and Technology, Hubei Wuhan 430030, China.

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摘要: 目的:研究miR-106a在骨肉瘤组织和MG-63细胞中的表达水平及其对MG-63细胞增殖和凋亡的影响及机制。方法:用荧光实时定量PCR法检测20对骨肉瘤和相邻正常组织及MG-63和成骨细胞hFOB 1.19中miR-106a的表达。用miR-106a mimics、miR-106a antagomir及两者相应的对照物转染MG-63细胞,然后分别用CCK-8法检测四组细胞增殖活性和FCM法检测细胞凋亡率。miR-106a mimics和mimics control与野生型或突变型PTEN 3'-UTR重组载体共转染后,应用荧光素酶基因报告系统检测miR-106a是否与PTEN基因3'-UTR结合。利用Western blot技术检测PTEN蛋白在上述四组转染MG-63细胞和骨肉瘤标本中的表达水平。结果:与相邻正常组织(1.19±0.15)相比,肿瘤组织miR-106a的表达水平(2.60±0.86)显著升高;同时,miR-106a在MG-63中的表达水平(2.60±0.92)明显高于hFOB 1.19(1.19±0.39),以上差异均有统计学意义(P<0.05)。CCK-8和FCM检测结果显示,与mimics control组相比,miR-106a mimics组的增殖率明显增加,而细胞凋亡率下降;反之,miR-106a antagomir组与antagomir control组相比,增殖率前者低于后者,而凋亡率前者高于后者,上述差异均有统计学意义(P<0.05)。荧光素酶报告实验显示,miR-106a mimics和wt PTEN 3'-UTR共转染组的荧光强度值明显低于mimics control和wt PTEN 3'-UTR组(P<0.05)。Western blot发现,与对照组相比,miR-106a mimics组PTEN表达下调,而miR-106a antagomir表达上调;临床标本,肿瘤组织PTEN表达明显低于正常组织,差异均有统计学意义(P<0.05)。结论:miR-106a在骨肉瘤组织及细胞中过表达,并靶向负调控PTEN表达,促进骨肉瘤细胞增殖并抑制其凋亡,从而发挥促癌作用。因此,miR-106a可为骨肉瘤的诊治提供新的潜在分子靶点。

Abstract: Objective: To study the expression level of miR-106a in osteosarcoma and MG-63 cells and its effect on proliferation and apoptosis of MG-63 cells and the underlying mechanism. Methods: The expression level of miR-106a in 20 pairs of osteosarcoma and adjacent normal tissues, MG-63 and osteoblasts hFOB 1.19 was detected by real-time quantitative PCR. MG-63 cells were transfected with miR-106a mimics, mimics control, miR-106a antagomir and antagomir control, respectively. Then the cell viability of four groups was detected by CCK-8 and their apoptosis rate was detected by FCM. After co-transfection of miR-106a mimics and mimics control with wild or mutant PTEN 3'-UTR recombinant vectors, the luciferase gene report system was used to detect whether miR-106a was combined with the 3'-UTR of gene PTEN. The expression level of PTEN protein in the four groups of MG-63 cells transfected and osteosarcoma samples was detected by Western blot. Results: Compared with the adjacent normal tissues (1.19±0.15), the relative expression level of miR-106a in tumor tissues (2.60±0.86) was significantly higher. Meanwhile, the expression level of miR-106a in MG-63 (2.60±0.92) was significantly higher than that of hFOB 1.19 (1.19±0.39), and the above differences were statistically significant (P <

0.05).The results of CCK-8 and FCM assay showed that compared with the mimics control group,the proliferation rate of miR-106a mimics group increased significantly,while its apoptosis rate decreased.On the contrary,the proliferation rate in miR-106a antagomir group was obviously lower than that of antagomir control group,and its apoptosis rate was higher than that of the latter,and the all difference was statistically significant ($P < 0.05$).The fluorescence intensity of miR-106a mimics and wt PTEN 3'-UTR co-transfection group was significantly lower than that of mimics control and wt PTEN 3'-UTR group ($P < 0.05$).Western blot found that compared with the corresponding control group,the protein expression of PTEN in the miR-106a mimics group was significantly down-regulated,but the expression of PTEN in miR-106a antagomir was up-regulated,and the expression of PTEN in the tumor tissue was significantly lower than that of the adjacent normal tissue,and all difference was statistically significant ($P < 0.05$).Conclusion:miR-106a was overexpressed in the osteosarcoma tissues and MG-63 cells and negatively regulated the expression of PTEN,which promotes the proliferation of osteosarcoma cells and inhibits its apoptosis,and thus may play an important role in promoting tumorigenesis.Therefore,miR-106a likely becomes a new potential molecular target for the diagnosis and treatment of osteosarcoma.

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