Research Article

MicroRNA-143 suppresses oral squamous cell carcinoma cell growth, invasion and glucose metabolism through targeting hexokinase 2

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miRNAs are non-coding RNAs that have functions to regulate gene expression and play essential roles in a variety of biological processes of cancers. In the present study, we report miR-143 acts as a tumor suppressor in human oral squamous cell carcinoma (OSCC). The expressions of miR-143 are down-regulated in both OSCC cell lines and patient samples compared with normal adjacent tissues. We found overexpression of miR-143 in oral cancer cell lines suppresses cell migration, cellular glucose metabolism and proliferation. Moreover, overexpression of miR-143 promoted apoptosis and significantly caused cell cycle arrest at G1 stage. The colony formation of oral cancer cells was also suppressed by miR-143. We identified hexokinase 2 (HK2) as a direct target of miR-143. We show that miR-143 complementary pairs to the 3'-UTR of HK2 in oral cancer cells, leading to the inhibition of glycolysis in vitro and in vivo. Moreover, knockdown of HK2 by siRNA in oral cancer cells inhibited glucose metabolism, proliferation and migration. Recovery of glucose metabolism by overexpression of HK2 in miR-143 overexpressing cells restores the cell migration and proliferation, suggesting that the miR-143-mediated cancer suppression is through the direct inhibition of HK2. In summary, the present studies highlight miR-143 as a tumor suppressor in OSCC by the suppression of cell migration, glucose metabolism and proliferation through directly targeting HK2, rendering miR-143 a therapeutic strategy for the treatment of clinical OSCC patients.

Introduction

Oral squamous cell carcinoma (OSCC) is the most common head and neck neoplasm, representing 95% of all forms of head and neck cancer and patients with OSCC display a disappointing 5-year survival rate [1,2]. Recently, the clinical outcome of OSCC has gradually improved, however, patients with OSCC develop chemoresistance reflecting limits in our understanding of the pathogenesis of this cancer [3,4]. Therefore, a better understanding of the molecular mechanisms for oral cancer carcinogenesis contributes to development of novel diagnostic and therapeutic approaches to improve the treatments and prognosis of OSCC patients.

miRNAs are a group of short non-coding RNAs (18–22 nt), which negatively regulate protein translation by binding to complementary sequences in the 3'-UTR of target mRNAs [5]. Cumulative studies reported that miRNAs regulate a wide range of biologic processes of cancers such as proliferation, apoptosis, metastasis, chemosensitivity, tumor development, and stem cell maintenance [6,7]. Moreover, abnormalities of miRNAs have been implicated in the pathogenesis of multiple cancers, suggesting either oncogenic or tumor suppressive roles of miRNAs...
Therefore, identification of aberrantly expressed miRNAs is an important step toward the study of the functions of miRNAs in oral cancer carcinogenesis. In particular, miR-143 has been extensively studied as a tumor suppressor in several cancers such as breast cancer [9], colon cancer [10], lung cancer [11], pancreatic cancer [12], brain tumor [13], and melanoma [14]. However, the roles of miR-143 and its target gene in regulating human OSCC development are poorly understood. In the present study, the roles of miR-143 in the human OSCC will be investigated. The potential target of miR-143 and the regulator mechanisms of miR-143 in oral cancer proliferation, invasion, and metabolism will be assessed. Our study will contribute to the development of the miRNAs-based therapeutic agents for the clinical treatments of oral cancer patients.

Materials and methods

Cell culture and tissue specimens

The collection of tumor specimens from OSCC patients was approved by the Institutional Review Board (IRB) of Tianjin Third Central Hospital. Human oral cancer cell lines (OECM-1 and Tca8113) were obtained from the cell bank of type culture collection of Chinese Academy of Sciences (Shanghai, China). Cells were routinely cultured in Dulbecco’s modified Eagle’s medium (DMEM; Gibco BRL, Paisley, U.K.) containing 10% FBS (HyClone, Logan, UT, U.S.A.), and 100 units/ml penicillin, 100 mg/ml streptomycin (HyClone, Logan, UT, U.S.A.) at 37°C in a humid atmosphere with 5% CO₂.

miRNAs and plasmid DNA transfection

miR-143 mimic and control mimic were obtained from ThermoFisher Scientific (Waltham, MA, U.S.A.). The miRNAs and plasmid DNA for overexpressing HK2 (hexokinase 2) were transfected using Lipofectamine® 2000 (Invitrogen Life Technologies). After 48 h following transfection, the expression of miR-143 was detected by quantitative-reverse transcription polymerase chain reaction (qRT-PCR), and the expression of hexokinase was measured by Western blotting.

