MicroRNA-338 inhibits migration and proliferation by targeting hypoxia-induced factor 1α in nasopharyngeal carcinoma

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Abstract. Nasopharyngeal cancer (NPC) is an endemic type of head and neck cancer with a high rate of cervical lymph node metastasis. An increasing number of studies have shown that microRNAs (miRNAs) play a key role in the development and progression of NPC. miR-338-3p has been demonstrated as an anti-oncogene in different solid tumors. The aim of the present study was to investigate the potential role of miR-338-3p in the development and progression of NPC. Compared with normal samples, our data showed that miR-338-3p were downregulated in NPC tissues and cells. The luciferase assay demonstrated that HIF-1 α was a direct target of miR-338-3p. We also found that miR-338-3p regulated the expression levels of HIF-1a, respectively. Overexpression of miR-338-3p in NPC cells significantly inhibited cell proliferation, and migration. Conversely, miR-338-3p knockdown in cells with lower endogenous expression levels significantly reduced antitumor behavior. Furthermore, enforced expression of miR-338-3p led to a decline in ERK phosphorylation as well as inhibited the hypoxia induced epithelial to mesenchymal transition. Cells pre-transfected with miR-338-3p can overcome hypoxiamediated cisplatin resistance. Taken together, we found that miR-338-3p directly targeted HIF-1 α , and we provide insight into NPC initiation and progression, possibly representing a novel therapeutic target.

Introduction

Nasopharyngeal carcinoma (NPC) is a common head and neck cancer derived from epithelium cells which were located

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in the nasopharynx (1). NPC is notorious for its metastatic potential at the early stages of the disease via both lymph and blood vessels. Due to the recurrence and distant metastasis, the prognosis of NPC patients is very poor (1). Furthermore, drug resistance may hamper the efficacy of anti-cancer drugs (2). Therefore, there is a great and urgent need to develop early diagnostic or predictive markers for NPC and to elucidate the mechanisms that would allow the development of efficient treatment options.

Recent studies of microRNAs (miRNAs) suggest a potential role of these regulatory molecules as candidate oncogenes or tumor suppressors in cancer progression (3-5). miRNAs are small non-coding RNA molecules ~19-25 nucleotide, which exist in many organisms and regulate gene expression through degradation of the corresponding mRNAs and/or inhibiting their translation by binding to their 3'-UTRs (1,6,7). miRNAs have been reported to participate in a variety of pathways in physiological and pathological processes such as cellular differentiation, proliferation and apoptosis (1,8), metastasis and resistance to therapy (6). Dysregulated expression of miRNAs has been reported in most tumor types, including NPC (1). At present, several miRNAs have been shown to target specific mRNAs to regulate the progression of NPC (9). For example, miR-29c suppressed metastasis by targeting TIAM1 and enhanced the sensitivity to cisplatin based chemotherapy and radiotherapy in nasopharyngeal carcinoma (10,11). miR-451, by targeting MIF, inhibited cell growth and invasion and was associated with survival of NPC (1). Furthermore, miR-218 (12), miR-26a/b (13,14), miR-216b (15), miR-10b (16), miR-141 (17) and miR-200a (18) have been shown to have tumor-suppressive functions in NPC. The dysregulated miRNAs are involved in NPC development and progression by regulating cell growth, proliferation, apoptosis, invasion and metastasis (1), indicating that miRNAs play important roles in NPC tumorigenesis.

Recently, there has been increasing interest in miR-338-3p as an anti-oncogene in different solid tumors, including gastric cancer (19), hepatocellular carcinoma (20) and malignant melanoma (21). For example, miR-338-3p suppressed liver cancer cell invasion by targeting smoothened (7). Moreover, miR-338 can decrease proliferative and migratory, invasive behavior by attenuating the expression of NRP1 in gastric cancer (22). However, there is no study on the functions and mechanisms of miR-338-3p in NPC development and progression.

