

Radiation-resistance increased by overexpression of microRNA-21 and inhibition of its target PTEN in esophageal squamous cell carcinoma

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Abstract

Objective: Overexpression of microRNA-21 (miR-21) increases the radiation resistance of esophageal squamous cell carcinoma (ESCC). However, the molecular mechanism responsible for this action is still unclear. In the present study, we investigated the role of phosphatase and tensin homolog deleted on chromosome 10 (PTEN) in miR-21-enhanced radiation resistance in patients with ESCC.

Methods: We evaluated the association between miR-21 levels and radiation resistance in patients with ESCC. We also investigated the role of PTEN in the proliferation and apoptosis of ESCC cells transfected with miR-21 inhibitor during irradiation, using PTEN small interfering RNA (siRNA).

Results: MiR-21 levels were significantly higher in radiation-resistant patients. Downregulation of miR-21 during irradiation suppressed the radiation resistance of ESCC cells, demonstrated by decreased cell proliferation and increased cell apoptosis. PTEN siRNA attenuated miR-21-induced suppression of radiation resistance in ESCC cells.

Conclusions: These results suggest that miR-21 enhanced the radiation resistance of ESCC by inhibiting PTEN. MiR-21 and PTEN are potential therapeutic targets for ESCC.

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Keywords

miR-21, PTEN, radiation resistance, esophageal squamous cell carcinoma, cell proliferation, apoptosis

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Introduction

Esophageal cancer is a relatively common high-risk tumor worldwide and a major cause of cancer-related death. About half of all cases of esophageal cancer occur in China.¹ Esophageal cancer can be divided pathologically into squamous cell carcinoma and adenocarcinoma, with squamous cell carcinoma mainly prevalent in developing countries and adenocarcinoma more common in developed countries, while other types are rare. Most cases in China are esophageal squamous cell carcinoma (ESCC), which accounts for around 90% of all cases in this country.² The survival rate of patients with ESCC is less than 10%,³ and the 5-year overall survival rate is 20% to 40%.⁴ Surgery, radiotherapy, and chemotherapy are the three main routine treatments for esophageal cancer. However, owing to the physiological anatomy of the esophagus, the atypical early symptoms, and a lack of effective early-diagnosis methods, most patients with esophageal cancer are diagnosed at an advanced stage and are therefore unsuitable for radical surgery.⁵ Radiotherapy is thus the main treatment method for advanced esophageal cancer, and more than half of all patients receive radiation during their treatment. However, surviving tumor cells may develop radiotolerance, resulting in tumor recurrence after the radiosensitive cells are killed by radiation therapy.⁶

Activation of oncogenes and inactivation of tumor suppressor genes leading to abnormalities in intercellular signaling pathways may be one mechanism leading

to radiation resistance in tumor cells.⁷ Numerous recent studies have shown that micro RNAs (miRNAs) are involved in many cellular regulatory processes, including activation of different signaling pathways and induction of apoptosis.^{8–10} Furthermore, miRNA expression is involved in the regulation of radiation resistance in tumor cells.^{11,12} MiRNAs are a group of small, conservative, non-coding RNAs that degrade or repress their translation target mRNAs by direct binding to the 3'-untranslated region.¹³ MiR-21 is overexpressed in many solid tumors.¹⁴ Meng et al. reported that phosphate and tension homology deleted on chromosome 10 (PTEN) was a target gene of miR-21,¹⁵ and miR-21 inhibited PTEN and promoted the growth and invasion of non-small cell lung cancer and pancreatic cancer cells.^{16,17} Liu et al. reported that miR-21 promoted the growth, metastasis, and chemoresistance or radioresistance of non-small cell lung cancer cells by targeting PTEN,¹⁸ and subsequent studies confirmed that miR-21 could regulate the radioresistance of tumor cells through PTEN.^{19,20} Li et al.²¹ also reported that abnormal regulation of miR-21 promoted the progression of esophageal cancer.

Based on the above studies, we hypothesized that miR-21 might enhance the radioresistance of tumor cells by inhibiting the expression of PTEN in ESCC. We therefore investigated the involvement of miR-21 in mediating radioresistance in ESCC. We also examined the role of PTEN as a target gene of miR-21 in the regulation of radiosensitivity, and considered the potential of

miR-21 as a therapeutic target for radiosensitization in patients ESCC.