Luciferase assays

The 3′-UTR luciferase vector was constructed using the pMIR-report luciferase vector containing wild-type or mutant 3′-UTR of HK2 mRNA, which carries a putative miR-143 complementary site. OECM-1 and Tca8113 cells (3 × 10⁴ per well) were pre-seeded in a 24-well plate the day before transfection for overnight. Cells were transfected with 0.5 μg of the 3′-UTR luciferase vector and 50 nM miR-143 mimics or negative control using Lipofectamine RNAiMAX (Invitrogen). Assays were performed using the pMIR-report luciferase vector system after 48 h of co-transfection.

Cell proliferation assay

The cancer cells were transfected with miR-143 mimic, or control mimic for 48 h. Cells were then seeded in a 96-well plate, at a density of 3000 cells/well for overnight incubation. The cell proliferation rates were measured with MTT assay (Sigma–Aldrich, Inc., St. Louis, MO, U.S.A.). Briefly, cells were treated with MTT at 50 mg per well. The generated formazan was dissolved in DMSO, and the absorbance was recorded by measuring the absorbance at 590 nm with a plate reader. The same experiment was repeated three times.

Scratch and cell migration assays

For wound-healing assays, 1 × 10⁵ cells were seeded on glass coverslips and cultured until confluence. Cells were scratched with micropipette tips, and images were captured at 0 and 24 h after wounding. The transwell assay was done by using a transwell chamber consisting of 8 mm membrane filter inserts (Corning, Corning, NY, U.S.A.) according to the previous description [15]. For each experiment, the number of cells in three random fields on the underside of the filter was counted, and three independent filters were analyzed.

Cell cycle analysis

The cell cycle was analyzed using the Cell Cycle Assay Kit (Abcam, #ab112116) according to the manufacturer’s protocol.
Colony formation assay
One hundred oral cancer cells were placed into each 6 cm cell culture dish and cultured for 2 weeks for allowing the colony formation. Cells were fixed by methanol and stained by 0.05% Crystal Violet for 5 min. After washing by PBS, the dishes were recored.

Measurements of glucose metabolism
Glucose uptake was detected using a Glucose Uptake Colorimetric Assay Kit (#K676-100, BioVision, Inc., Milpitas, CA, U.S.A.) and lactate production was detected using a Lactate Colorimetric Assay Kit II (#K627-100, BioVision, Inc., Milpitas, CA, U.S.A.) according to the manufacturer’s instructions. Results were normalized by the protein quantities of experimental group to the control group. All experiments were repeated three times.

LDH activity assay
The LDH activity assay was performed using the Lactate Dehydrogenase Activity Assay Kit (Sigma, # MAK066, St. Louis, MO, U.S.A.) according to the manufacturer’s protocol.

Xenograft experiments
Athymic nude mice were purchased from Vital River (Beijing, China) and housed in the certified animal facility in the Tianjin Third Central Hospital. All experimental procedures were performed according to the Guide for the Care and Use of Laboratory Animals published by the U.S. National Institutes of Health (NIH Publication No. 85-23, revised 1996). For developing human oral cancer xenografts, OEOM-1 cells (3 × 10⁶) in were implanted by subcutaneous (s.c.) injection into each mouse. After tumor development, the mice were randomly divided into two groups: control miRNA and miR-143 mimic injection. The injection was given twice a week for 2 weeks. Tumors were taken after injections for the downstream experiments.

Tissue immunohistochemistry
Human oral tumor IHC was performed according to previous descriptions [16]. Briefly, sections of miR-143 low or high expression were deparaffinized in xylene. Then, samples were rehydrated through a gradient concentration of alcohol followed by incubation by 5% normal goat serum to block non-specific staining. Sections were then incubated with rabbit anti-HK2 antibody (1:200, Cellsignaling, #2867, Danvers, MA, U.S.A.) overnight at 4°C. The slides were incubated with biotin-labeled goat anti-rabbit IgG and further incubated with streptavidin peroxidase solution (SABC kit, Boster Biological Technology, Ltd., Wuhan, China). The staining was visualized by reaction with 3,3’-di-aminobenzidine (Boster Biological Technology, Ltd.) in PBS for 5 min at room temperature. All of the IHC staining results were reviewed independently by two pathologists.

