

Overexpressing miR-335 inhibits DU145 cell proliferation by targeting early growth response 3 in prostate cancer

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Abstract. MicroRNA-335 (miR-335) was reported to suppress cell proliferation in prostate cancer (PC), a common malignancy in males. The expression of early growth response 3 (EGR3) was determined to be elevated in human PC tissues; however, the possible effects and underlying mechanism of miR-335 on PC remains unknown. In the present study, miR-335 mimics and miR-335 inhibitors were respectively transfected into DU145 cells. Stable silencing of EGR3 was observed in DU145 cells following transfection with small interfering RNA. We also used Cell Counting Kit-8 and *in vitro* angiogenesis assays to determine the viability and revascularization potential of DU145 cells. The expression levels of EGR and caspase-3 activity were analyzed by immunohistochemistry and immunocytochemistry, respectively. We predicted the target of miR-335 by bioinformatics analysis and a dual-luciferase reporter gene assay. Western blot and quantitative real-time polymerase chain reaction analyses were performed to determine the protein and mRNA expression of molecules. miR-335 expression was downregulated in PC tissues and cell lines. Overexpression of miR-335 significantly reduced the viability and the formation of regenerative tubes of DU145 cells, and inhibited the expression of inflammatory factors. EGR3 was proposed as a possible target of miR-335, and was negatively regulated by miR-335. Silencing EGR3 suppressed the viability and angiogenesis of DU145 cells, and reduced the activity of caspase-3 and inflammatory factor expression. miR-335 inhibition along with EGR3 silencing

EGR3 inhibited the cell proliferation. Furthermore, miR-335 inhibited the formation of a PC solid tumor xenograft *in vivo*. Thus, miR-335 may exert an antitumor effect on DU145 cells by regulating the expression of EGR3. The findings of the present study may provide insight into a novel therapeutic strategy for the treatment of prostatic carcinoma.

Introduction

Prostate cancer (PC) is a common malignant cancer that often occurs in aged males and is the second leading cause of cancer-associated mortality (1,2). As of its notable heterogeneity, PC is difficult to diagnose according to current diagnostic and prognostic indexes; for example, an underestimated Gleason score and elevated levels of prostate specific antigen in the serum of patients with prostatitis and benign prostatic hyperplasia (3,4). Given the molecular heterogeneity of PC (5), its diagnosis, treatment and prognosis may be improved by further investigations into the translation of several molecular markers in specific stages of disease (6). Recently, the roles of micro (miRNAs/miRs) in PC have been widely studied, and aberrant miRNAs expression has been regarded as crucial biomarkers for the diagnosis, grading and prognosis of PC (7,8). miRNAs can regulate carcinogenesis of prostate tissue by interacting with their target genes (9,10). Therefore, it is necessary to explore the regulatory mechanism of miRNA, which may provide insight into novel therapeutic strategies for the treatment of PC.

As small, non-coding and single-stranded RNAs, the endogenously expressed miRNAs can reversely mediate the expression of target genes at the post-transcriptional level, and inhibit translation or induce mRNA degradation to activate various pathways mainly by binding to the 3'-untranslated region (3'-UTR) (11-13). It has been demonstrated that miR-335 is an important miRNA associated with numerous types of cancer (14). For instance, the overexpression of miR-335 has been reported to suppress the metastatic invasion of breast cancer (BCa) cells (15). It was revealed that reduced miR-335 expression in BCa tissues is associated with the clinicopathological characteristics of BCa (16). Additionally, miR-335 has been demonstrated to be related

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to the chemoresistance of ovarian cancer cells (17). On the contrary, in astrocytoma cells, miR-335 can promote tumor growth and invasion by targeting a potential tumor inhibitor, while miR-335 inhibition could notably induce growth arrest and apoptosis *in vitro* and *in vivo* (18). Furthermore, miR-335 expression is dysregulated in several cancers, including hepatocellular carcinoma, meningioma, PC and colorectal cancer (19-22). Additionally, miR-335 was reported to possess diverse targets in the same cancer (23,24). Therefore, exploring the target genes of miR-335 may reveal novel therapeutic targets.