Hypoxia or low oxygen tension has emerged as a specific and general feature of the microenvironment of many malignant tumors (23), including nasopharyngeal cancer (24). Tumor hypoxia is known to be mainly responsible for tumor resistance to radiotherapy and chemotherapy as well as to promote tumor phenotype influencing invasiveness, metastasis and poor prognosis (24). Studies showed that 100% of primary NPC and 58% of cervical nodal metastases of NPC were found with hypoxic regions which increased distant metastases, as well as resistance to chemotherapy in advanced NPC patients (25). There is emerging evidence shows that the adaptive response to hypoxia is also mediated by HIF-1dependent pathways in cancer as well as in nasopharyngeal carcinoma (24,26). Among the proteins associated with tumor hypoxia, hypoxia-inducible factor 1-alpha (HIF-1α) is an important hypoxia regulatory molecule promoting angiogenesis and metastasis (25). HIF-1a is commonly found overexpressed in NPC (27). HIF-1 α has been reported as a prognostic factor as well as a potential therapeutic target of NPC (25). HIF-1 α can be transcriptionally and translationally regulated by signaling molecules such as cytokines growth factors and microRNAs (28). However, the association of HIF-1 α with miR-338-3p has not been established in NPC.

In the present study, we found that decreased expression of miR-338-3p had a causal role as a tumor suppressor in NPC by target HIF-1 α . miR-338-3p was downregulated in NPC tissues and cells. We found that miR-338-3p regulated the expression levels of HIF-1 α of CNE2 cells under hypoxia, respectively. The luciferase assay showed that HIF-1 α was a direct target of miR-338-3p. Overexpression of miR-338-3p suppressed the proliferation and migration of CNE2 cells under hypoxia, whereas the inhibition of miR-338-3p promoted these processes. Furthermore, our results suggested that the chemosensitizing effect of miR-338-3p may be an important feature for its potential therapeutic roles in NPC.

Materials and methods

Clinical specimens and cell culture. All NPC specimens and normal nasopharyngeal epithelium samples were obtained from the Affiliated Hospital of Nantong University. No patients had received any antitumor treatments before biopsy. The patients were diagnosed via histopathological evidence. The research protocols were approved by the Academic Committee of Nantong University.

The human immortalized nasopharyngeal epithelial cell line NP69 and NPC cell line CNE2 were received as a kind gift from the Sun Yat-Sen University of China. The NPC cell lines CNE1 and 5-8F, 6-10B were a kind gift from the Xiang-Ya School of Medicine, Central South University, China. The human immortalized nasopharyngeal epithelial cell line NP69 was maintained in keratinocyte/serum-free medium (Invitrogen) supplemented with bovine pituitary extract (BD Biosciences), human NPC cell lines (CNE-1, CNE-2, 5-8F and 6-10B) were cultured in RPMI-1640 (Gibco) supplemented with 10% FBS (Gibco). 5-8F and 6-10B cells were cultured in completed medium with 100 U/ml penicillin and 100 μ g/ ml streptomycin (Shanghai Genebase Gen-Tech Co., Ltd., Shanghai, China). All the cells were cultured at 37°C in a 5% CO₂ incubator. Hypoxic condition was induced by exposing the cells in a hypoxia modular incubator chamber (Billups-Rothenberg) that provided hypoxic conditions with $1\% O_2$, $5\% CO_2$ and $94\% N_2$.

Cell transfection. CNE2 cells were seeded in 6-well plates and cultured overnight at 37°C in a humidified atmosphere of 95% air and 5% CO₂. Subsequently, transfections were conducted for miR-338-3p mimics, miR-338-3p inhibitor and non-specific control (NC) using HiPerFect transfection reagent (Qiagen) according to the manufacturer's instructions. Following culture for a further 48 h, total RNA and cellular protein lysates were collected and used for reverse transcription-quantitative PCR (RT-qPCR) and western blot analysis, respectively.

Quantitative real-time PCR (qRT-PCR) analysis. miR-338-3p expression in NPC cells and tissues compared with normal was measured with SYBR qRT-PCR. Total RNA was extracted with TRIzol (Invitrogen) according to the manufacturer's instructions. miR-388-3p cDNA was synthesized from 2 μ g of total RNA. Subsequently, mRNA expression was analyzed by quantitative PCR using primers designed and synthesized by Guangzhou RiboBio Co., Ltd. (Guangzhou, China). Briefly, 20 μ l reactions containing 2 μ l RT product, 9 μ l of SYBR-Green PCR Master Mix and 200 nM of primers were subjected to 1 cycle of 95°C for 20 sec, and then 40 cycles of 95°C for 10 sec, 60°C for 20 sec and 70°C for 10 sec. miR-338-3p expression was normalized to U6 RNA. The 2^{- $\Delta\Delta$ CT} method was used to quantify the expression changes of target genes. Three independent experiments were performed.

