Original Article

MicroRNA-665 suppressed the invasion and metastasis of osteosarcoma by directly inhibiting RAB23

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Abstract: MicroRNAs (miRNAs) are small, short and noncoding RNAs that regulate gene expression posttranscriptionally. Increasing evidences have demonstrated that deregulated expression of miRNAs is found in osteosarcoma. In this study, we demonstrated that miR-665 was downregulated in osteosarcoma tissues compared to non-tumor tissues. The overall survival (OS) of osteosarcoma patients with low miR-665 expression was lower than that of these patients with high miR-665 expression. Ectopic expression of miR-665 suppressed the osteosarcoma cell proliferation, EMT and invasion. We identified Rab23 as a direct target gene of miR-665. Rab23 was downregulated in osteosarcoma tissues and cell lines. The expression of miR-665 was inversely associated with the expression of Rab23 in the osteosarcoma tissues. These results suggested that miR-665 acted as a tumor suppressor gene in the development of osteosarcoma.

Keywords: Osteosarcoma, microRNAs, miR-665, Rab23

Introduction

Osteosarcoma, which is the most frequent tumor of bone, is the leading cause of cancer-related death in adolescents and children [1-4]. Despite the development treatment approaches of osteosarcoma such as chemotherapy and curative resection, the 5-year overall survival rate of osteosarcoma patients is still low [5-8]. Therefore, it is important to find new biomarkers for prognosis and diagnosis and a new therapy for osteosarcoma.

miRNAs (MicroRNAs) are non-coding, small, endogenous RNAs that induce mRNA translational inhibition and/or degradation by binding to 3’UTR (3’-untranslated region) of the target mRNAs [9-13]. Increasing evidences showed that play important roles in complicated cell processes including cell metabolism, proliferation, differentiation [14-16]. Moreover, deregualtion expression of miRNA has been found in multiple cancers such as gastric cancer, hepatocellular carcinoma, lung cancer, colorectal cancer, breast cancer and ovarian carcinoma [17-23]. Both the gain and loss of miRNAs expression contribute to tumor development through the downregulation of tumor suppressor genes and upregulation oncogenes [24-27].

In the present study, we demonstrated that miR-665 expression was downregulated in the osteosarcoma tissues compared to non-tumor tissues. Ectopic expression of miR-665 suppressed the osteosarcoma cell proliferation, EMT and invasion. We identified Rab23 as a direct target gene of miR-665.

Materials and methods

Clinical samples and cell lines

Thirty paired osteosarcoma tissues and matched non-tumor tissues were obtained from the Third Affiliated Hospital of Third Military Medical University. The tissues were obtained during surgery and frozen in liquid nitrogen immediately. This research was approved by the Ethics Committee of The Third Affiliated Hospital of Third Military Medical University and all patients have written informed consent. Osteosarcoma cell lines (U2OS, SOSP9607, Saos-2, and MG-63) and osteoblastic cell
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TRlzo1 reagent (TaKaRa, Japan) was used to extract total RNA. The miR-665 expression was measured by the TaqMan assay (Applied Biosystems, USA) and the expression of Rab23 was detected by the qRT-PCR. U6 or GAPDH was used for miRNA or Rab23 expression control respectively. The primers were used as follows: Rab23, sense 5’-AGCGAGAC- TCCGTCTTCAA-3’; antisense, 5’-CACCCCTAAAGG TACGCATGT-3’; GAPDH, sense 5’-GCACC- GTCAAGGCTG AGA AC-3’; antisense, 5’-GGTGAAGACGCAC- GTGGA-3’.

**Plasmid and transfection**

MiR-665 mimic and scramble oligonucleotide and pCDNA were synthesized by RiboBio (Guangzhou, China). 20 nM miR-665 mimic or scramble was transfected into cells used Lipofectamine 2000 (Invitrogen) according to manufacturer’s advising.