Materials and methods

Patients and specimens

Tumor specimens were obtained from 40 consecutive patients with ESCC admitted to Hebei Medical University between September 2015 and December 2017. All patients received radiotherapy and were divided into a radiation-resistant group (20 patients with <40% reduction of tumor size after radiation treatment) and a radiation-sensitive group (20 patients with >60% reduction of tumor size after radiation treatment). The tissue specimens were stored at -193°C until use. The study was approved by the Ethics Research Committee of Fourth Hospital, Hebei Medical University (HU406AF). Signed informed consent, in accordance with the Declaration of Helsinki, was obtained for each donated tumor sample.

Cell culture and ionizing radiation

The human ESCC TE-1 cell line was purchased from the Chinese Academy of Sciences (Shanghai, China). The cells were grown in RPMI-1640 medium with 10% fetal bovine serum (HyClone, Logan, UT, USA), 100 U/mL penicillin, and 100 $\mu\text{g}/\text{mL}$ streptomycin at 37°C in 5% CO_2 . Cells in logarithmic growth phase were used for all experiments. Cells were exposed to 0, 2, 4, 6, and 8 Gy irradiation, respectively, using a Model 143 $^{137}\text{cesium}$ γ -irradiator (JL Shepherd & Associates, San Fernando, CA, USA) at a rate of 2.4 Gy/minute.

RNA isolation and quantitative real-time PCR (qRT-PCR)

Total RNA was isolated using an Absolutely RNATM RT-PCR Miniprep

Kit (Stratagene, China), in accordance with the manufacturer's instructions. Reverse transcription was carried out using 10 ng of RNA in a volume of 5 μL , with 3 μL of primer. RT-PCR was performed using the TaqMan probe method (PrimeScript RT reagent Kit, TaKaRa Biotechnology, Dalian, China), in accordance with the manufacturer's instructions. PCR was performed with an ABI 7500 real-time PCR system (Applied Biosystems, Foster City, CA, USA). The primers used in this study were as follows: miR-21-5p: UAGCUUAUCAGACUGAUGUUGA, miR-21-5p R: CTCAACTGGTGTCGTG GAGTC GGCAATTCAGTTGAGTCAA CATC, miR-21-5p F: ACACTCCAGCT GGGTAGCTTATCAG ACTGATG, all R: CTCAACTGGTGTCGTGGA, U6F: CTCGCTTCGGCAGCACACA, U6R: AAC GCTTCACGAATTTGCGT. U6 was used for standardization and relative expression levels were calculated using the $2^{-\Delta\Delta\text{Ct}}$ method.

Cell transfection

We constructed an miR-21 inhibitor vector and PTEN small interfering RNA (siRNA). A negative control (NC) group was treated with a scrambled sequence with no significant homology to any rat, mouse, or human gene. The cells were transfected using LipofectamineTM 2000 (Invitrogen, Carlsbad, CA, USA), in accordance with the manufacturer's instructions. MiR-21 expression was assessed by qRT-PCR at 48 hours after transfection.

Cell proliferation assay

We evaluated the effect of miR-21 expression on cell proliferation 48 hours after transfection using a Cell Counting Kit-8 (CCK-8, Beyotime, China). The radioreistance of the transfected cells was assessed by plating in 96-well plates and exposure to

8 Gy γ -radiation. Cells were removed at 1, 2, 3, and 4 days, CCK-8 (10 μ L) was added, and the cells were then incubated at 37°C for 1.5 hours. Finally, cell viability was measured at 450 nm using a microplate reader.

Cell apoptosis

Apoptosis was detected by annexin V-fluorescein isothiocyanate (FITC) staining. The cells were irradiated for 8 hours at 8 Gy and then collected. The supernatant was removed and the cell monolayer was gently washed with phosphate-buffered saline (PBS). The cells were then digested with trypsin, digestion was terminated with PBS containing 10% serum, and the cells were resuspended in ice-cold PBS, fixed with 70% ethanol, and stained using an Annexin V-FITC Apoptosis Detection Kit (BD Biosciences, San Jose, CA, USA), in accordance with the manufacturer's instructions. Fluorescence was measured using a flow cytometer (BD Biosciences) and the rate of apoptosis was analyzed.