Western blot analysis. HIF-1a proteins were measured by western blot analysis. Total protein was extracted from transfected cells using the RIPA lysis buffer (Beyotime Institute of Biotechnology, Haimen, China) according to the manufacturer's instructions. For western blot analysis, equal amounts of protein samples (20 μ g) were separated by 10% SDS-PAGE and transferred onto PVDF membranes (Millipore, Billerica, MA, USA). Membranes were blocked using with 5% skim milk in TBST. After 2 h at room temperature, the membranes were incubated overnight with polyclonal primary antibodies. The antibodies used were as follows: anti-HIF-1a (Abcam; 1:2,000 dilution), anti-p-ERK (Santa Cruz Biotechnology; 1:500 dilution), anti-ERK (Santa Cruz Biotechnology; 1:500 dilution) and mouse anti-β-actin (Santa Cruz Biotechnology; 1:1,000 dilution). Blots were then incubated with goat anti-rabbit or anti-mouse secondary antibody conjugated to horseradish peroxidase (Santa Cruz Biotechnology; 1:1,000 dilution) and visualized by enhanced chemiluminescence (ECL; Cell Signaling Technology). The values are representative of at least three independent experiments.

Luciferase reporter assay. HEK293 cells were seeded in the inner wells of 24-well plates. Then, cells were co-transfected using Lipofectamine 2000 (Invitrogen) with 80 ng of the pGL3-vector plasmid harboring the wild-type or mutant 3'-UTR of HIF-1 α and 4 ng of the pRL-TK. For each plate, the hsa-miR-338-3p mimics or mimics NC (Biomics Biotechnologies Co., Ltd., Nantong, China) was co-transfected at a final concentration of 50 pmol. Luciferase activities were measured

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consecutively 48 h post-transfection using the Dual-Luciferase reporter assay system (Promega, Southampton, UK) according to the manufacturer's instructions. Luminescence signals served as a measure for reporter activity normalized for transfection efficiency.

Wound healing assay. Wound healing assay was conducted 48 h after cell transfection. An artificial homogeneous wound was created on the monolayer using a sterized 200 μ l micropipette tip when the cells reached ~90% confluency. Cell debris was removed by washing with RPMI-1640 twice. Wound closure was recorded and observed by light microscopy and images were captured at the indicated time-points.

Immunocytochemical analysis. Cells cultured on glass coverslips were transfected with miR-338-3p mimics, miR-338-3p inhibitor and non-specific control (NC) and then cultured under hypoxia condition for 24 h. Glass coverslips were fixed with 4% paraformaldehyde and washed in PBS. The cells were incubated with rabbit anti-HIF-1 α antibody (Abcam; 1:200 dilution) overnight at 4°C. PBS washed and incubated with fluorescein isothiocyanate (FITC) labeled secondary antibodies (EarthBox, Lancaster, PA, USA; 1:1,000) and at the same time the nuclei were labeled with Hoechst (Invitrogen, Carlsbad, CA, USA). The coverslips were then observed under an Olympus camera.

Cell viability assay (Cell Counting kit-8). Cells were seeded into 96-well plates $(1 \times 10^4 \text{ cells/well})$ and treated as indicated. Cell viability was assessed by Cell Counting kit-8 assay (Beyotime Institute of Biotechnology, Shanghai, China). The absorbance of each well was read on a microplate reader (F-2500 fluorescence spectrophotometer; Hitachi) at 450 nm. Three independent experiments were performed in quintuplicate.

Migration assays. For the Transwell migration assays, $5x10^4$ CNE2 cells were plated in a serum-free medium in the top chamber with a non-coated membrane (24-well insert; 8 μ m pore size; Millipore) and a medium supplemented with 10% serum was in the lower chamber. The cells were incubated for 24 h under hypoxia conditions. The non-migrated cells were removed from the upper sides of the Transwell membrane filter inserts using cotton-tipped swabs. The migrated/invaded cells on the lower sides were stained with crystal violet and the cells were counted.

Flow cytometry. Cells were plated in 6-well plates at a specific density. The cells were treated with miR-338-3p mimics, inhibitor or miR-NC under hypoxic conditions for 24 h. After 24 h, the cells were collected and analyzed using an Annexin V-FITC apoptosis detection kit (BD Biosciences, Oxford, UK). The apoptotic cells were detected by flow cytometry.

Statistical analysis. All the data are expressed as mean \pm SEM. Data were compared using the Student's t-test. P<0.05 was considered to indicate a statistically significant difference. Each experiment consisted of at least three replicates per condition.