Early growth response (EGR) transcription factors can be induced in a variety of cells to respond to stimuli, including stress, hypoxia, injury, cytokines and growth factors (25). In the EGR family of proteins, EGR1 has been reported to be associated with inflammation, ischemic injury and atherosclerosis (26), in addition to its antitumor effects by inducing apoptosis in cancer (27-29). However, to the best of our knowledge, the number of the studies on EGR3 is limited. A recent study has suggested that high expression of EGR3 in A549 cells could inhibit cell growth and is associated with improved prognosis in lung adenocarcinoma (30). Conversely, elevated EGR3 expression was determined to be associated with poor prognosis in PC (31). Furthermore, EGR3 can regulate the levels of inflammatory cytokines, including interleukin-6 (IL-6) and IL-8 in PC cells, and these cytokines can exert critical effects on the development of PC (32). Considering the roles of miR-335 and EGR3 in the PC, we aimed to examine whether miR-335 exhibits anticancer effects by regulating its potential target gene EGR3.

Materials and methods

Prostate tissue specimens. The present study was approved by the Ethics Committee of The Second Affiliated Hospital of Xi'an Jiaotong University and informed consent was obtained from each patient. A total of 53 male patients in our hospital from December 2017 to January 2018 were enrolled in this study: 36 patients with prostate cancer (mean age, 68.2; age range, 55-82 years) and 18 patients with metastatic prostate cancer (mean age, 72.4; age range, 58-86 years). Patients with prostate cancer included in the present study received no preoperative medication and had no history of surgical castration or radiotherapy. Fifty-three PC tissues and matched normal tissues from patients were frozen for reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analysis and immunohistochemistry (IHC).

Cell culture. Human umbilical vein endothelial cells (HUVECs) were purchased from Promocell (Jiangyin, China) and cultured in endothelial cell growth medium (Promocell). DU145 and PC-3M cells (human prostate carcinoma cell line) were obtained from the American Type Culture Collection (Manassas, VA, USA) and maintained in RPMI-1640 medium (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) containing 10% fetal bovine serum (Thermo Fisher Scientific, Inc., Waltham, MA, USA), 2 mM L-glutamine (Gibco; Thermo Fisher Scientific, Inc.) and 1% antibiotic mixture (penicillin and streptomycin; Gibco;

Thermo Fisher Scientific, Inc.) at 37°C in a humidified incubator with 5% CO₂.

Cell transfection. The prostate cancer DU145 Cells (2x10⁵) were transfected with miR-335 mimics (5'-UCAAGAGCAA UAACGAAAAAUGU-3'), miR-335 inhibitors (5'-ACAUUU UUCGUUAUUGCUCUUGA-3') and corresponding controls (Thermo Fisher Scientific, Inc.) using Lipofectamine® 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) at a final concentration of 50 nM according to the manufacturer's protocols. The PC-3M cells were only transfected with miR-335 mimics and corresponding controls. To knockdown EGR3, DU145 cells were transfected with 30 nM small interfering RNA (siRNA) against EGR3 (siEGR3; #115514; Ambion, Austin, TX, USA) and negative controls (siNC; #4642; Ambion) using siPORT NeoFX (Ambion) according to the manufacturer's protocol and incubated for 24 h. The sequence of siRNAs for EGR3 was 5'-CCAACACAACAGAUAGAAUtt-3'. Puromycin was used to select transfected cells. In addition, siEGR3 and miRNAs were co-transfected into DU145 cells at a concentration of 50 nM/well. At 48 h after transfection, cells were harvested for further analyses. Non-transfected cells were served as another control.