Western blot

Protein expression was detected by western blot. Total protein was extracted by conventional radioimmunoprecipitation assay buffer (Beyotime, Shanghai, China), in accordance with the manufacturer's instructions. The extracted protein was then quantified by the bicinchoninic acid assay (Cell Signaling Technology, Danvers, MA, USA), in accordance with the manufacturer's instructions. Proteins were separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis, electroporated to polyvinylidene difluoride membranes, blocked in 5% skim milk for 1.5 hours, and then incubated with primary antibody on a shaker at 4°C overnight. The primary antibodies used were anti-PTEN (1:200, Abcam, Cambridge, MA, USA)

and anti-cleaved-poly (ADP-ribose) polymerase (PARP), and anti-cleaved-caspase 3 (1:1000; Cell Signaling Technology). The blots were then incubated with horseradish peroxidase-conjugated secondary PARP antibody (1:100; Cell Signaling Technology) for 1 hour. Glyceraldehyde 3-phosphate dehydrogenase antibody was used as a control. Finally, the blots were visualized by enhanced chemiluminescence with horseradish peroxidase-conjugated donkey anti-rabbit secondary antibody (Cell Signaling Technology) for 1 hour at room temperature.

Statistical analyses

The data were presented as mean \pm standard deviation. All statistical analyses were performed using SPSS for Windows, Version 15.0 (SPSS Inc., Chicago, IL, USA) in at least three independent experiments. Two-tailed Student's *t*-tests or one-way ANOVA were used to analyze the differences among the groups. $P < 0.05$ was considered significant.

Results

MiR-21 levels were increased in radiation-resistant compared with radiation-sensitive patients

MiR-21 was overexpressed in ESCC, and downregulation of miR-21 suppressed the radiation resistance of ESCC.²² We detected miR-21 expression levels in 20 radiation-resistant and 20 radiation-sensitive ESCC patients using qRT-PCR. MiR-21 levels were significantly higher in the radiation-resistant patients ($P < 0.05$) (Figure 1), confirming that high miR-21 levels were significantly correlated with radiation resistance in ESCC.

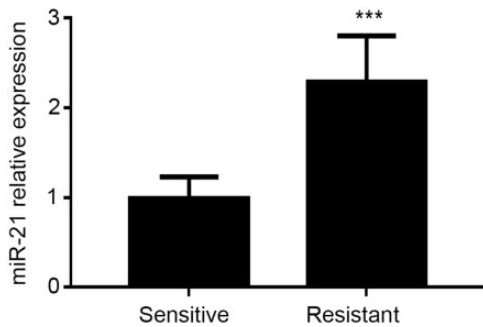


Figure 1. Expression of miR-21 in radiation-resistant and radiation-sensitive esophageal squamous cell carcinoma, detected by qRT-PCR. *** $P < 0.001$.

MiR-21 was associated with radiation resistance in ESCC cells

MiR-21 expression was examined in TE-1 cells after radiation with 0, 2, 4, 6, and 8 Gy. MiR-21 expression was increased by radiation in a dose-dependent manner in these cells (Figure 2a), indicating that miR-21 was associated with radiation in ESCC cells. The viability of the TE-1 cells was not significantly inhibited after 24 hours of continuous irradiation at 8 Gy, compared with cells without continuous irradiation (Figure 2b). These findings imply that miR-21 was associated with radiation resistance in ESCC cells. After cumulative radiation exposure, the cells developed a distinct radioresistant phenotype, with increased expression levels of the tumor radioresistance factors heat shock protein90 and HER3 (data not shown), indicating the establishment of a radioresistant cell line.

Downregulation of miR-21 suppressed proliferation of irradiated ESCC cells via PTEN pathway

To determine the role of miR-21 in ESCC during irradiation, we transfected ESCC cells with miR-21 inhibitor or NC vector

