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Clinical significance of microRNA-130b in osteosarcoma and in cell growth and invasion

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ABSTRACT

Objective: To investigate clinical significance of microRNA-130b (miR-130b) in osteosarcoma and its role in cell growth and invasion.

Methods: miR-130b expression was detected in 68 samples of surgically resected osteosarcoma and matched normal tumor-adjacent tissues by qRT-PCR. The expression of miR-130b was altered by corresponding vectors in osteosarcoma cells, and then Western blot was used to detect the expression of PPAR γ . BrdU cell proliferation and Transwell assays were performed to determine cell proliferation and invasion.

Results: The expression of miR-130b in osteosarcoma tissues was significantly higher than that in normal tumor-adjacent tissues. Its expression in patients with metastasis was significantly higher than that in those without metastases. miR-130b expression in tumor tissues was significantly associated with tumor size, clinical stage and distant metastasis. And its expression was significantly correlated with overall survival and disease free survival. miR-130b overexpression obviously repressed the expression of PPAR γ , and resulted in significant increase of Saos-2 cell proliferation and invasion. On the contrast, repressing miR-130b expression with its inhibitor significantly increased PPAR γ expression, and inhibited MG-63 cell proliferation and invasion.

Conclusions: The high-expression of miR-130b is correlated with the adverse clinicopathological features and poor prognosis in osteosarcoma. miR-130b may regulate proliferation and invasion of osteosarcoma cells by targeting PPAR γ , suggesting miR-130b may play a key role in the progression of osteosarcoma.

1. Introduction

Osteosarcoma is the most common type of primary malignant tumor of bone, accounting for 60% of all malignant bone tumors in teenage [1]. With the advancement of multiple therapeutic strategies for osteosarcoma including surgical resection, adjuvant chemotherapy and radiotherapy, the 5 year survival of the non-metastatic patients has improved to approximately 60%–70% [2,3]. However, for patients with metastasis or recurrence, the efficacy of chemotherapy is significantly decreased, and the long-term prognosis remains poor with a survival rate of 5%–20% [3]. Therefore, identifying novel biomarkers and clarifying the molecular mechanisms of the metastasis and recurrence of osteosarcoma may greatly improve the prognosis of osteosarcoma patients.

MicroRNAs (miRNAs) are a large group of non-coding RNAs containing 18–25 nucleotides and post-transcriptionally regulate the expression of multiple genes by interacting with their 3'-untranslated regions. Numerous deregulated miRNAs have been found to play fundamental roles in the development and progression of malignant diseases [4–10] including cell growth, differentiation, invasion, metastasis, and angiogenesis. Studies of melanoma [11], gastric carcinoma [12,13], bladder cancer [14] and colorectal cancer [15] have shown that aberrantly elevated miR-130b expression in these malignancies was correlated with poor prognosis of patients, indicating an oncogenic role of miR-130b in human cancers. In colorectal cancer, miR-130b has been found to promote the migration and invasion of cancer cells by targeting PPAR γ [15]. However, the clinical significance of miR-130b expression in osteosarcoma tissues and its functional role on the migration and invasion of osteosarcoma cells have not been examined.

In this study, we evaluated the expression of miR-130b in osteosarcoma tissues by qPCR. Our results confirmed that miR-130b expression was significantly increased in osteosarcoma tissues. And its elevated expression was closely associated with

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poor clinical features of osteosarcoma patients including large tumor size, advanced TNM stage, and distal metastasis. Furthermore, *in vitro* experiments showed that miR-130b could promote the progression of osteosarcoma by promoting cell proliferation and invasion. This study proposes that miR-130b may be a promising biomarker and attractive therapeutic target of osteosarcoma.

2. Material and methods

2.1. Patients and specimens

Cancerous tissues and adjacent non-tumor tissues (>3 cm distance to the resection margin) were collected from 68 patients who underwent curative resection of osteosarcoma. The clinicopathological data of all enrolled patients were demonstrated in Table 1. All clinical samples were used after obtaining informed consent. All patients did not receive any blood transfusion, radiation treatment, or chemotherapy. The protocols of this study have been approved by the Ethics Committee of University.