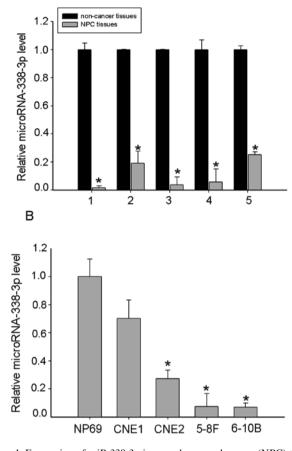


Figure 1. Expression of miR-338-3p in nasopharyngeal cancer (NPC) tissue samples and cells. (A) The expression levels of miR-338-3p in NPC and non-cancer tissues were measured by RT-qPCR. U6 snRNA was used as an internal control for miR-338-3p. Data are the means \pm SD. ^{*}P<0.05. (B) miR-338-3p was decreased in the NPC cell lines (CNE1, CNE2, 6-10B and 5-8F) compared with the immortalized normal nasopharyngeal epithelial NP69 cells.

Results

miR-338-3p is downregulated in NPC clinical specimens and cell lines. To determine whether miR-338-3p is involved in regulation of human NPC tumorigenesis, we firstly tested miR-338-3p expression in 5 freshly-frozen NPC and normal nasopharyngeal epithelial tissue samples and found that the miR-338-3p expression was significantly downregulated in NPC tissues (Fig. 1A). We also found miR-338-3p underexpressed in NPC cell lines CNE1, CNE2, 6-10B and 5-8F, compared to the normal nasopharyngeal epithelial cell line NP69 (Fig. 1B). Taken together, these findings suggested that miR-338-3p was downregulated in NPC.

miR-338-3p directly targets oncogenic HIF-1a. miRNAs have been shown to target specific genes to regulate the progression of NPC. To identify miR-338-3p target genes, we used the target prediction program, Bioinformatic analysis using the TargetScan (http://www.targetscan.org) and miRanda algorithm (http://www.microrna.org) indicated that the 3'-UTR of HIF-1a contained a predicted binding site for miR-338-3p (Fig. 2A). HIF-1a increased in the NPC cells under hypoxic conditions. Compared with the other NPC cell lines CNE1,

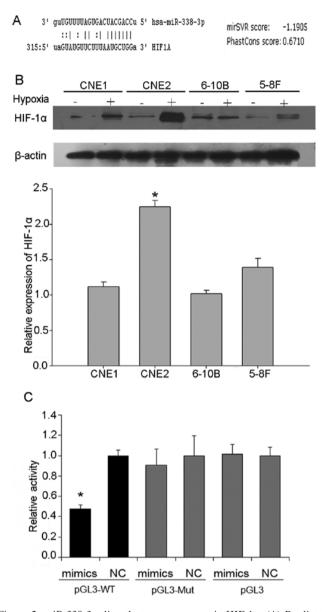


Figure 2. miR-338-3p directly targets oncogenic HIF-1 α . (A) Predicted miR-338-3p target sequences in the HIF-1 α 3'-UTR. (B) Western blot analysis of HIF-1 α levels in NPC cells. β -actin was used as control. Cells were cultured under normal or hypoxic conditions at 24 h. (C) Luciferase reporter assay of cells transfected with the wild-type (wt) or mutant (mut) HIF-1 α 3'-UTR luciferase reporter plasmid of negative control miRNA (NC) or miR-338-3p in NPC cells two days post-transfection.

6-10B and 5-8F, as shown by western blot analysis, HIF-1 α increased most in CNE2 cells (Fig. 2B). Then CNE2 cells were chosen for the following experiments. HIF-1 α has been reported to be an important molecule that drives cancer cell proliferation, migration and invasion including NPC. Using prediction tools, we predicted that the target of miR-338-3p is HIF-1 α . To demonstrate that miR-338-3p directly targets HIF-1 α by interacting with its 3'-UTR, we co-transfected the pGL luciferase reporter plasmid harboring the wild-type or mutant 3'-UTR of HIF-1 α , along with miR-338-3p or NC miRNA (Fig. 2C). Overexpressed miR-338-3p resulted in significant reduction of HIF-1 α 3'-UTR firefly luciferase reporter activity containing wild-type but not mutant binding sites compared to that of NC-miRNA. In summary, these results indicate that HIF-1 α was a direct target gene of miR-338-3p in NPC cells.

