

MicroRNA-15a Inhibits Proliferation and Induces Apoptosis in CNE1 Nasopharyngeal Carcinoma Cells

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Nasopharyngeal carcinoma (NPC) is a highly metastatic cancer, frequently occurring in Southeast Asia and Southern China. Several microRNAs (miRNAs) have been shown to have an inhibitive effect on NPC, while the effect of miR-15a on NPC remains unclear. Thus, our study aimed to investigate the potential effect of miR-15a on NPC cell proliferation, apoptosis, and possible functional mechanism. Human NPC CNE1 cells were transfected with miR-15a mimics, miR-15a inhibitors, or a control. Afterward, cell viability and apoptosis were assayed by using CCK-8, BrdU assay, and flow cytometry. Moreover, Western blot was used to detect the expression changes of proliferation and apoptosis of related proteins. As a result, miR-15a overexpression significantly reduced cell proliferation ($p < 0.01$ or $p < 0.001$) and induced cell apoptosis ($p < 0.001$), while miR-15a suppression got the opposite result for cell proliferation and apoptosis. In addition, miR-15a overexpression upregulated the protein levels of p27, GSK-3 β , Bax, procaspase 3, and active caspase 3, whereas miR-15a suppression downregulated these proteins. The protein level of p21 was not significantly regulated by miR-15a overexpression or suppression. These results indicated that miR-15a played a role for inhibition of proliferation and induction of apoptosis in CNE1 cells.

Key words: miR-15a; Nasopharyngeal carcinoma (NPC); Cell proliferation; Apoptosis; GSK-3 β ; p27

INTRODUCTION

Nasopharyngeal carcinoma (NPC) is an invasive and highly metastatic cancer arising in nasopharyngeal epithelial cells, with high incidence in Southeast Asia and Southern China (1). The exact cause of NPC is not yet well understood, but genetic susceptibility, Epstein-Barr virus (EBV) infection, and living environment have been identified as the three major etiologic factors during NPC pathogenesis (2). Accordingly, a variety of treatment methods have been investigated to play a certain role in the clinical treatment of NPC. Among these methods, chemo- and radiotherapy are two primary therapeutic strategies of NPC, while the 5-year overall survival rate remains less than 50% (3,4). Hence, a better understanding of the molecular mechanisms of NPC will be helpful for the development of novel therapeutic and diagnostic strategies.

MicroRNAs (miRNAs) are small noncoding RNAs that tune gene expression and modulate target gene functions (5–7) and have already been implicated in a growing number of cancers (8). Several miRNAs, such as miR-29c, miR-24, and miR-608, have been shown to target specific mRNAs to regulate NPC development and

progression (9). Thus, it seems that modulation of miRNA could be a new approach for gene therapy of NPC (10). miR-15a has been reported as an inhibitor of DNA synthesis (11) and a tumor suppressor (12) in a variety of cancers. For example, miR-15a, when associated with the other two miRNAs, inhibits pituitary tumor cell proliferation (13) and induces cell apoptosis in non-small cell lung cancer (14). However details of the function and molecular mechanism of miR-15a on NPC cell proliferation and apoptosis have not been well investigated.

Therefore, in the current study, either miR-15a mimics, miR-15a inhibitors, or controls were transfected into human NPC cells (CNE1). The effect of miR-15a on human CNE1 cell proliferation and apoptosis was investigated in vitro. Moreover, the protein levels of p27, p21, glycogen synthase kinase-3 β (GSK-3 β), BCL2-associated X protein (Bax), procaspase 3, and active caspase 3 in cells were detected, to reveal the possible molecular mechanism of miR-15a in CNE1 cells. This study provided a partial understanding about the role of miR-15a in NPC and evidenced that miR-15a might be used in NPC treatment as a novel gene therapeutic target.

