

## miR-183 Modulates Cell Apoptosis and Proliferation in Tongue Squamous Cell Carcinoma SCC25 Cell Line

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This study was designed to investigate the role of miR-183 in modulating cell growth and apoptosis of tongue squamous cell carcinoma SCC25 cell line. Human squamous epithelial cell and squamous cell carcinoma cell line SCC25 was used, and miR-183 was inhibited. Cell growth, colony formation, and apoptotic rate, as well as the expression of caspase 3 and BCL-xL, were detected. Results showed that miR-183 was significantly overexpressed in the SCC25 cell line when compared with normal control. The miR-183 inhibitor reduced cell growth and colony formation, while the apoptosis percentage was significantly increased. The expression of activated caspase 3 and BCL-xL was obviously up- and downregulated in siRNA-transfected cells, respectively. In conclusion, miR-183 contributed to cell growth and proliferation, and suppressed cell apoptosis in SCC25 cells. Therefore, miR-183 might serve as a therapeutic target in tongue squamous cell carcinoma (TSCC).

**Key words:** Tongue squamous cell carcinoma (TSCC); miR-183; Cell proliferation; Apoptosis; SCC25 cells

### INTRODUCTION

Tongue squamous cell carcinoma (TSCC) is an aggressive head and neck squamous cell carcinoma with a high recurrence rate (1). Over 50% of TSCCs were diagnosed in an advanced stage, III and later (2), and more than 600,000 patients were registered with TSCCs every year (3). Surgery treatment is the first choice for patients diagnosed with TSCC due to metastasis and the high rate of cell proliferation (2). However, the quality of life for patients receiving surgery is low. Thus, it is crucial for the early diagnosis and intervention of TSCC.

MicroRNAs (miRNAs) are gene-specific regulators, and their dysregulation is common in cancers including TSCC, such as miR-184 (2), miR-7 (4), and miR-21 (5). For example, miR-184 was overexpressed in SCC patients, and inhibition of miR-184 in the TSCC cell line reduced the cell proliferation rate and oncogene c-Myc expression (2). Recently, another oncogene, miR-183, which had been reported to be overexpressed in several types of cancers (6–9), was detected to be obviously overexpressed in esophageal squamous cell carcinoma (OSCC), and the overexpression of miR-183

promoted cell proliferation of OSCC (10). miR-183 promotes cell growth and proliferation and inhibits cell apoptosis of cancer cells by modulating its target genes (6,9). However, the related mechanism is still unclear.

miR-183-modulated cell proliferation and apoptosis are related to numerous factors (6,10,11). For instance, Ren et al. (10) identified that a target of miR-183, programmed cell death 4 (PDCD4), could be modulated by the P13K/AKT signaling pathway, which is associated with cell proliferation (12) and related to caspase 3 activation. Caspase 3 is commonly activated by death-related signals (13,14). Moreover, BCL-x molecules, including antiapoptotic (BCL-xL) and proapoptotic (BCL-xS) members, are tightly associated with cell apoptosis (15). Overexpression of BCL-xL could reduce apoptosis (16,17) and block the expression of caspase 3 (17). However, the relationship between BCL-xL, activated caspase 3, miR-183 expression, and cell growth and the apoptotic rate in TSCC cells has not been reported until now.

This study aimed to investigate the association of miR-183 with cell growth, proliferation, and apoptosis.

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Moreover, miR-183 was silenced by a small miR-183-specific inhibitor to determine the effect of miR-183 on cell growth, proliferation, and apoptosis. This study would provide us with more information on the treatment and cure of TSCC.

## MATERIALS AND METHODS

### *Cell Line and Culture Conditions*

Human squamous epithelial cell and squamous cell carcinoma cell line SCC25 was obtained from the American Type Culture Collection (Manassas, VA, USA). Cell cultures were performed as previously reported (18). Cells were cultured in DMEM (Dulbecco's modified Eagle's medium) and Ham's F12 medium (1:1; both were obtained from Corning Cellgro Inc., Herndon, VA, USA) supplemented with 1.2 g/L sodium bicarbonate, 2.5 mM L-glutamine (Gibco-BRL, Gaithersburg, MD, USA), 15 mM HEPES, 0.5 mM sodium pyruvate, 400 ng/mL hydrocortisone (Sigma-Aldrich, St. Louis, MO, USA), and 10% heat-inactivated fetal bovine serum (FBS; Thermo Fisher Scientific Inc., Waltham, MA, USA). Cells were cultured at 37°C in a humidified incubator with 5% CO<sub>2</sub>.