To examine the functional significance of miR-338-3p in NPC, we infected the CNE2 cells with miR-338-3p mimics and inhibitor, using miR-67 as control. Using RT-qPCR, we confirmed that the miR-338-3p level had increased >20-fold after transfection with miR-338-3p mimics compared with the expression of cont-miR, while the level was downregulated after the miR-338-3p inhibitor was delivered into CNE2 cells (Fig. 3A). To further investigate the antitumor effect of miR-338-3p under hypoxic conditions and to monitor the effect of miR-338-3p overexpression on HIF-1a expression in target cells expressing low levels of miR-338-3p, NPC cells transfected with miR-338-3p mimics, inhibitors and scrambled control were exposed to hypoxic conditions for 24 h. To confirm that miR-338-3p regulates HIF-1α expression, we assessed HIF-1 α protein levels in CNE2 cells expressing ectopic miR-338-3p, using western blot analysis. The results showed that HIF-1 α levels, under hypoxia, were consistently downregulated by overexpressed miR-338-3p in CNE2 cell lines (Fig. 3B). These results were substantiated by immunofluorescence microscopy (Fig. 3C). Immunofluorescence staining revealed that HIF-1 α expression was inhibited by transfection with miR-338-3p mimics and was promoted by transfection with miR-338-3p inhibitor.

miR-338-3p regulates the metastasis potential of human NPC cell lines in vitro. To examine the functional significance of overexpressed miR-338-3p in NPC, we first investigated the metastasis potential of CNE2 cells with exceptional expression of miR-338-3p. Wound healing tests demonstrated that hypoxic condition increased the metastasis capacity of CNE2 cells. However, the overexpression of miR-338-3p significantly reduced the metastasis of the CNE2 cells under hypoxia (Fig. 4A), the results were confirmed by Transwell analysis (Fig. 4B).

To further explore the molecular mechanism of miR-338-3p on hypoxic signaling in NPC cells, the factors responsible for hypoxia were investigated after the treatment of hypoxic CNE2. Our previous study by western blot analysis disclosed that miR-338-3p downregulated the expression level of HIF-1 α , showed that hypoxia selects for survival the more aggressive tumor cells and induces epithelial to mesenchymal transition (EMT) (29). We suspected that miRNAs would affect EMT under hypoxia since hypoxia is a key factor in the process of EMT. As shown here, exposured CNE2 cells to hypoxia resulted in the transition to a mesenchymal morphology, significant loss of E-cadherin and increased expression of vimentin. To assess the effect of miR-338-3p on hypoxiamediated EMT, miR-338-3p mimics, inhibitor and scrambled control was transfected into CNE2 cells, and the cells were moved to the hypoxic incubator (1% O₂, 5% CO₂, 37°C) for another 24-h incubation. As shown in Fig. 4C, the downstream E-cadherin was upregulated and vimentin was downregulated in miR-338-3p mimic-pretreated CNE2, respectively.

miR-338-3p suppresses NPC cell viability under hypoxic conditions. In order to investigate the effect of miR-338-3p on NPC cell proliferation, CCK8 assays were employed to analyze cell proliferation. As shown in Fig. 5A, overexpression of miR-338-3p in CNE2 cells significantly inhibited cell proliferation at first 24 h under hypoxic conditions (P<0.05), compared