**Cell growth, colony formation assay and invasion assays**

The cell proliferation was detected by using CCK-8 (DOJINDO, Japan) following to manufacturer’s advising. The cell proliferation rate was measured at 0, 24, 48 and 72 hours. For colony formation assay, cells were cultured in Petri dish and were maintained with 10% FBS. The colonies were stained with crystal violet after 14 days and counted. For invasion analysis, Transwell chamber was used. The membranes were coated with Matrigel (BD Biosciences, USA) and cells were seeded on the upper chamber with no-FBS medium. 10% FBS medium was put on the lower chamber. After cultured for 24 hours, the invaded cells were stained with crystal violet and counted.

**Western blot**

Protein was extracted from cell or tissue and was measured using BCA Kit (Thermo Scientific). Protein was separated by 12% SDS-PAGE and
transferred onto PVDF membrane. The membrane was probed with primary antibodies (Rab23, GAPDH, abcam, USA). Then membrane was incubated with secondary antibodies and was detected with ECL reagent (PerkinElmer Inc.).

**Dual-luciferase reporter assay**

MiR-665 mimic, pRL-CMV Renilla luciferase reporter was cotransfected into the cells using Lipofectamine 2000 (Invitrogen). After 48 hours, luciferase data was measured by using a luciferase assay kit (Promega, USA).

**Statistical analysis**

All data are shown as mean ± SD. Student t test was performed for two group's comparisons and one-way ANOVA was performed for more than two groups’ comparisons. P<0.05 was considered statistically significant.

**Results**

**MiR-665 was downregulated in osteosarcoma tissues**

The expression of miR-665 in osteosarcoma tissues and non-tumorous tissues was shown in the Figure 1A and 1B. Furthermore, miR-665 was downregulated in osteosarcoma tissues compared to non-tumorous tissues (Figure 1C). Moreover, the overall survival (OS) of osteosarcoma patients with low miR-665 expression was lower than that of these patients with high miR-665 expression (Figure 1D).

**MiR-665 was downregulated in osteosarcoma cell lines**

We also demonstrated that miR-665 was downregulated in osteosarcoma cell lines (U2OS, SOSP-9607, Saos-2, and MG-63) compared to osteoblastic cell line (hFOB) (Figure 2A). To study the biological function of miR-665, miR-665 mimics and scramble mimics were transfected into both MG-63 and U2OS cells. The ectopic expression of miR-665 was confirmed using qRT-PCR (Figure 2B).

**MiR-665 overexpression suppressed the osteosarcoma cell proliferation, EMT and invasion**

Ectopic expression of miR-665 suppressed osteosarcoma cell (MG-63). Overexpression of
miR-665 promoted the E-cadherin mRNA expression and inhibited the expression of N-cadherin and vimentin mRNA in the MG-63 cell (Figure 3B). MiR-665 overexpression increased the protein expression of E-cadherin and suppressed the protein expression of N-cadherin and vimentin in the MG-63 cell (Figure 3C). Overexpression of miR-665 suppressed MG-63 cell invasion (Figure 3D).

**MiR-665 downregulates Rab23 expression through targeting its 3’UTR**

Rab23 was found to have a potential binding site of miR-665 within its 3’UTR using bioinformatics (Figure 4A). To confirm whether Rab23 is a direct target gene of miR-665, we performed dual-luciferase reporter containing mutant 3’UTR and wild type of Rab23. Co-transfection with miR-665 suppressed the luciferase activity of the reporter containing the mutant 3’UTR but not the wild type in MG-63 cell (Figure 4B) and U2OS cell (Figure 4C). Overexpression of miR-665 suppressed the expression of Rab23 in both MG-63 and U2OS cell (Figure 4D and 4E).