qRT-PCR
Total RNA was extracted from cells or human OSCC tumor tissues using TRIzol reagent (Invitrogen) according to the manufacturer’s protocol. Total RNA was used for reverse transcription using the PrimerScript RT-PCR kit (TaKaRa Biotechnology, Dalian, China). miRNAs were reverse transcribed using sequence-specific stem-loop primers (Invitrogen). Quantitative RT-PCR was conducted according to the previous description [15]. Briefly, qRT-PCR was conducted using a standard SYBR Green PCR kit (Roche) protocol with a CFX96 Touch™ Real-Time PCR Detection System (Biorad). The relative expression was calculated using the 2−ΔΔCt method. The transcription levels of GAPDH or U6 were used as an internal control. The specific Primers are listed as follows: GAPDH: forward: 5′-GCACCGTCAAGGCTGAGAAC-3′; reverse: 5′-TGGTGAAGACGCCAGTGGA-3′; HK2: forward: 5′-CAAAGTGACAGTGGGTTGGG-3′; reverse: 5′-GCCAGGTCTCTCTACGTCTC-3′.

Western blotting
Cells were collected and lysed using RIPA buffer (Pierce, Waltham, MA, U.S.A.) followed by protein concentration measurements by Bradford assay. Equal amount of protein samples were separated by 10% SDS/PAGE, and were electrophoretically transferred on to a PVDF membrane (Millipore Corporation). The membrane was blocked with 5% low-fat milk in Tris buffered saline for 1 h at room temperature, and then incubated with primary antibodies (β-actin and HK2: cell signaling) at 1:1000 dilution at 4°C overnight. The primary antibodies were detected with horseradish peroxidase-conjugated secondary antibodies and a chemiluminescence detection kit (Pierce, Waltham, MA, U.S.A.).
Figure 1. miR-143 is downregulated in patient OSCC species and suppress oral cancer cells growth

(A) Expression level of miR-143 in each individual case of OSCCs and adjacent normal tissues was measured by qRT-PCR. snRNA U6 was used as an internal control. (B) OECM-1 and Tca8113 cells were transfected with miR-143 mimic or control miRNAs for 48 h, the expressions of miR-143 were measured by qRT-PCR. (C) OECM-1 and Tca8113 cells were transfected with miR-143 mimic or control miRNAs for 48 h, cells were plated into 24-well plate and the cell proliferation rates were measured at 0, 24, 48, 72, and 96 h by MTT assay. (D) The transfected OECM-1 and Tca8113 cells were subjected to cell cycle analysis, (E) colony formation assay and (F) apoptosis assay. Data are presented as mean ± S.D.: *, *P* < 0.05; **, **P** < 0.01.

Statistical analysis
The statistical significance was assessed by Student’s *t* test using GraphPad Prism program 5.0 (GraphPad Software, U.S.A.). *P* < 0.05 was considered statistically significant.

Results

miR-143 levels are significantly down-regulated in human OSCC tumors

Previous studies revealed that miR-143 possesses tumor-suppression function in multiple tumor types [9-14]. To evaluate the function of miR-143 in human oral tumor tissues, we analyzed the expressions of miR-143 from 15 human normal health oral tissues and 15 human oral cancer tissues using real-time PCR. As we expected, the expression levels of miR-143 in OSCC tumor species were significantly lower than that in normal oral tissues (Figure 1A), suggesting miR-143 act as a tumor suppressor in OSCC.

miR-143 suppresses OSCC cells proliferation, cell cycle and colony formation, promotes apoptosis

To further validate the tumor-suppression functions of miR-143 in OSCC, the effects of miR-143 on cell invasion, glucose metabolism, and tumor proliferation were measured. We transiently transfected miR-143 mimic into two human OSCC cell lines, OECM-1 and Tca8113 (Figure 1B). Overexpression of miR-143 markedly suppressed the proliferation ratios of these two oral cancer cell lines (Figure 1C) compared with negative miRNA transfection. Moreover, we found overexpression of miR-143 caused a cell cycle arrest of oral cancer cells in G1-phase (Figure 1D). The colony formation assay demonstrated that overexpression of miR-143 significantly suppressed colony of OCEM-1 and Tca8113 (Figure 1E). The apoptosis was induced by miR-143 (Figure 1F).
**miR-143** suppresses OSCC cells migration, invasion and glucose metabolism

In addition, the invasive abilities of oral cancer cells were significantly inhibited by *miR-143* but promoted by *miR-143* inhibitor (Figure 2A–C). It has been reported that dysregulated glycolysis reflected one of the features of cancer cells [17], we measured the glucose uptake and lactate production both of which are indicators of glucose metabolism. Consistently, the glucose metabolism was attenuated by overexpression of *miR-143* but activated by *miR-143* inhibition (Figure 2D), supporting the above results that *miR-143* may play a suppressive role in oral cancer.