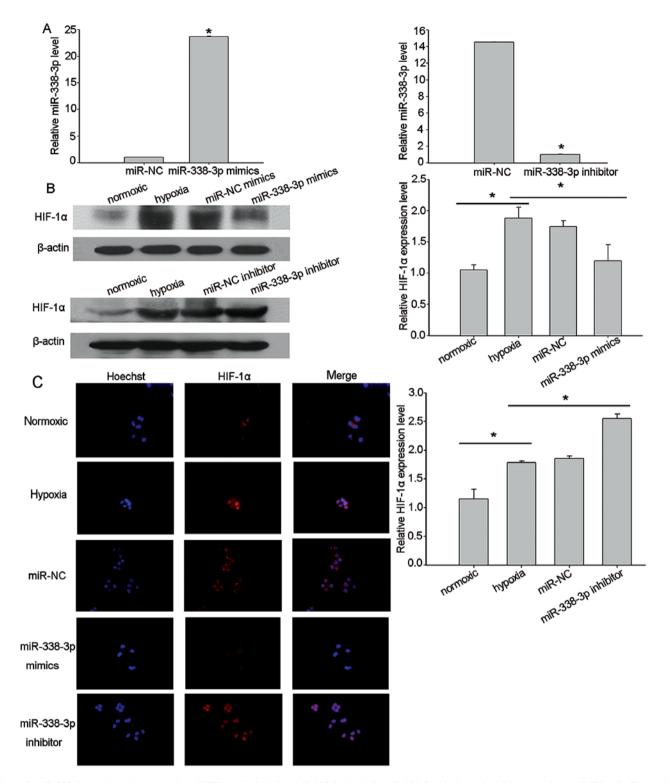


Figure 3. miR-338-3p regulates the expression of HIF-1 α . (A) Relative miR-338-3p levels in miR-338-3p mimics and inhibitor transfected CNE2 cells. Transcript levels were normalized to U6 expression. (B) Western blot analysis of HIF-1 α levels in NC- or miR-338-3p-transfected cells. β -actin was used as loading control. Cells were cultured under hypoxia at two days post-transfection with cells cultured under normoxia as reference, and 24 h later protein levels were analyzed. (C) Immunofluorescence staining analysis of cytoplasmic and nuclear expression of HIF-1 α in NC- or miR-338-3p-transfected CNE2 cells. Cells were incubated under hypoxia for 24 h. Red is HIF-1 α staining. Blue is the nuclear staining by Hochest. Data are shown as mean ± SEM of three independent experiments. "P<0.05.

with the NC transfection. To further demonstrate the effect of miR-338-3p on cell proliferation, miR-338-3p inhibitor was transfected into CNE2 cells. Compared with the mock group, the proliferation of the miR-338-3p inhibitor transfected cells significantly increased (Fig. 5B). Previous evidence showed

that miR-338-3p regulated the phosphorylation of ERK1/2 to mediate tumor cell metastasis and proliferation in gastric cancer (22). Thus, we assumed that miR-338-3p could decrease the expression of the phosphorylation of Erk1/2 by targeting HIF-1 α under hypoxic conditions. Our data showed that the

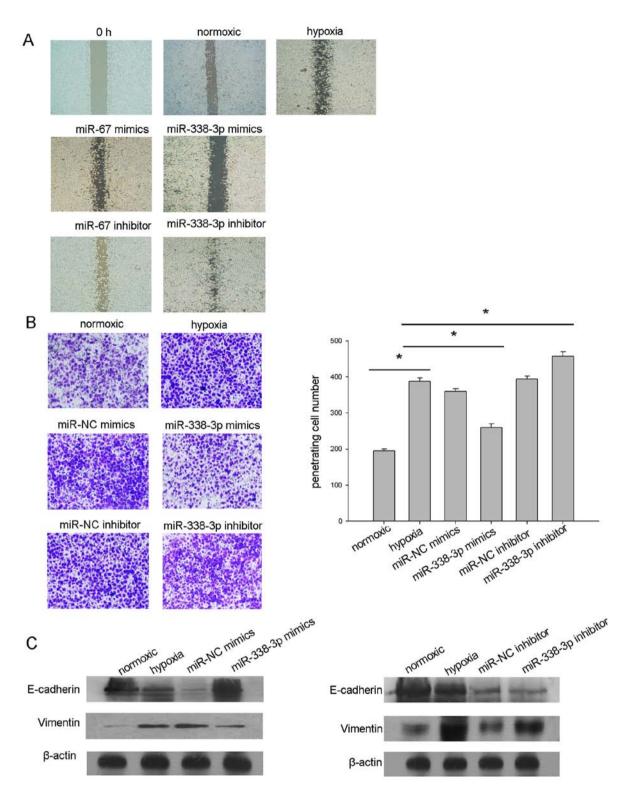


Figure 4. miR-338-3p regulates the metastasis potential of human NPC cell lines *in vitro*. (A) Cells transfected with miR-338-3p mimics showed a slower wound closure rate than the cells transfected with the non-specific control or miR-338-3p inhibitor under hypoxic conditions. (B) Transwell analysis confirmed the results above that the overexpression of miR-338-3p significantly reduced the metastasis capacity of the CNE2 cells under hypoxic conditions. Data are presented as mean number of migrated cells per HPF (five random fields were analyzed), compared with the control. $^{\circ}P<0.05$. (C) Western blot analysis of E-cadherin, and vimentin expression in parental cells, miR-338-3p and miRNA control transfected cells after 24 h under hypoxic conditions compared with cells cultured under normal conditions. A representative blot from three independent experiments is shown. β -actin was used as control.