2.2. Real-time quantitative PCR

Total RNA was isolated from clinical tissues by TRIZOL[®] reagent (Life Technologies, Carlsbad, CA, USA) under the guidance of operation instructions. The first strand cDNA was synthesized by the Revertid[™] First Strand cDNA Synthesis Kit (TaKaRa, Shiga, Japan). Two microliters of cDNA obtained from each sample was amplified and quantified by real-time PCR of TaqMan Human MiRNA Assay (Applied Biosystems). miR-130b expression relative to U6 was calculated by the method of $2^{-\Delta\Delta Ct}$.

2.3. Cell culture

Human osteosarcoma cell lines, Saos-2 and MG-63, were purchased from American Type Culture Collection, and was cultured in Dulbecco's modified Eagle medium (DMEM, Gibco, USA), containing 10% fetal bovine serum (Gibco) in a humidified 5% CO₂ incubator at 37 °C. Cells at logarithmic growth phase were collected for further experiments.

2.4. Transfection of miRNA mimics and inhibitors

MiRNA vectors including miR-130b expression vector (miR10004680-1-5), the control vector for miR-130b (miR01201-1-2), miR-130b inhibitor (miR20004680-1-5) and the negative control for the miR-130b inhibitor (miR02201-1-2) bought from RiboBio (RiboBio, Guangzhou, China). Cells were seeded at 1×10^5 cells per well in a six-well plate and transfected with miRNA vectors at a final concentration of 50 nM by Lipofectamine 2000 (Invitrogen, Carlsbad, CA) under the guidance of operation instructions. Total RNA and protein were collected 3 d post-transfection for experimental analyses.

2.5. Western blot

RIPA buffer (50 mM Tris pH 7.5, 150 mM NaCl, 1% TritonX-100, 5 mM ethylenediaminetetraacetic acid) was used to extract the proteins of cell lysates. Protein concentration was determined using the BCA Kit (Pierce, IL, USA). Protein samples (20 µg) were separated by sodium dodecyl sulfate–polyacrylamide gel electrophoresis and transferred to a nitrocellulose membrane. The blots were then probed with antibodies against PPAR-γ (1:1 000; Cell Signaling Technology, MA, USA) and β-actin (1:1 500, Boster, Wuhan, China). Blots were then incubated with the appropriate fluorescent secondary antibody (1:5 000; Boster). Images were acquired by the Bio-Rad Gel imaging system and analyzed by the software program as specified by Bio-Rad.

2.6. Cell proliferation assay

Saos-2 or MG-63 cells were seeded at 5×10^3 cells per well in a 96-well plate, and was transfected with miRNAs vectors. Twenty-four after transfection, cell proliferation ability was assessed with Proliferation ELISA, BrdU (5-bromodeoxyuridine) (chemiluminescent) (Roche, USA).

2.7. Transwell assay

The invasion assay was performed in 24-well Transwell units (BD Biosciences) with 8 µm porosity polycarbonate filters. Each

Table 1

Correlation between clinicopathological characteristics and expression of miR-130b in the osteosarcoma patients ($n = 68$).

Characteristics	No. of cases	miR-130b		P
		High expressing no.	Low expressing no.	
Gender	Male	40	29	0.108
	Female	28	15	
Age (years)	<24	49	34	0.194
	≥25	19	10	
Clinical stage	I	27	13	<0.001*
	II	34	24	
	III	7	7	
T classification	T1	26	12	0.012*
	T2	42	32	
M classification	M0	61	38	0.020*
	M1	7	6	
Histology	Conventional osteosarcoma	58	39	0.487
	Others	10	5	
Histological differentiation	G1	27	13	0.407
	G2	41	31	

* $P < 0.05$.

filter was coated with 70 μ L matrigel (BD Biosciences, Franklin Lakes, NJ, USA) at 1 mg/mL on the inner layer. 1.5×10^5 Saos-2 or MG-63 cells suspended in 200 μ L reduced serum DMEM medium were added into the upper chamber, and 800 μ L DMEM medium containing 20% FBS was added into the lower chamber. After incubating for 24 h, cells were fixed in 4% paraformaldehyde for 3 min, and then permeabilized in methanol for 20 min. Cells on the inner layer were removed with a cotton swab, and the adherent cells on undersurface of the insert were stained with 0.3% crystal violet dye for 10 min. The filters were washed with PBS and images were taken. Invaded cells on undersurface were counted under a light microscope.