MATERIALS AND METHODS

Cell Culture and Transfection

Human NPC CNE1 cells were obtained from Cell Bank of the Chinese Academy of Sciences (Shanghai, China). The cells were cultured in RPMI-1640 medium (Gibco-BRL, Gaithersburg, MD, USA) containing 10% fetal bovine serum (FBS; Hyclone, Logan, UT, USA) at 37°C in humidified 5% CO₂ incubator.

Cells were seeded into six-well plates at 5 × 10⁴ per well and incubated for 24 h at 37°C. After the cells were grown to about 60% confluence, cells were transfected with miR-15a mimics, miR-15a inhibitors, or control for 48 h, by using Lipofectamine RNAiMAX reagent (Thermo Fisher Scientific, Cleveland, OH, USA), according to the manufacturer's protocol. miR-15a mimics, miR-15a inhibitors, and control were designed and produced from GenePharma (Shanghai, China).

Cell Viability Assay

Cell viability was determined using Cell Counting Kit-8 (CCK-8; Dojindo, Kumamoto Prefecture, Kyushu, Japan), according to the manufacturer's instruction. In brief, after 48 h of transfection, the cells were seeded in 96-well plates at 2 × 10³ cells per well and incubated for 24 h, then 20 µl of CCK-8 was added to each well and incubated for another 4 h at 37°C. Afterward, the absorbance at 450 nm was measured by a microplate reader (Bio-Rad Laboratories, Hercules, CA, USA).

BrdU Staining Assay

Cell viability was further measured using bromodeoxyuridine (BrdU) assay kit (Roche Diagnostics GmbH, Germany). For BrdU assay, miR-transfected cells were seeded in six-well plates at 2 × 10⁴ cells per well on sterilized coverslips. After 72 h, 10 µM of BrdU solution was added to each well, and the cells were incubated for 5 h. BrdU integration was detected by labeling with 100 µl of anti-BrdU-POD monoclonal antibody and incubating at room temperature for 30 min. Afterward, 100 µl of peroxidase substrate with substrate enhancer was added to each well and incubated at room temperature for another 15 min. Finally, immune complexes were detected by reading absorbance at 490 nm (OD₄₉₀).

Apoptosis Assay

Cell apoptosis was performed by the Annexin-V/FITC and PI apoptosis detection kit (Becton Dickinson, Franklin Lakes, NJ, USA), according to manufacturer's recommendations. Briefly, miR-transfected cells were collected and resuspended in 200 µl of binding buffer containing 10 µl Annexin-V-FITC and 5 µl of PI, then incubated at room temperature for 30 min in the dark.

Apoptotic cells were discriminated by FACS Calibur flow cytometer (Becton Dickinson).

Western Blot

The protein expression changes in miR-transfected cells were detected by Western blot analyses. The cells were lysed, and protein concentration was quantified using the Bradford method. Equal amounts of protein were resolved by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to polyvinylidene fluoride (PVDF) membranes. Blots were blocked for 1 h in 5% nonfat dry milk and then incubated overnight with the primary antibodies: GSK-3β (ab93926), Bax (ab32503), procaspase 3 (ab32150), active caspase 3 (ab2302), p27 (ab54563), p21 (ab7960), and actin (ab3280) (Abcam, Cambridge, MA, USA). The blots were then incubated for 1 h with horseradish peroxidase (HRP)-conjugated secondary antibodies and visualized by chemiluminescence using the enhanced chemiluminescence (ECL) reagent (GE Healthcare, Little Chalfont, UK), and data were analyzed using Image Lab (Bio-Rad) software.

Statistical Analysis

All data were expressed as means ± standard deviations (SD). The data were analyzed using one-way analysis of variance (ANOVA). Results with a value of $p < 0.05$ were considered statistically significant. GraphPad Prism 5 (GraphPad Software Inc., San Diego, CA, USA) was used for these analyses.