overexpression of miR-338-3p inhibited the phosphorylation of Erk1/2, but the relative expression level of total Erk1/2 was not significantly altered (Fig. 5C). Taken together, these results suggested anti-cell growth properties of miR-338-3p in CNE2 cells. miR-338-3p sensitizes NPC cells to cisplatin. Because recent studies have reported that destabilization of HIF-1 α can overcome hypoxia-mediated cisplatin resistance in non-small cell lung carcinoma (30), we then tested whether miR-338-3p could sensitize NPC cells to chemotherapy

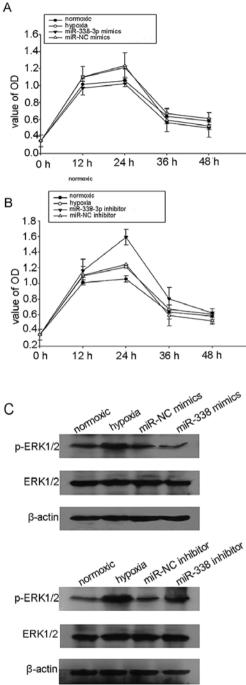
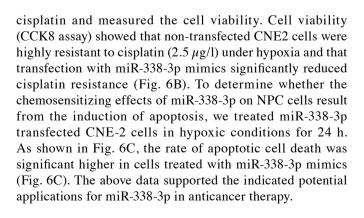


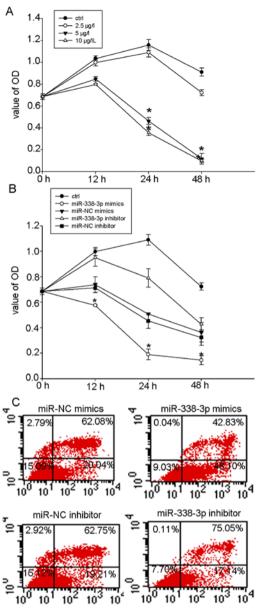
Figure 5. miR-338-3p inhibited CNE2 cell proliferation *in vitro*. (A) The CCK8 assay was performed after transfection of miR-338-3p mimics and miR-NC into the

treatment of cisplatin. CCK8 assay showed that 5 and 10 μ g/l cisplatin inhibited the proliferation of CNE-2 cells in a dose- and time-dependent manner (Fig. 6A). However, the inhibition effect of 2.5 μ g/l cisplatin on hypoxic CNE2 cells was not obvious. To determine the chemo-sensitizing effects of miR-338-3p on hypoxic NPC cells, we chose the treatment dose at 2.5 μ g/l for the following experiments. We treated miR-338-3p transfected cells and NC cells with

CNE2 cells. (B) The cell viability of CNE2 cells treated with miR-338-3p inhibitor compared to normaxic, hypoxia or miR-NC transfected cells. (C) Western blot analysis showed downregulation of p-ERK induced by miR-338-3p in

hypoxic CNE2 cells. β-actin was used as an internal control.





Discussion

The present study focused on the antitumor effect of miR-338-3p in NPC. We demonstrated that the overexpressed miR-338-3p was able to inhibit the proliferation and migration of hypoxic CNE2 cells as well as overcome hypoxia-mediated cisplatin resistance.

NPC is a highly invasive malignancy, and up to 70% of patients with NPC present with a locally advanced stage or with cervical lymph node metastasis at the time of diagnosis (2,25). Those patients have a poor outcome, and drug resistance may hamper the efficacy of anticancer drugs (2). Therefore, clarification of the molecular pathogenesis and mechanism of NPC is crucial for developing effective therapy strategies to improve the outcome of patients with this disease.

Growing evidence indicates that miRNAs hold great promise for novel therapeutic approaches for treating human cancers (28) and that hypoxia regulated miRNAs exhibit induction in response to HIF activation and participate in the development of angiogenesis and tumorigenesis (31). Previous studies have demonstrated that miR-15b, miR-16, miR-20a, and miR-20b are sharply downregulated in CNE cells during hypoxia (31).

Many studies have reported on the tumor suppressive effects of miR-338-3p in the malignant processes of various cancers, such as gastric cancer (19) hepatocellular carcinoma (20) and malignant melanoma (21). However, there is little knowledge on miR-338-3p and its targets in NPC. Our data showed that miR-338-3p expression is markedly downregulated in NPC patient samples and NPC cell lines as compared to immortalized nasopharyngeal epithelial cells.