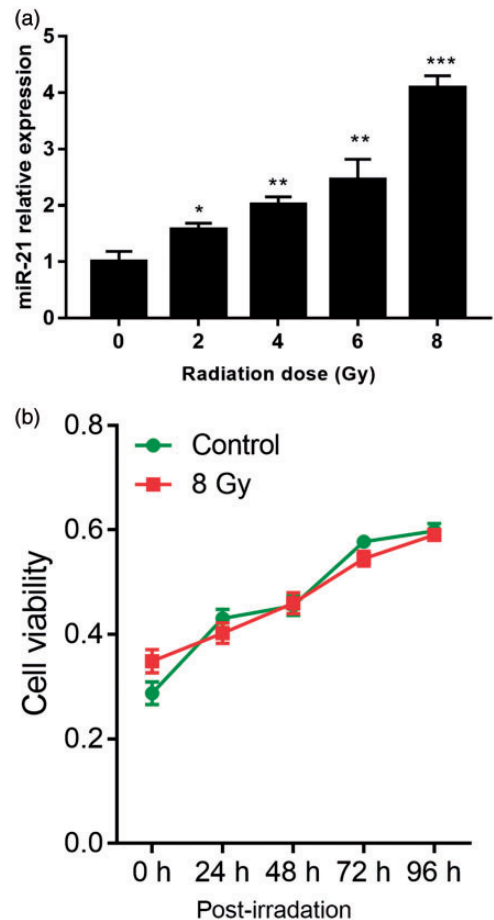


Figure 2. Effects of different radiation doses on miR-21 levels and viability in TE-1 cells. Cells were exposed to different doses (0, 2, 4, 6, 8 Gy) of radiation at a rate of 2.4 Gy/minute, cultured for 48 hours, and then collected. (a) MiR-21 levels were detected by qRT-PCR. (b) After 48 hours irradiation at 8 Gy, irradiation was removed or not. Viability of TE-1 cells was detected by CCK8 assay. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ vs. 0 Gy.

for 48 hours, and confirmed the downregulation of miR-21 by qRT-PCR (Figure 3a).

After transfection for 48 hours, cells were exposed to 8 Gy irradiation at a rate of 2.4 Gy/minute for 48 hours, followed by assessment of cell proliferation (Figure 3b). Downregulation of miR-21 significantly inhibited cell viability under irradiation

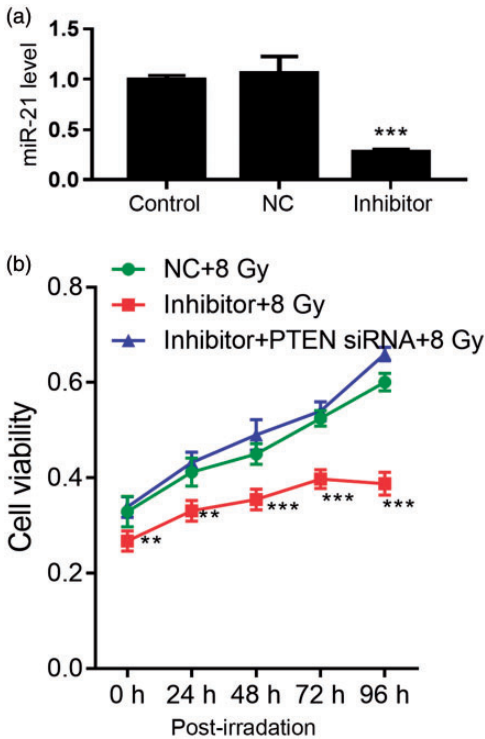


Figure 3. Viability of miR-21 inhibitor- or PTEN siRNA-transfected TE-1 cells after exposure to 8 Gy for 48 hours. After transfection, cells were irradiated with 8 Gy for 48 hours, followed by removal of irradiation or not. Cell viability was then detected by CCK8 assay. (a) Expression of miR-21 was detected by qRT-PCR after transfection for 48 hours. *** $P < 0.001$ vs. control. (b) Cell viability. ** $P < 0.01$, *** $P < 0.001$ vs. control+8 Gy. NC, negative control.

compared with NC transfection ($P < 0.05$). The effect of miR-21 inhibitor transfection on cell viability was reversed by PTEN siRNA (Figure 3b). These results indicated that miR-21 mediated radiation resistance in ESCC cells via the PTEN pathway.

MiR-21 inhibitor increased expression of cell apoptosis markers via PTEN pathway

The effects of miR-21 inhibitor and PTEN on cell apoptosis were confirmed by detecting the expression levels of caspase 3 and

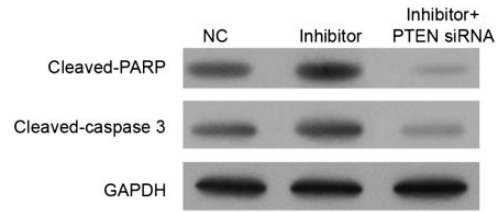


Figure 4. Expression levels of cleaved-caspase 3 and cleaved-PARP in miR-21 inhibitor- and/or PTEN siRNA-transfected TE-1 cells after exposure to 8 Gy for 48 hours. After transfection, cells were irradiated with 8 Gy for 48 hours and then collected, and PTEN, cleaved-caspase 3, and cleaved-PARP levels were detected by western blot. NC, negative control.