RESULTS

miR-15a Overexpression Reduced CNE1 Cell Proliferation

To test the effect of miR-15a on NPC cells proliferation, CNE1 cells were transfected with miR-15a mimics, miR-15a inhibitors, or control, and then cell viability was measured by CCK-8 kit after the cells were cultured for 1, 2, 3, 4, and 5 days, respectively. Results shown in Figure 1 indicate that miR-15a overexpression could significantly reduce cell proliferation at 3 days ($p < 0.01$), 4 days ($p < 0.001$), and 5 days ($p < 0.001$) when compared with the control group. miR-15a suppression could significantly promote cell proliferation at the same time points ($p < 0.001$, $p < 0.001$, and $p < 0.01$) when compared with the control group. These results revealed that miR-15a might play an important role in NPC cell proliferation.

miR-15a Overexpression Reduced Cell Proliferation via Regulating p27

To further confirm the role of miR-15a in NPC cell proliferation, BrdU staining was performed, and the expression

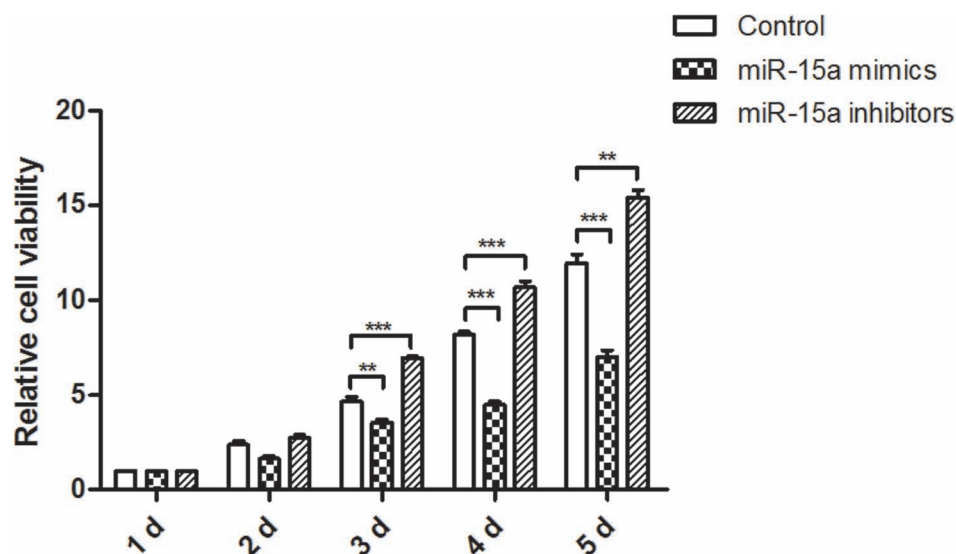


Figure 1. miR-15a overexpression reduced proliferation in CNE1 cells. After transfection with miR-15a mimics, miR-15a inhibitors, or control, cell viability was measured by CCK-8 methods at 1, 2, 3, 4, and 5 days. CCK-8, Cell Counting Kit-8. ** $p < 0.01$ compared with control group; *** $p < 0.001$ compared with control group.

levels of p21 and p27 were conducted by Western blot assay. BrdU assay results (Fig. 2A, B) confirmed that miR-15a overexpression could significantly reduce cell proliferation ($p < 0.001$), and miR-15a suppression could significantly promote cell proliferation ($p < 0.05$) when compared with the control group. Western blot results (Fig. 2C) showed

that the expression level of p27 was significantly upregulated by miR-15a overexpression and significantly downregulated by miR-15a suppression, whereas the expression level of p21 was unaffected. Hence, miR-15a might affect cell proliferation via regulating the expression of p27 but not p21.