2.8. Statistical analysis

All data are presented as mean \pm SEM. The GraphPad Prism 5 software (GraphPad Software, Inc., San Diego, CA, USA) was used for Pearson chi-square test and Two-tailed Student's *t* test. Kaplan–Meier method was employed to plot overall survival and disease-free survival curves, and the prognostic effect of miR-130b was evaluated using the log-rank test. $P < 0.05$ was considered statistically significant.

3. Results

3.1. Elevated expression of miR-130b expression in osteosarcoma tissues

Compared with that in matched non-tumor tissues, miR-130b expression in tumor tissues was significantly elevated ($P < 0.05$,

Figure 1A). The level of miR-130b in metastases group was significantly higher than that in non-metastases group ($P < 0.05$, Figure 1B). These results suggest that miR-130b plays an oncogenic role in osteosarcoma and is involved in the metastasis of osteosarcoma.

3.2. Correlation between clinicopathological characteristics and expression of miR-130b

As shown in Table 1, clinical association analysis by the Pearson chi-square test demonstrated that increased expression of miR-130b was significantly correlated with large tumor size ($P = 0.020$), advanced tumor stage ($P < 0.001$) and distal metastasis ($P = 0.020$). These results indicate increased expression of miR-130b is correlated with poor clinicopathological features in osteosarcoma. Patients with high miR-130b level had significantly decreased overall survival ($P < 0.05$, Figure 2A) and disease free survival ($P < 0.05$, Figure 2B).

3.3. Overexpression of miR-130b promotes proliferation and invasion of osteosarcoma cells

As shown in Figure 2, transfection of miR-130b expressing plasmid resulted in significantly increased level of miR-130b in Saos-2 cells (Figure 3A, $P < 0.05$). Moreover, PPAR γ , a well-defined downstream target of miR-130b in colorectal cancer cells, was significantly down-regulated accordingly (Figure 3B, $P < 0.05$). And Brdu incorporation assay and transwell assay demonstrated overexpression of miR-130b in Saos-2 cells resulted in significantly increased proliferation (Figure 3C,

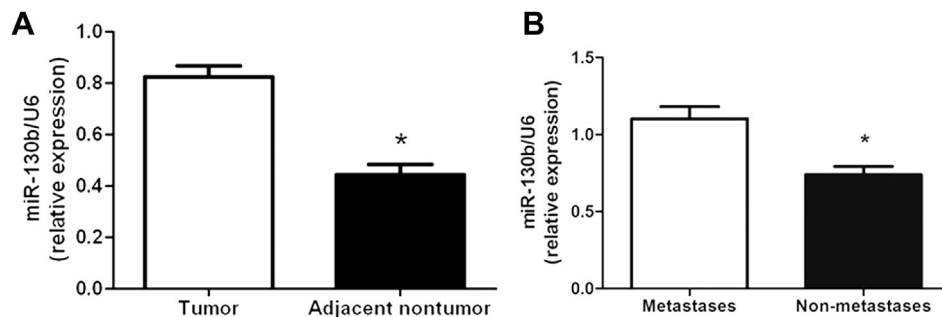


Figure 1. Expression level of miR-130b in osteosarcoma tissues.

A) osteosarcoma tissues and matched non-tumor tissues; B) osteosarcoma tissues arising from metastases and non-metastases groups. * $P < 0.05$.

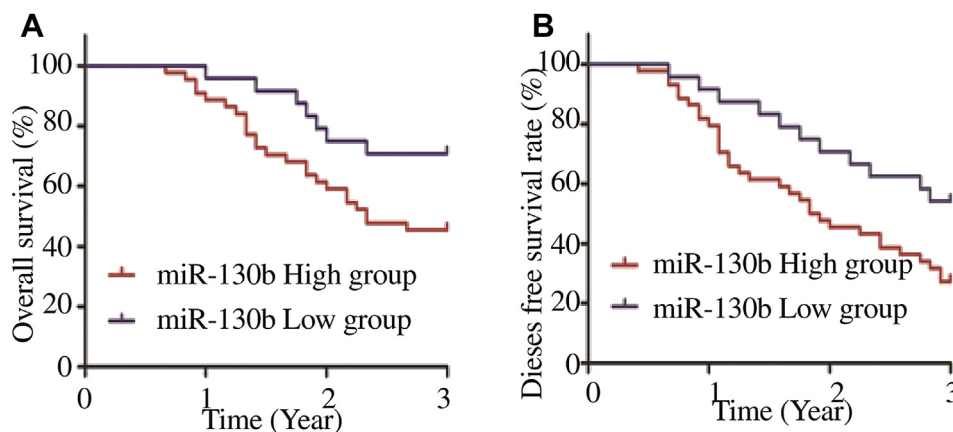


Figure 2. Kaplan–Meier survival curves for osteosarcoma patients based on miR-130b level.