Predicted targets of miR-338-3p are elements involved in many biological processes, such as cell proliferation, differentiation, metastasis and cell death in various types of cancer (28,32). Similar to previous studies, our data identified HIF-1 α as a key target of miR-338-3p. HIF-1 α is an important hypoxia transcription molecule promoting angiogenesis and metastasis.

Hypoxia is one of the fundamental biological properties that are associated with the development and aggressiveness of a variety of solid tumors (33,34), including NPC (24). Previous study shows that all primary NPC tumors contain hypoxic regions which result in decreased local control and increased distant metastases, as well as resistance to chemotherapy in NPC patients (25). It was recently reported that hypoxia inducible factor (HIF) results in global transcriptional changes in gene expression, playing an important role in promoting tumor progression, angiogenesis and metastasis (34,35). In the present study, we showed evidence that the HIF-1 α levels were consistently downregulated by overexpressed miR-338-3p in CNE2 cell lines under hypoxia conditons. Previous studies have shown that the oxygen-sensitive HIF-1α subunit accumulates and translocates into the nucleus under hypoxia conditions, where it binds to the constitutively expressed HIF-1 β , forming the active HIF-1 heterodimer (36). In the present study, immunofluorescence staining results revealed that HIF-1 α was accumulated into the nuclei in hypoxia cells. However, the overall staining and nuclear accumulation of HIF-1a was reduced after miR-338-3p transfection. Furthermore, overexpressed miR-338-3p inhibited cell viability as well as metastasis ability by directly targeting HIF-1α.

HIF-1 α activates some important oxygen modulated genes critically involved in tumor angiogenesis and metastasis (37). EMT is clearly associated with pathological processes requiring epithelial cell migration and invasion (2,38). Furthermore, HIF-1 α also initiates endothelial to mesenchymal transition (39). Our data showed that the over-expressed miR-338-3p inhibited the hypoxia-induced EMT.

HIF-1 α has attracted considerable interest as a potential target in cancer therapy (37). Notably, growing evidence suggests that higher HIF-1 α levels in tumors are related to radioresistance and poor clinical outcome in different tumor entities (40). Some evidence shows that HIF-1 α -deficient cells are more sensitive to radiotherapy and chemotherapy compared with normal cells (41), however, the mechanism needs further investigation. Therefore, the specific inhibition of HIF may enhance cancer radiosensitivity in clinical settings. Our results showed that miR-338-3p downregulated the level of HIF-1 α in CNE2 under hypoxic conditions and could overcome hypoxia-mediated cisplatin resistance.

Previous studies have shown that other cell regulatory elements such as cyclin D, and smoothened, are also targets of miR-338-3p that are aberrantly expressed due to the downregulation of miR-338-3p in HCC (28). Undoubtedly, regulation of these other targets may contribute to the inhibitory effects of miR-338-3p on NPC. However, considering our observation that HIF-1 α overexpression rescued the cell from the anti-NPC activity of miR-338-3p, it is likely that regulation of HIF-1 α by miR-338-3p is a key antitumor aspect in NPC. Our further studies will focus on other targets of miR-338-3p and their specific roles under both normoxic and hypoxic conditions.

In conventional clinical therapy of NPC, radiotherapy is the primary therapeutic approach, while using radiotherapy combined with chemotherapy is recommended for the treatment of advanced carcinoma (42). Treatment failure rates remain high and the 5-year survival rate is extremely low (42). One reason for the treatment failure may be attributed to the drug resistance and distant metastasis (42). Evidence shows that destabilization of HIF-1α can overcome hypoxia-mediated cisplatin resistance in non-small cell lung carcinoma (30). We want to determine whether miR-338-3p potentiates sensitivity of NPC cells to cisplatin, which is clinically used as adjuvant therapy for NPC in order to induce tumor cell death (42). Our results showed that under hypoxic conditions, CNE2 cells are highly resistant to cisplatin in vitro. Moreover, cells pretransfected with miR-338-3p can overcome hypoxia-mediated cisplatin resistance. These results enriched the function of miR-338-3p in addition to its role as a tumor suppressor.

In conclusion, we found that miR-338-3p was downregulated in NPC clinical samples and cell lines. HIF-1 α was verified as a direct target of miR-338-3p, and involved in NPC cell growth and invasion. Furthermore, our data suggest that miR-338-3p and/ or its target gene HIF-1 α could represent important therapeutic targets in NPC. We explored its effects on cell growth, invasive and chemotherapy resistance. The miR-338-3p/HIF-1 α pathway may help to understand the mechanisms of NPC progression and would provide a novel therapeutic strategy for NPC.

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