### *Inhibition of miR-183 and Cell Transfection*

The miRNA inhibitors of miR-183 were chemically synthesized by GeneChem (Shanghai, P.R. China). To inhibit the expression of miR-183, we seeded SCC25 cells into 24-well plates and transfected them with miR-183-specific inhibitor and the negative control for 48 h. The transfection was done via Lipofectamine 2000 (Invitrogen Life Technologies, Carlsbad, CA, USA).

### *MTT Assay*

SCC25 cell proliferation was assessed using MTT (3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide; Sigma-Aldrich) assay (19). Briefly, cells at  $5.0 \times 10^3$  cells/well were seeded into 96-well plates at 0 and 72 h and 7 days posttransfection. The medium was changed every 3 days. Cells were additionally incubated in MTT solution for 2 h. The optical density of cells was determined at 570-nm absorbance (A<sub>570</sub>) using a microplate reader (Bio-Rad Labs, Sunnyvale, CA, USA). Each experiment was performed in triplicate.

### *Soft Agar Colony Formation Assay*

SCC25 cells transfected with siRNA were seeded into six-well plates at 1,000 cells/well containing DMEM medium plus 10% FBS for 24 h, followed by soft agar colony formation assay (20). Briefly, cells were cultured with fresh medium supplement at 37°C with 5% CO<sub>2</sub> for 3 weeks. The medium was changed every 3 days. Cell clones were stained with crystal violet, and colonies were counted under a microscopy.

### *Apoptosis Assay*

For cell apoptosis assay, SCC25 cells at  $1.0 \times 10^5$  cells/well were seeded into six-well plates containing DMEM. Next, cell apoptosis was determined by using the Annexin-V apoptosis detection kit (Bender MedSystems, Vienna, Austria) as previously reported (21). Cells were stained with annexin V/propidium iodide (PI) according to the manufacturer's instruction. FACScalibur flow cytometer (Becton Dickinson Immunocytometry Systems, San Jose, CA, USA) was used to determine apoptosis percentage. Each experiment was repeated three times.

### *Quantitative Real-Time Polymerase Chain Reaction (qRT-PCR)*

miRNA isolation of cells was performed using the miRvana miRNA isolation kit (Ambion, Austin, TX, USA). The cDNA were synthesized from miRNA with Mir-X miRNA First-Strand Synthesis Kit (Takara Bio Inc., Kyoto, Japan). Quantitative PCR analysis was conducted using the SYBR Premix Ex Taq (Takara Bio Inc.). U6 small RNA was used as the reference RNA to normalize relative expression of miR-183.

### *Western Blot Analysis*

Cultured SCC25 cells were harvested, washed in PBS, and lysed in RIPA buffer (Pierce, Rockford, IL, USA). BCA protein assay kit (Pierce) was used to measure protein concentration (10). Next, cell lysates were separated by 12% SDS-PAGE gels and electrophoretically transferred to polyvinylidene difluoride (PVDF) membranes (Invitrogen Corp., Carlsbad, CA, USA). The PVDF membranes were blocked with 5% skim milk and incubated with the primary antibody against activated caspase 3, BCL-xL and  $\beta$ -actin (1:1,000 dilution; Epitomics, Burlingame, CA, USA) at 4°C overnight. Next, membranes were incubated with secondary antibodies at room temperature for 1 h. The polypeptide bands were detected using an enhanced chemiluminescence (ECL) detection system (Pierce) and quantified with AlphaEase software (Alpha, USA).