**HK2 is a direct target of miR-143 in OSCC cells**

To investigate the mechanisms by which *miR-143* suppresses OSCC cells proliferation, invasion and glucose metabolism, we searched the potential targets of *miR-143* by the web-based target analysis tools www.microRNA.org and www.targetscan.org. The analysis software predicted that HK2 might be a target for *miR-143* and the 3’-UTR of HK2 contains a highly conserved binding site for *miR-143* (Figure 3A). In addition, the binding of *miR-143* on HK2 3’-UTR is conserved in multiple species (Figure 3B), further supporting our results that *miR-143* directly targets HK2 and this binding has important functions. Currently, there is no publication reported that HK2 was a *miR-143* direct target in OSCC cells. To determine whether *miR-143* could target HK2 in endothelial cells, we transfected control miRNA, *miR-143* mimic or *miR-143* inhibitor into OECM-1 and Tca8113 cells. Our results showed overexpression of *miR-143* inhibited HK2 protein and mRNA levels and inhibition of *miR-143* could increase HK2 protein and mRNA levels (Figure 3C–E). To further validate targeting of HK2 by *miR-143*, we investigated whether *miR-143* directly interacted with the 3’-UTR of HK2 using a dual-luciferase reporter assay. A reporter plasmid harboring a mutated *miR-143* binding site was used as a control. Overexpression of *miR-143* significantly suppressed luciferase activity of 3’-UTR of the wild-type HK2 reporter constructs in both OECM-1 and Tca8113 cells (Figure 3F), while the suppressive effect of *miR-143* mimic was abrogated with mutant HK2 3’-UTR. These results demonstrated that HK2 was indeed a direct downstream target of *miR-143* in human OSCC cells.
Knockdown of HK2 suppresses OSCC cells growth, invasion, and glucose metabolism

To investigate whether the miR-143 mediated OSCC cell suppression through HK2, we specifically knocked down HK2 by siRNA (Figure 4A). Consistently, knockdown of HK2 in Tca8113 cells significantly suppressed cells proliferation (Figure 4B), invasion (Figure 4C) and glucose metabolism (Figure 4D) compared with control siRNA transfection, indicating deregulation of HK2 by miR-143 might be a therapeutic approach for OSCC tumor treatments.

Restoration of HK2 rescues the miR-143-mediated inhibitory effects on OSCC cells

To validate whether the tumor-suppressive function of miR-143 was through direct inhibition of HK2 in OSCC cells, an HK2 ectopic expression vector was transfected into Tca8113 cells which were pre-transfected with pre-miR-143. The restoration of HK2 in miR-143 overexpressing cells was measured by Western blot (Figure 5A). As we expected, rescue of HK2 in miR-143 overexpressing cells led to marked enhancement of the cellular proliferation (Figure 5B), invasion (Figure 5C) and tumor glucose metabolism (Figure 5D). Taken together, the above data further supported that miR-143 inhibited OSCC cellular proliferation, invasion and glucose metabolism through direct targeting HK2.

miR-143 suppresses glucose metabolism and HK2 expression in a xenograft mouse tumor model

To further support the above in vitro results that miR-143 suppresses oral cancer cells glucose metabolism through targeting HK2, we performed in vivo xenograft experiments. OECM-1 cells were inoculated into the mammary fat pads of nude mice to develop tumor. After tumors formed, miR-143 mimic or control miRNA was injected into the tumors twice a week for 2 weeks. As shown in Figure 6A, injection of miR-143 mimic significantly down-regulated HK2 expression in nude mice. Moreover, the mRNA levels of LDHA (Figure 6B) and activity of LDH (Figure 6C) were
Figure 4. Knockdown of HK2 inhibits proliferation, invasion, and glucose metabolisms of oral cancer cells
(A) Tca8113 cells were transfected with control siRNA or siHK2 for 48 h, cells were collected and subjected to Western blot analysis. β-Actin was a loading control. (B) Tca8113 cells were transfected with control siRNA or siHK2 for 48 h, then cell proliferation assay was performed at 0, 24, 48, 72, and 96 h. (C) The transwell assay was performed for the measurements of cell invasion ability in control or miR-143 mimic transfected Tca8113 cells. (D) The glucose uptake and lactate production were measured in control or miR-143 mimic transfected Tca8113 cells. Data are presented as mean ± S.D.: *, P < 0.05; **, P < 0.01.