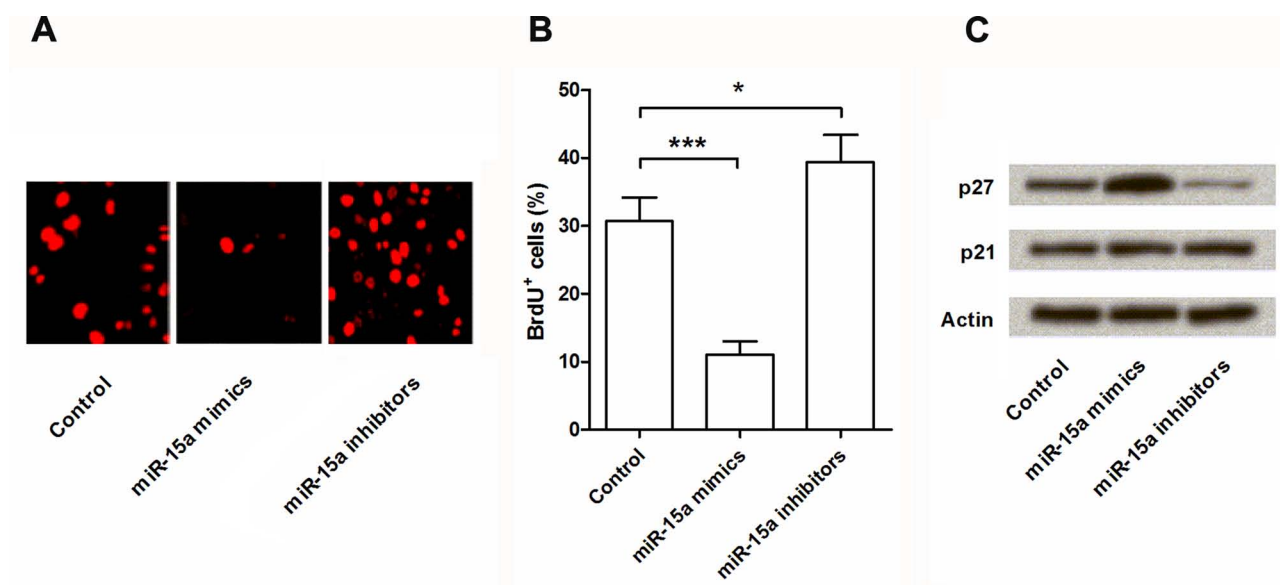


Figure 2. miR-15a overexpression reduced cell proliferation by regulating p27. After transfection with miR-15a mimics, miR-15a inhibitors, or control, cell viability was detected by bromodeoxyuridine (BrdU) assay (A, B). The protein expressions of p27 and p21 in the transfected cells were measured by Western blot analysis (C). Actin acted as an internal control. BrdU+ cells, BrdU-positive cells. * $p < 0.05$ compared with control group; *** $p < 0.001$ compared with control group.

miR-15a Overexpression Promoted CNE1 Cell Apoptosis

To address the effect of miR-15a on NPC cells apoptosis, CNE1 cells were transfected with miR-15a mimics, miR-15a inhibitors, or the control, and then apoptotic cells were classified by flow cytometry. Results in Figure 3A and B showed that miR-15a overexpression could significantly increase the rate of apoptosis ($p < 0.001$) when compared with the control group. However, miR-15a suppression could significantly decrease the rate of apoptosis ($p < 0.05$) when compared with the control group. These results indicated that miR-15a might be involved in NPC cells apoptosis.

miR-15a Overexpression Induced Apoptosis by Regulating GSK-3 β

To define the molecular mechanism of the impact of miR-15a on NPC cell apoptosis, miR-15a mimics, miR-15a inhibitors, or the control were transfected with CNE1 cells, and protein levels of GSK-3 β , Bax, procaspase 3, and active caspase 3 in cells were measured by Western blot analysis. As shown in Figure 4A and B, these four proteins were significantly upregulated by miR-15a overexpression ($p < 0.001$, $p < 0.01$, $p < 0.001$, and $p < 0.001$) when compared with the control group. These four proteins were significantly downregulated by miR-15a suppression ($p < 0.01$, $p < 0.05$, $p < 0.05$, and $p < 0.05$) when