Cell viability assay. Transfected cells (1x10⁴) were cultivated for 24, 48 and 72 h, respectively. Subsequently, 10 µl Cell Counting Kit-8 solution (Beyotime Institute of Biotechnology, Shanghai, China) was added into the cells at each time point. After the cells were incubated for 2 h at 37°C, the optical density at 450 nm was measured with a FlexStation3 microplate reader (Molecular Devices, LLC, Sunnyvale, CA, USA).

Apoptosis analysis. Apoptosis was detected according to the protocols of the Annexin V-fluorescein isothiocyanate/propidium iodide apoptosis detection kit (Nanjing KeyGen Biotech, Co., Ltd., Nanjing, China). The results were analyzed with a FACSCalibur flow cytometer (BD Biosciences, San Jose, CA, USA).

In vitro angiogenesis. HUVECs (3x10⁵) were seeded with Matrigel (BD Biosciences) in a 96-well plate in the respective presence of supernatants comprising non-transfected DU145 cells, DU145 cells transfected with miR-NC or miR-335 mimics, DU145 cells transfected with miR-335 inhibitors, or DU145 cells transfected with siNC or siEGR3, and then incubated at 37°C for 15 h. Then, the cells were fixed with 3.7% formaldehyde for 15 min at room temperature. The images of capillary-like structures from six replicates of each group were captured using an IX81 Olympus microscope (Olympus Corporation, Tokyo, Japan; magnification, x100). The overall length and cell junctions in each frame from five randomly selected fields were calculated using ImageJ v1.48 software (National Institutes of Health, Bethesda, MD, USA). Data were normalized to the controls.

Immunocytochemistry. For caspase-3 expression, the cells were immobilized in 4% paraformaldehyde for 24-48 h at room temperature and permeabilized in Tris-buffered saline with 0.1% Triton X-100 (TBST; pH 7.4; Sigma-Aldrich; Merck KGaA). The cells were treated with 0.1% H₂O₂

Table I. Sequences of primers used for reverse transcription-quantitative polymerase chain reaction.

Gene	Forward (5'-3')	Reverse (5'-3')
miR-335	GTCGTATCCAGTGCAGGGTCCG	GTGCAGGGTCCGACCT
EGR3	TTCGCTTTTCGACTCTCC	CTCCGAGTAGAGATCGC
IL-6	AAATTCGGTACATCCTCGACGGCA	AGTGCCTCTTTGCTGCTTTTCACAC
IL-8	AGGACAAGAGCCAGGAAGAAACCA	AGAGCTGCAGAAATCAGGAAGGCT
IL-1 β	AACAGGCTGCTCTGGGATTCTCTT	AACAGGCTGCTCTGGGATTCTCTT
U6	CTCGCTTCGGCAGCACA	AACGCTTACGAATTTGCGT
β -actin	GTGACGTTGACATCCG	GAGCGTTTGTGTACCT

miR, microRNA; EGR3, early growth response 3; IL, interleukin.

for 30 min at room temperature to inhibit endogenous peroxidase, followed by incubation with 5% normal goat serum (Vector Laboratories, Inc., Burlingame, CA, USA) for 40 min at room temperature to block non-specific binding sites. The cells were then incubated with anti-caspase-3 (1:50; cat. no. 9661, Cell Signaling Technology, Inc., Danvers, MA, USA) overnight at 4°C and then with a biotinylated goat anti-rabbit IgG (1:200) as a secondary antibody (BP-9100, Vector Laboratories, Inc.) for 1 h at room temperature. Subsequently, the cells were washed in TBST and incubated with avidin-biotin-peroxidase complex (1:1:100; Strept ABC complex/HRP; Dako; Agilent Technologies, Inc., Santa Clara, CA, USA) for 45 min in dark at room temperature. The results was visualized with 3'3'-diaminobenzidine (DAB; Vector Laboratories, Inc.) for 10 min at room temperature and stained with hematoxylin (Sigma-Aldrich; Merck KGaA) for 30 sec at room temperature.

Target prediction. To predict the potential targets of miR-335, TargetScan 7.2 (<http://www.targetscan.org>) was utilized and the candidate targets were identified based on the phenotype of miR-335.