PARP (Figure 4). After 8 Gy irradiation for 48 hours, miR-21 inhibitor significantly induced PARP and caspase 3 activities ($P < 0.05$), and this effect was reversed by PTEN siRNA. miR-21 inhibition thus increased PARP activity and caspase 3 activity via the PTEN pathway.

Discussion

In the current study, we used an miR-21 inhibition vector and PTEN siRNA to examine the roles of miR-21 and PTEN in the radioresistance of ESCC. We collected pathological specimens from patients with ESCC and divided the patients into radiation-sensitive and radiation-resistant groups based on their response to radiation. We then tested the hypothesis that miRNA-21 promoted radiation resistance of tumor cells by inhibiting the expression of PTEN, both *in vivo* and *in vitro*.

Radiation therapy is widely used as a routine clinical treatment for cancer, especially for some tumors with a specific physiological anatomy.^{23,24} However, inter-individual differences result in a relatively high frequency of radiation-induced radioresistance,²⁵ though the mechanism responsible is not yet clear. Recent miRNA research has revealed that radiation affects

the expression profiles of miRNAs, and evidence suggests that radiation resistance may be caused by the abnormal expression of single miRNAs.^{26,27} miRNA-21 was widely overexpressed in solid tumors, associated with a poor tumor prognosis.²⁸ In the present study, we detected miR-21 expression levels in radiosensitive and radioresistant patients with ESCC, and showed that levels were significantly higher in the radiation-resistant group. This result was consistent with previous studies and further corroborates the role of miR-21 in ESCC.²¹ Overexpression of miR-21 was correlated with the generation of radioresistance, and the current *in vitro* cell experiments showed that miRNA-21 expression levels increased with increasing radiation dose. Overall, the above experiments suggest that miRNA-21 is related to the generation of radiation resistance in ESCC, and that high expression levels of miRNA-21 can promote radioresistance in ESCC.

Li et al.²¹ reported that inhibition of miR-21 increased the radiosensitivity of esophageal cancer cells by downregulating the expression of PTEN.²¹ PTEN is a tumor suppressor gene located on chromosome 10q23, which is mutated or deleted in a variety of tumors.^{29,30} The most important substrate for PTEN is phosphatidylinositol (3,4,5)-trisphosphate (PIP3), which is the product of phosphatidylinositol-3 kinase (PI3K) and mediates the activation of Akt (also known as protein kinase B). PTEN dephosphorylates PIP3 to maintain a low level of PIP3, thereby downregulating the PI3K/Akt pathway as part of the PTEN-PI3K/Akt signal transduction pathway.³¹ Furthermore, PTEN targeting by miR-21 has been assessed by luciferase assay.^{15,32,33} In this study, we transfected ESCC cells with miR-21 inhibitor or NC vector and found that downregulation of miR-21 significantly inhibited cell viability, but this effect was reversed by PTEN siRNA. We therefore speculate that

miRNA-21 induces radioresistance in ESCC by inhibiting PTEN. We further confirmed our hypothesis by examining the expression of apoptotic proteins in the PTEN pathway after irradiation of TE-1 cells with miR-21 inhibitors, and found that miR-21 inhibitors significantly induced PARP activity and caspase 3 activity, and that this inhibition was also reversed by the addition of PTEN siRNA. Overall, the above experimental studies suggest that miR-21 acts via the PTEN pathway in ESCC.

Conclusion

This study demonstrated that miR-21 is abnormally overexpressed in patients with radioresistant ESCC. Furthermore, experimental studies revealed that this radioresistance was achieved by miR-21 regulation of its target gene PTEN. These results thus confirm that miR-21 and its target gene PTEN may be potential molecular targets for the future treatment of patients with ESCC.

Availability of data and materials

The raw data used and analyzed for the current study (not including any personal statements or information) are available from the corresponding author on reasonable request.

Declaration of conflicting interest

The authors declare that there is no conflict of interest.

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