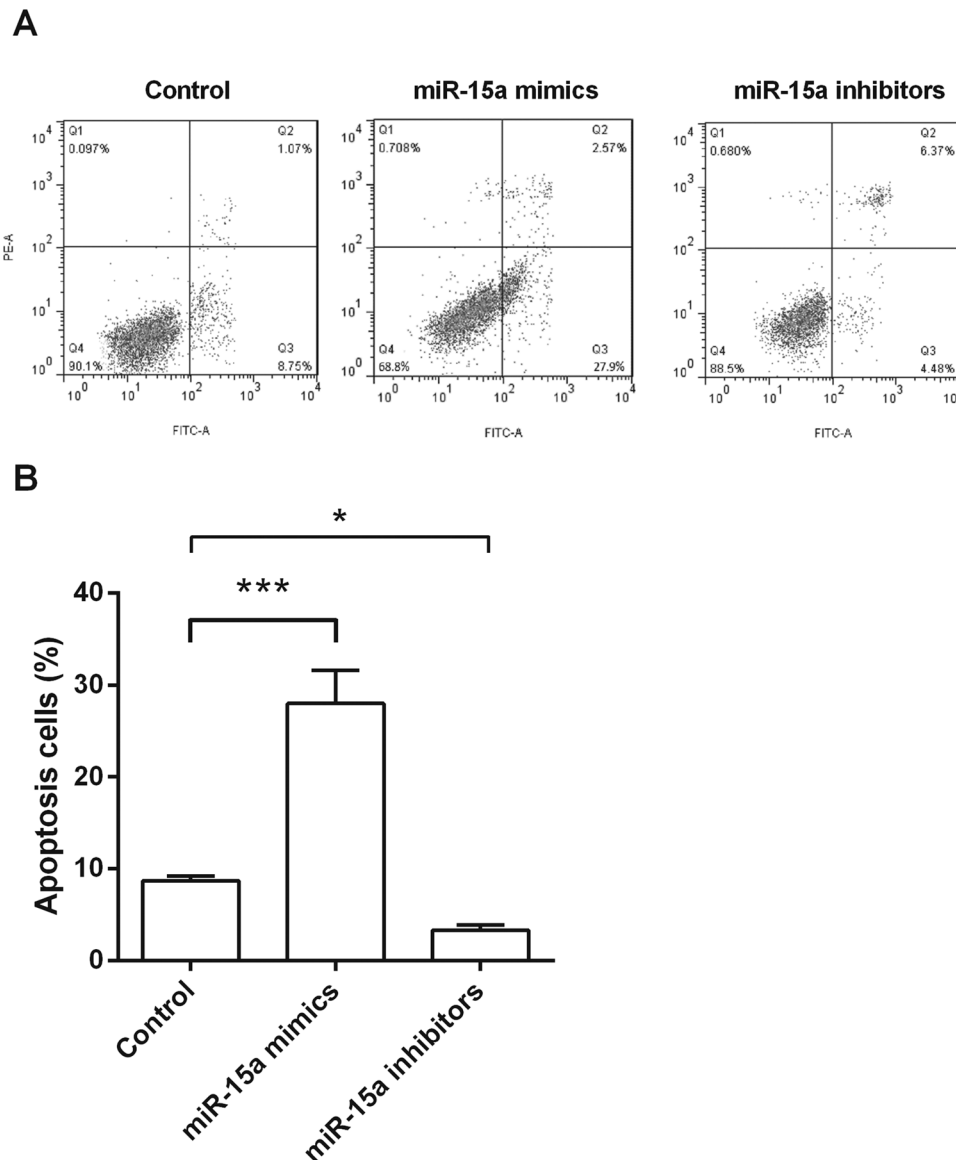


Figure 3. miR-15a overexpression induced apoptosis in CNE1 cells. After transfection with miR-15a mimics, miR-15a inhibitors, or control, cell apoptosis was measured by flow cytometry. * $p < 0.05$ compared with control group; *** $p < 0.001$ compared with control group.

compared with the control group. These results indicated that the effect of miR-15a on NPC cell apoptosis might be through regulating the protein expression of GSK-3 β , Bax, procaspase 3, and active caspase 3.