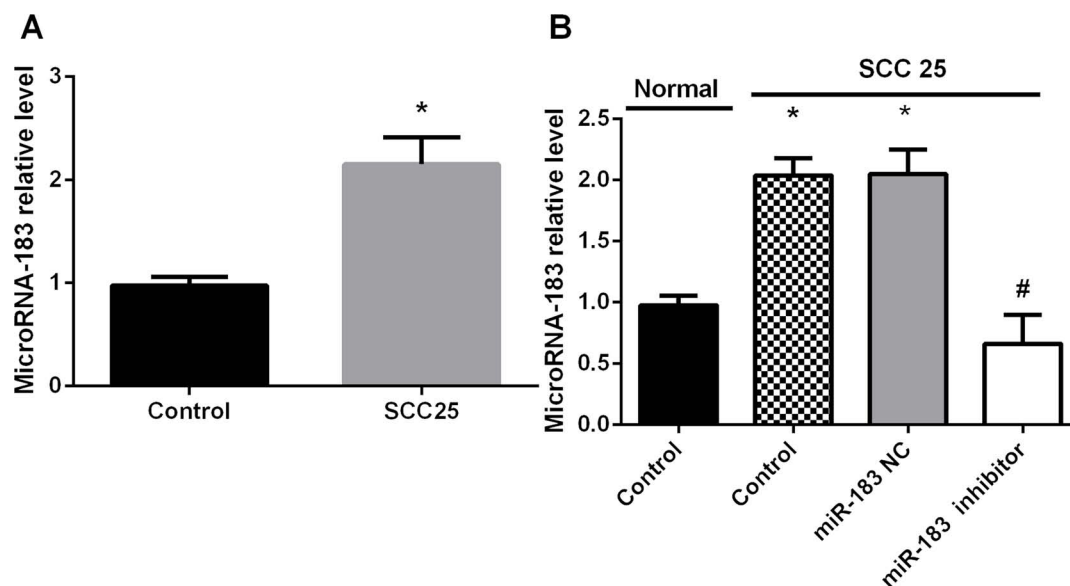
### *Statistical Analysis*

Quantitative data are expressed as mean  $\pm$  standard deviation (SD). SPSS 18.0 software (SPSS, Chicago, IL, USA) was used for statistical analysis. One-way analysis of variance (ANOVA) test was compared among groups. A statistical significance was defined as a value of  $p < 0.05$ .

## RESULTS

### *The Expression of miRNA-183 in SCC25 Cells*

QRT-PCR analysis showed that the expression of miR-183 was significantly higher in SCC25 cells than that in normal squamous epithelial cell ( $p < 0.05$ ) (Fig. 1A).



**Figure 1.** Expression of miR-183 in SCC25 cells. (A) Expression of miR-183 in normal squamous epithelial cells (Control) and SCC25 cells before miR-183 inhibition. (B) Expression of miR-183 in control, SCC25 cells before (SCC25) and after miR-183 inhibition. \* $p < 0.05$  versus Control; # $p < 0.05$  versus SCC25.

After being transfected with miR-183 inhibitors, the expression of miR-183 in transfected cells was obviously decreased compared with that in SCC25 without transfection ( $p < 0.05$ ) (Fig. 1B). This suggested that the miR-183 inhibitor successfully inhibited miR-183 expression.

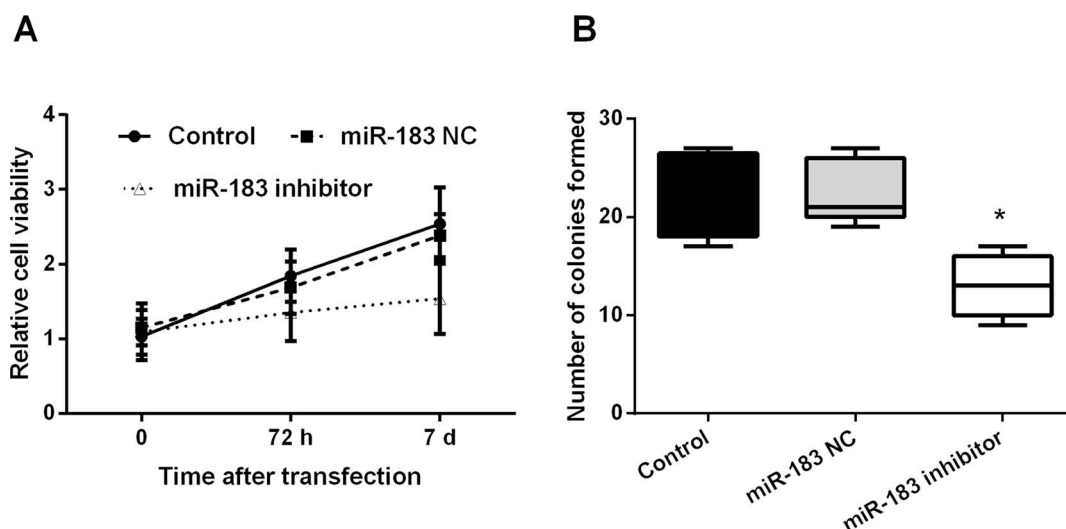
#### miR-183 Inhibition Reduces SCC25 Cell Viability and Colony Formation

To investigate the effect of miR-183 on SCC25 cell viability and colony formation ability, we performed MTT