A) Overall survival rate ($P < 0.05$). B) Survival rate ($P < 0.05$).

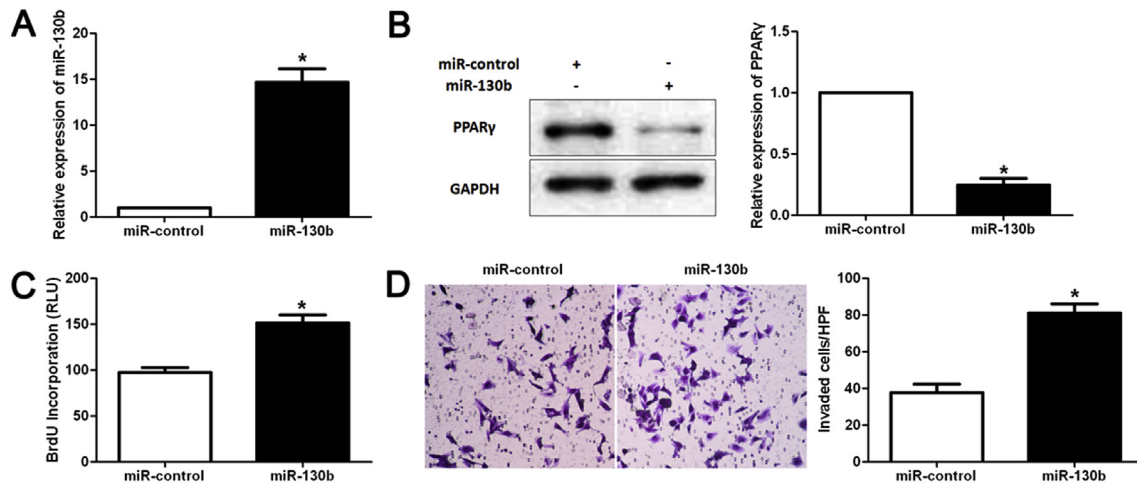


Figure 3. PPAR γ expression, cell proliferation and invasion in Saos-2 cells.

A) miR-130b level in Saos-2 cells. $n = 6$, $*P < 0.05$; B) PPAR γ expression in Saos-2 cells. $n = 6$, $*P < 0.05$; C) Cell proliferation and invasion in Saos-2 cells. $n = 6$, $*P < 0.05$; D). Cell invasion in Saos-2 cells. $n = 6$, $*P < 0.05$.

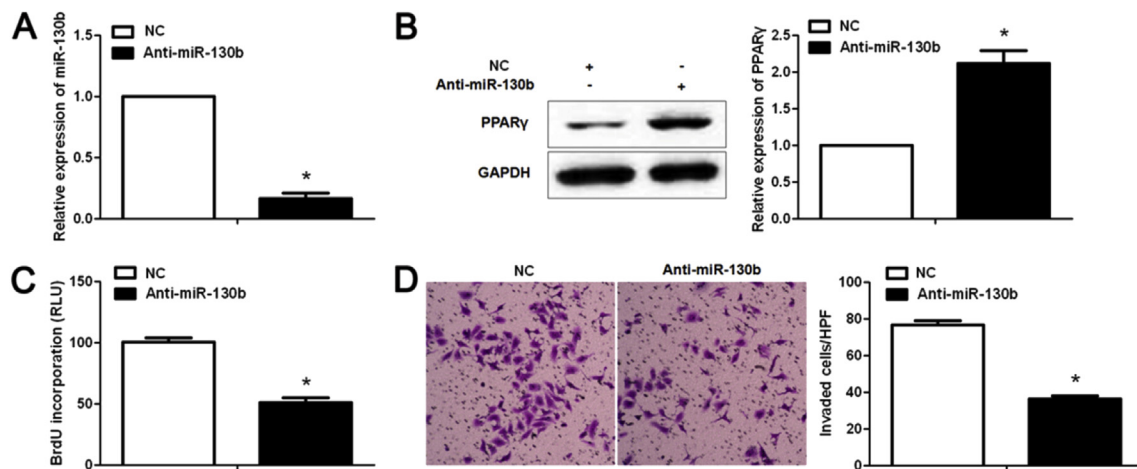


Figure 4. PPAR γ expression, cell proliferation and invasion in MG-63 cells.