DISCUSSION

NPC is known as a prevalent malignant neoplasm frequently occurring in Asia, and there are many patients suffering with a poor prognosis. Recent studies have confirmed that several miRNAs could regulate the development and progression of NPC. In the current study we presented in vitro data suggesting a possible

role for miR-15a in proliferation inhibition and apoptosis induction of NPC. Our study found that miR-15a overexpression could significantly reduce proliferation and induce apoptosis of CNE1 cells. In addition, the protein expression of p27, GSK-3 β , Bax, procaspase 3, and active caspase 3 in CNE1 cells was upregulated by transfected with miR-15a.

miR-15a is a highly conserved RNA, which is located at chromosome 13q14 and has been implicated in multiple biological processes in recent years. It has been reported that miR-15a overexpression inhibited cell growth by arresting cells at G₁ phase (15). Moreover, it has been

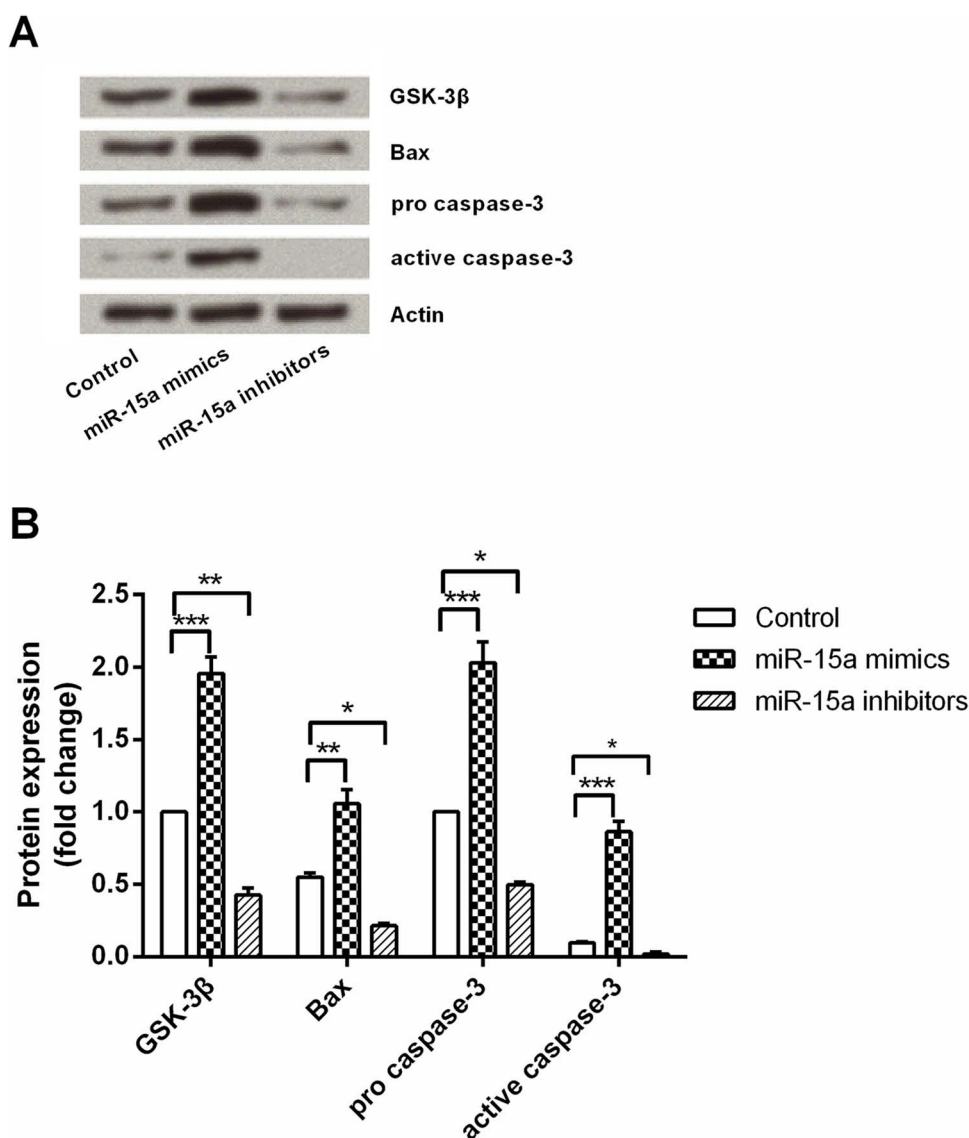


Figure 4. miR-15a overexpression induced cell apoptosis by regulating GSK-3 β , Bax, and caspase 3. After transfection with miR-15a mimics, miR-15a inhibitors, or control, the protein expression of GSK-3 β , Bax, procaspase 3, and active caspase 3 in the transfected cells were measured by Western blot analysis. Actin acted as an internal control. GSK-3 β , glycogen synthase kinase-3 β ; Bax, BCL2-associated x protein. * p <0.05 compared with control group; ** p <0.01 compared with control group; *** p <0.001 compared with control group.

found to induce apoptosis by targeting B-cell lymphoma/leukemia-2 (Bcl-2) both in vitro and in vivo (16). In the study of Zhang et al. (17) miR-15a inhibited cell proliferation and increased the apoptosis of CNE2 cells by regulating Bcl-2, caspase 2, and caspase 3. Consistent with these previous investigations, our study demonstrated that miR-15a could significantly inhibit cell proliferation and induce apoptosis in NPC cells.

In order to reveal the underlying molecular mechanisms by which of miR-15a affects NPC cell proliferation and apoptosis, the protein expression levels of p27, p21, GSK-3 β , Bax, procaspase 3, and active caspase 3 were assessed. Our results indicated that miR-15a overexpression upregulated the level of p27, GSK-3 β , Bax, procaspase 3, and active caspase 3, but not