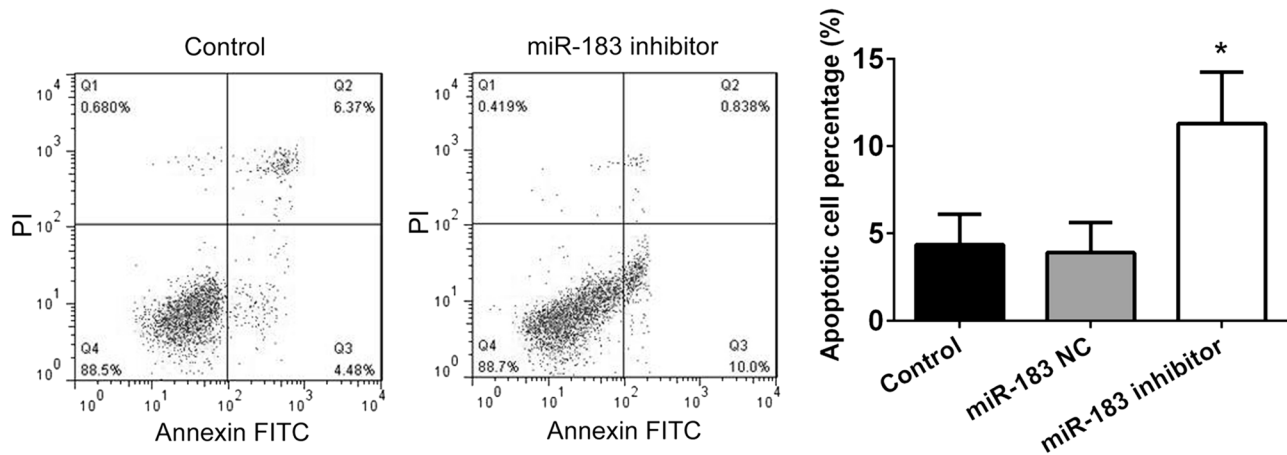
assay and soft agar colony formation assay. The results showed that miR-183 significantly reduced cell viability ( $p > 0.05$ ) (Fig. 2A) and colony formation ( $p < 0.05$ ) (Fig. 2B) of transfected cells, compared with controls. This suggested that miR-183 contributes to SCC25 cell growth, and the miR-183 inhibitor inhibited its function.

#### miR-183 Inhibitor Induces SCC25 Cell Apoptosis

The annexin V/PI method was conducted on SCC25 cells to explore the effect of miR-183 on SCC25 cell



**Figure 2.** The effect of miR-183 silence on SCC25 cell growth and colony formation. (A) miR-183 inhibition suppressed the cell viability of SCC25 cells. (B) miR-183 inhibition inhibited the colony formation of SCC25 cells. \* $p < 0.05$  versus Control.



**Figure 3.** The effect of miR-183 silencing on cell apoptosis. miR-183 inhibition significantly increased the apoptotic percentage of SCC25 cells. A four-quadrant diagram was detected by FACScalibur flow cytometer. \* $p < 0.05$  versus Control.

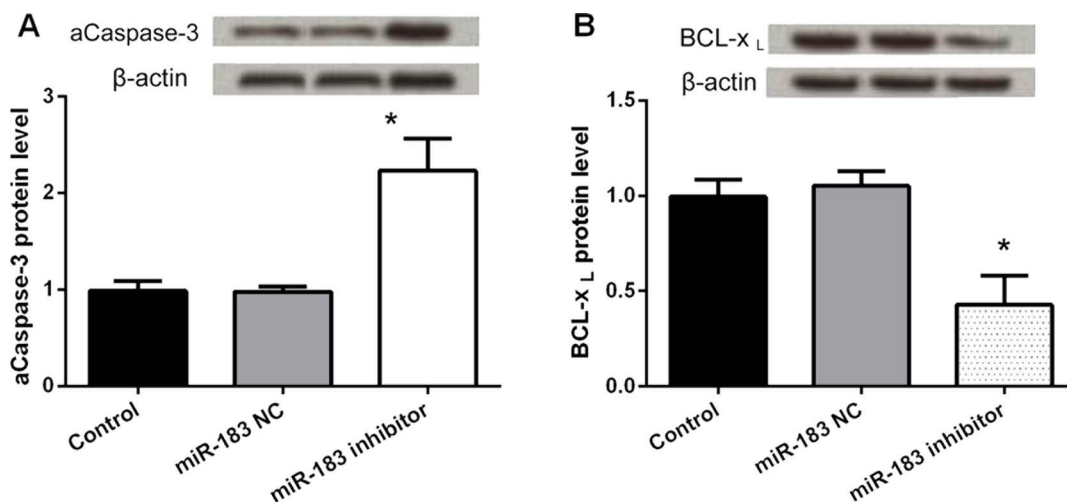
apoptosis. Compared with that in control cells, an obvious increment in apoptotic percentage was detected in SCC25 cells transfected with inhibitor ( $p < 0.05$ ) (Fig. 3). This fact demonstrated that miR-183 inhibition induced the apoptosis of SCC25 cells. For the determination of cell apoptosis factors, we detected the expression of activated caspase 3 and BCL-xL in transfected cells by Western blot analysis. Results showed that the expression of activated caspase 3 and BCL-xL was remarkably up- and downregulated in miR-183 inhibitor-transfected cells ( $p < 0.05$ ), respectively (Fig. 4).