**Rab23 was upregulated in osteosarcoma tissues and cell lines**

The expression of Rab23 in osteosarcoma tissues and non-tumorous tissues was shown in the Figure 5A and 5B. Furthermore, Rab23 expression was upregulated in the osteosarcoma tissues compared to non-tumorous tissues (Figure 5C). The expression level of miR-665 was inversely associated with the expression level of Rab23 in osteosarcoma tissues (Figure 5D). Rab23 was upregulated in the osteosarcoma cell lines (U2OS, SOSP-9607, Saos-2, and MG-63) compared to osteoblastic cell line (hFOB) (Figure 5E).

**Discussion**

In this study, we demonstrated that miR-665 was downregulated in osteosarcoma tissues compared to non-tumorous tissues. The OS of osteosarcoma patients with low miR-665 expression was lower than that of these patients with high miR-665 expression. Ectopic expression of miR-665 suppressed the osteosarcoma cell proliferation, EMT and invasion. We identified Rab23 as a direct target gene of miR-665. Rab23 was downregulated in the osteosarcoma tissues and cell lines. The expression of miR-665 was inversely associated with the expression of Rab23 in the osteosarcoma tissues. These results suggested that miR-665 acted as a tumor suppressor gene in the development of osteosarcoma.

miRNAs are small, endogenous and noncoding RNAs that regulate gene expression by post-transcriptionally [10, 28-30]. Recently, evidences also have showed that miRNAs acts an important role in the development of osteosarcoma [31-35]. For example, Cheng et al. demonstrated that miR-320 was downregulated in osteosarcoma tissues. Moreover, overexpression of miR-320 suppressed osteosarcoma cell proliferation and tumor growth through targeting fatty acid synthase (FASN) expression [33].
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Huang et al. showed that miR-100 was downregulated in osteosarcoma tissues and enforced expression of miR-100 suppressed the osteosarcoma cell proliferation through repressing Cyr61 expression [36]. Li et al. found that miR-145 was downregulated in osteosarcoma tissues and cells. Overexpression of miR-145 repressed the cell invasion, migration and proliferation by inhibiting Rho-associated protein kinase 1 (ROCK1) expression [37]. Shen et al. demonstrated that miR-128 was upregulated in osteosarcoma tissues and overexpression of miR-128 increased the cell proliferation by suppressing the PTEN expression [38]. However, the role of miR-665 in the development of osteosarcoma still remains unknown. In our study, we demonstrated that miR-665 was downregulated in the osteosarcoma tissues compared to non-tumorous tissues. The OS of osteosarcoma patients with low miR-665 expression was lower than that of these patients with high miR-665 expression. Moreover, ectopic expression of miR-665 inhibited the osteosarcoma cell proliferation, EMT and invasion. These results supported that miR-665 played a suppressor miRNA in the development of osteosarcoma.

Furthermore, Rab23 was identified as a direct target gene of miR-665 in osteosarcoma cells. Rab23 was overexpressed and acted an onco-genic role in many tumors such as squamous cell carcinoma, gastric cancer, bladder cancer, breast cancer and pancreatic duct adenocarcinoma [39-43]. Overexpression of Rab23 promoted tumor cell proliferation, invasion, metastasis, and migration [39, 40]. In our study, we revealed that Rab23 was upregulated in the osteosarcoma tissues compared to non-tumorous tissues. We also found that the expression of Rab23 was increased in osteosarcoma cell lines. Here, we demonstrated an inverse correlation between miR-665 and Rab23 expression in osteosarcoma tissues. These results suggested that increased Rab23 was involved in osteosarcoma progression. Moreover, Rab23 was regulated by some miRNAs such as miR-367 and miR-200b [43-45]. In line with these results, we demonstrated that miR-665 suppressed osteosarcoma cell proliferation and invasion through targeting Rab23.

In conclusion, we revealed that miR-665 was downregulated in osteosarcoma tissues and cell lines and ectopic expression of miR-665 suppressed osteosarcoma cell proliferation and invasion through targeting Rab23. These results suggested that miR-665 might be a potential target for osteosarcoma therapy.
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Disclosure of conflict of interest

None.

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