p21. p27 and p21 are two cyclin-dependent kinase inhibitors, which block the cell cycle at the G₀/G₁ phase and thus restrain cell proliferation (18). Thus, our study indicated that miR-15a might inhibit the proliferation in CNE1 cells via regulating the protein expression of p27 but not p21. Similarly, Kang et al. (19) demonstrated that miR-15a exerted an inhibitive function in gastric adenocarcinoma cell viability by targeting p27.

Recently, it has been reported that several signaling pathways and proteins were involved in the effect of miR-15a on cancer cells apoptosis, such as reactive oxygen species (ROS)–p53 pathway (20), Bcl-2, and caspase 3 (17,21). However, we suggested the first in vitro evidence that miR-15a induced cell apoptosis by regulating GSK-3 β . In fact, GSK-3 β has been implicated in apoptosis under a variety of conditions, including DNA damage, endoplasmic reticulum stress, and hypoxia (22). It has been identified as a target of Akt, which plays a role in the regulation of apoptosis (23), and the activated GSK-3 β activates downstream proapoptotic proteins (e.g., caspase 3 and Bax) (24,25). Sun et al. demonstrated that inhibition of GSK-3 β could inhibit apoptosis, leading to caspase 8 and caspase 3 activation in breast cancer cells (26). Bax is one of the subfamily member of Bcl-2, and it is identified as a proapoptotic protein (27). Usually, Bax translocates to the mitochondria and then involves homooligomerization and releases cytochrome c (28), finally activating caspase and inducing apoptosis (29). However, caspase 3 is the most prevalent caspase in the cell, and active caspase 3 is the one ultimately responsible for the majority of the apoptotic effects (30). Therefore, in the current study miR-15a might upregulate the expression of the three apoptotic relative proteins (i.e., GSK-3 β , Bax, and caspase 3) and thus induce NPC cell apoptosis.

Taken together, the results of this study found that miR-15a inhibited cell proliferation and induced apoptosis in CNE1 cells, possibly by regulating the protein expression of p27, GSK-3 β , Bax, and caspase 3. These results demonstrated that miR-15a might be a potential target for the

treatment of NPC in the future. However, further study of miR-15a on NPC still needs to be done.

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