### DISCUSSION

In this study, we detected that the expression level of miR-183 in SCC25 was relatively higher than that

in human squamous epithelial cells. To investigate the effect of miR-183 on cell growth and apoptosis, miR-183 expression in SCC25 cells was silenced by siRNA silencing technology. The results showed that miR-183 inhibition reduced cell viability and colony formation, and induced cell apoptotic percentage. Moreover, miR-183 inhibition significantly upregulated activated caspase 3 expression while downregulating BCL-xL expression. These data might suggest that miR-183 plays important role in SCC25 cell growth and apoptosis.

miR-183 was overexpressed and acted as an oncogene in several types of cancers, such as bladder cancer (6), gastric cancer (7,8), and colon cancer (9). Oncogenic miR-183 promotes cell growth and proliferation and inhibits cell apoptosis of cancer cells by modulating its



**Figure 4.** Expression of activated caspase 3 (aCaspase 3) and BCL-xL in transfected cells. Protein level of activated caspase 3 and BCL-xL was detected by Western blot analysis. The protein bands were detected using the ECL detection system and quantified with AlphaEase software. \* $p < 0.05$  versus Control.



target genes (6,9). For example, Li et al. (7) reported that miR-183 was markedly upregulated in gastric cancer. Ren et al. (10) identified miR-183 was obviously overexpressed in OSCC. These reports were in accordance with the results in this study that miR-183 was overexpressed in SCC25 when compared with normal human squamous epithelial cell. Moreover, data from long-term follow-up showed miR-183 to be associated with poor survival in patients with several cancers, including lung cancer (22) and glioma (23). This might suggest that miR-183 is involved in the pathogenesis of squamous cell carcinoma and might be a biomarker for overall survival of TSCC.

Furthermore, genetic methods were used to inhibit miRNA expression in cancer cell lines to explore the effect of miRNAs in cancer cell apoptosis and growth (7,10,24). For example, Li et al. (7) inhibited the endogenous miR-183 and found that cell proliferation and colony formation were remarkably inhibited, while cell apoptosis was significantly enhanced. Similarly, Ren et al. (10) found that overexpression of miR-183 promoted cell proliferation, while inhibition of it suppressed cell proliferation and invasion. In this study, we inhibited the expression of miR-183 and hence inhibited cell growth and colony formation while inducing cell apoptosis. These demonstrated that inhibition of miR-183 contributed to the inhibition of cancer cell growth; thus, the inhibition of miR-183 might be explored as a therapeutic target for TSCC.

BCL-x molecules including antiapoptotic (such as BCL-xL) and proapoptotic (such as BCL-xS) activity members belong to the BCL2 family (15). Overexpression of BCL-xL could reduce apoptosis, and the upregulation of BCL-xL had been determined in many types of cancer (16,17). Moreover, the expression of BCL-xL might block the expression of caspase 3 (17). The result in this study is that inhibition of miR-183 induced overexpression of activated caspase 3 in SCC25 cells, reduction of the BCL-xL protein, and decrease of the cell apoptosis rate. This demonstrated that proapoptotic factors including activated caspase 3 and antiapoptotic members, such as BCL-xL, responded to miR-183 dysregulation. However, the detailed mechanism related to miR-183-directed cell apoptosis should be explored with more experiments.

In summary, this study demonstrated that miR-183 was crucial for SCC25 cell growth, cell proliferation, and cell apoptosis. The inhibition of miR-183 by inhibitors suppressed cell growth and proliferation while inducing apoptosis of SCC25. Also, miR-183 inhibition upregulated activated caspase 3 expression while decreasing BCL-xL expression. Taken together, these results showed that miR-183 contributed to cell growth and proliferation and suppressed cell apoptosis in SCC25 cells. More animal and clinical experiments should be done to explore

whether miR-183 could be used as a biomarker for diagnosis and cure treatment for TSCC.

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