A) miR-130b level in MG-63 cells. $n = 6$, $*P < 0.05$; B) PPAR γ expression in MG-63 cells. $n = 6$, $*P < 0.05$; C) Cell proliferation in MG-63 cells. $n = 6$, $*P < 0.05$; D) Cell invasion in MG-63 cells. $n = 6$, $*P < 0.05$.

$P < 0.05$) and increased invasion (Figure 3D, $P < 0.05$), respectively.

3.4. Down-regulation of miR-130b inhibits proliferation and invasion of osteosarcoma cells

The expression of miR-130b was significantly reduced after transfecting its inhibitor into MG-63 cells (Figure 4A, $P < 0.05$). And the expression of PPAR γ was significantly increased after down-regulating miR-130b (Figure 4B, $P < 0.05$). Functionally, after repressing the expression of miR-130b, decreased proliferation (Figure 4C, $P < 0.05$) and invasion (Figure 4D, $P < 0.05$) of MG-63 cells were observed.

4. Discussion

MiRNAs can regulate gene expression at post-transcriptional level by binding to 3'-untranslated regions of target messenger RNAs (mRNAs), and thus participate in many critical biological processes including cell differentiation, morphogenesis and tumorigenesis [16,17]. The critical role of miRNAs in the development and progression of human cancers have been

widely accepted. miRNAs can play either oncogenic role or tumor suppressive role in human malignancies [18–20]. Moreover, miRNAs have been regarded as promising biomarkers and attractive therapeutic targets for human cancers [21].

MiR-130b is being actively investigated in many kinds of human cancers [11–15,22–25]. It has been found to be down-regulated in papillary thyroid carcinoma [22], endometrial cancer [23], pituitary adenomas [24] and pancreatic cancer [25]. However, in melanoma [11], gastric carcinoma [12,13], bladder cancer [14], and colorectal cancer [15], miR-130b expression was found to be aberrantly elevated. Increased level of miR-130b in the serum of colon cancer patients was associated with resistance to chemotherapy [26]. And miR-130b has been regarded as an effective biomarker for HCC patients [27,28]. In this study, we evaluated the expression status of miR-130b in 68 pairs of osteosarcoma tissues and adjacent non-tumor tissues using qRT-PCR. The result revealed a significant increase of miR-130b expression in osteosarcoma tissues. Furthermore, we investigated the clinical significance of miR-130b in osteosarcoma. Clinical association analysis proved that miR-130b was closely associated with large tumor size, advanced TNM stage and distal metastasis. Moreover, increased level of miR-130b

was correlated with poorer prognosis of osteosarcoma patients. These results indicate that miR-130b serves as an oncogenic miRNA in osteosarcoma and can potentially act as a novel biomarker of the prognosis of osteosarcoma patients.

Previous study demonstrated that miR-130b could down-regulate *PTEN*, E-cadherin, Snail and VEGF by inhibiting PPAR γ , and thus resulted in increased proliferation, EMT and angiogenesis of colorectal cancer [15]. In endometrial cancer, miR-130b could contribute to EMT of cancer cells by targeting DICER1 [29]. This study demonstrated that overexpression of miR-130b in MG-63 cells could inhibit the expression of PPAR γ which was found to be down-regulated in osteosarcoma and could regulate proliferation, invasion and resistance to chemotherapy [30,31]. Accordingly, our functional experiments confirmed that overexpressing miR-130b led to increased proliferation and invasion of Saos-2 cells. On the other way, down-regulating miR-130b resulted in elevation of PPAR γ expression, and decreased proliferation and invasion in MG-63 cells. These results indicate that miR-130b may promote the growth and metastasis of osteosarcoma by repressing PPAR γ .

In summary, our results demonstrate that miR-130b is up-regulated in osteosarcoma, and its elevated expression is associated with large tumor size, distal metastasis and advanced TNM stage. Patients with increased level of miR-130b have a significantly poorer prognosis. Functionally, miR-130b may promote the proliferation and invasion of osteosarcoma cells by targeting PPAR γ , and thus contributes to the development and progression of osteosarcoma. This study indicates that miR-130b may be a promising biomarker and therapeutic target in osteosarcoma.

Conflict of interest statement

We declare that we have no conflict of interest.

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