Figure 5. Restoration of HK2 rescues oral cancer cells proliferation, invasion, and glucose metabolism
(A) Tca8113 cells were transfected with control, miR-143 mimic or miR-143 mimic plus HK2 for 48 h followed by Western blot analysis. β-Actin was a loading control. (B) Tca8113 cells were transfected with miR-143 mimic or miR-143 mimic plus HK2 for 48 h, then cell proliferation assay was performed at 0, 24, 48, 72, and 96 h. (C) The transwell assay was performed for the measurements of cell invasion ability in control, miR-143 mimic or miR-143 mimic plus HK2 transfected Tca8113 cells. (D) The glucose uptake and lactate production were measured in miR-143 mimic or miR-143 mimic plus HK2 transfected Tca8113 cells. Data are presented as mean ± S.D.: *, P < 0.05; **, P < 0.01.
The correlation of HK2 mRNAs and miR-143 expressions of LDHA and (C) LDH activity. (D) The representative expressions of HK2 in miR-143 high and low OSCC patient species. (E) The correlation of HK2 mRNAs and miR-143 levels in OSCC tissues was shown. P<0.05 is considered statistical significance.

consistently decreased in the miR-143-injected mice tumors. Taken together, these in vivo results further support that miR-143 is a tumor suppressor through suppression of glucose metabolism.

Invert correlation between miR-143 and HK2 expressions in human OSCC tissues

In light of our observation that overexpression of miR-143 led to the down-regulation of HK2 in OSCC cells, we postulated an inverse correlation between miR-143 expression and HK2 in metastasis oral tumor tissues. As we expected, we detected a strong negative correlation between miR-143 and HK2 in oral tumor tissues: the expression of HK2, which is associated with tumor aggressiveness [17] was up-regulated in high metastasis oral tumor species (Figure 6D) compared with low metastasis tumors, indicating that HK2 may be associated with metastasis in OSCC; in contrast, the expressions of miR-143 were significantly down-regulated in high metastasis tumors (Figure 6E), indicating miR-143 might be a therapeutic target for the treatments of metastasis OSCC tumor by targeting HK2.

Discussion

Abnormal expressions of miRNAs have been implicated in the pathological processes of a variety of human cancers [6-8]. Overexpression of oncogenic miRNAs or down-regulation of tumor suppressor miRNAs plays an essential role in tumorigenesis. miR-143 has been reported as an under-expressed miRNA in various tumors [9-14,17-20]. In the current study, we identified a tumor suppressive role of miR-143 in human OSCC. We found miR-143 was significantly down-regulated in oral cancer patient specimens and oral cancer cell lines. In addition, overexpression of miR-143 suppressed oral cancer cell proliferation, migration, and invasion. Our results are consistent with the previously described functions of miR-143 in human cancers.

HK2 converts glucose to glucose-6-phosphate, which is the first committed step in anaerobic glycolysis [21]. It has been widely studied that HK2 regulates tumorigenesis and migration in multiple cancer types, yet the mechanisms are still poorly defined [22]. HK2 has been reported to be a transcriptional target of HIF-1, which is induced in response to hypoxic conditions [23]. We identified HK2 as a direct target of miR-143 in oral cancer cells and tumor patients, suggesting inhibition of HK2 by miR-143 might be a new therapeutic approach for the treatments of oral cancer.

Cancer cells predominantly generate ATP as well as metabolic intermediates by a high rate of glycolysis in the cytosol, rather than by a relatively high efficient approach through oxidation of pyruvate in mitochondria [24]. In the present study, we reported the glycolysis rate was suppressed by miR-143, supporting the tumor suppressive roles of
miR-143 in oral cancer cells. In addition, we demonstrated restoration of HK2 in miR-143 overexpressing oral cancer cells could increase the tumorigenesis and invasion of oral cancer cells in vitro. However, the mechanisms for the HK2-mediated tumor migration and invasion are still under investigation and more functional targets of miR-143 in OSCC require further discovery. In summary, we report a tumor suppressive role of miR-143 in human oral cancer cells and oral tumor patients. Overexpression of miR-143 inhibited cancer cells proliferation, migration, invasion, and glucose metabolism through direct targeting HK2. Restored HK2 expression in miR-143 overexpressing cells exhibited oncogenic effects in vitro. Our data suggest an important role of the miR-143 mediated glycolysis in the development of gene therapy for OSCC.

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Author contribution

X.H.S. designed research; X.H.S. and L.Z. performed research; X.H.S. and L.Z. analyzed data and wrote the paper.

Competing interests

The authors declare that there are no competing interests associated with the manuscript.

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Abbreviations

OSCC, oral squamous cell carcinoma; HK2, hexokinase 2; LDH, Lactate dehydrogenase; HIF-1α, Hypoxia inducible factor-1α.

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