Luciferase assay. DU145 cells were transfected with a firefly luciferase reporter pGL3 vector (Promega Corporation, Madison, WI, USA) containing the wild-type or mutant EGR3 3'-UTR, the control pRL-TK vector containing *Renilla* luciferase (Promega) with miR-335 mimics and miR-335 inhibitors using Lipofectamine 2000. The cells were maintained for 24 h and the activities of firefly and *Renilla* luciferase were determined by Dual-Luciferase Reporter Assay System (Promega Corporation).

RNA extraction, cDNA synthesis and RT-qPCR. Total RNA of prostate tissues and DU145 cells was extracted using TRIzol (Invitrogen; Thermo Fisher Scientific, Inc.). In brief, TRIzol reagent and 200 ml chloroform were added to samples and mixed for 5 min, followed by centrifugation (12,000 x g for 15 min at 4°C) to recover the supernatant. Then, the supernatant was incubated with an equal volume of isopropyl alcohol for 10 min at room temperature and centrifuged at 12,000 x g for 15 min at 4°C. After dislodging the supernatant, 75% ethanol was added to for precipitation and the RNA was eluted with nuclease-free water. For the RT, the concentration and purity

were detected by a NanoDrop 2000 spectrophotometer (NanoDrop Technologies; Thermo Fisher Scientific, Inc.). cDNA was obtained from 1 μ g RNA at 25°C for 10 min, 37°C for 120 min and 85°C for 5 min using the High Capacity cDNA Reverse Transcription Kit (Applied Biosystems; Thermo Fisher Scientific, Inc.). The qPCR reaction system was as follows: 2.5 μ l dNTPs (2.5 mM); 2.5 μ l 10X PCR buffer; 1.5 μ l MgCl₂ solution; 1 unit Taq polymerase; 0.25X SYBR Green I (Sigma-Aldrich; Merck KGaA); 1 μ l primers (10 μ M each); 1 μ l cDNA; water (to a total volume of 25 μ l). The qPCR reactions were conducted with a LightCycler system (Roche Applied Science, Indianapolis, IN, USA) using the following parameters: 95°C for 10 min, 40 cycles at 95°C for 10 sec, at 60°C for 20 sec and at 72°C for 30 sec. The specific primers were listed in Table I. U6 and β -actin served as the internal reference genes. Expression was calculated using the 2^{- $\Delta\Delta$ C_q} method (33).

Western blotting. The cells were lysed in lysis buffer (Beyotime Institute of Biotechnology, Shanghai, China) and centrifuged at 10,000 x g for 10 min at 4°C, and placed on ice for 30 min to obtain the supernatant. The protein concentration was determined using a BCA kit (Beyotime Institute of Biotechnology). After being separated by 10% SDS-PAGE (40 μ g protein was used), the proteins were transferred onto nitrocellulose membranes (EMD Millipore, Bedford, MA, USA) and blocked for 1 h. The membranes were incubated overnight at 4°C with anti-EGR3 (1:500, sc-390967, Santa Cruz Biotechnology, Inc., Dallas, TX, USA), anti-IL-6 (1:1,000, ab9324, Abcam, Cambridge, USA), anti-IL-8 (1:1,000, ab18672, Abcam), anti-IL-1 β (1:1,000, ab156791, Abcam) and anti- β -actin (1:1,000, sc-517582, Santa Cruz Biotechnology, Inc.) and incubated with secondary antibodies (1:1,000, HRP-labeled goat anti-mouse IgG, A0216, Beyotime Institute of Biotechnology) for 2 h at room temperature. Proteins were visualized with an ECL kit (GE Healthcare, Chicago, IL, USA). The Gray value was detected by ImageJ 1.48 software.

Xenograft experiment of prostate cancer. Female severe combined immunodeficient mice (20-22 g; n=6; 6 weeks old) were purchased from the Animal Experimental Center of Zhejiang Academy of Medical Sciences. Mice were housed in

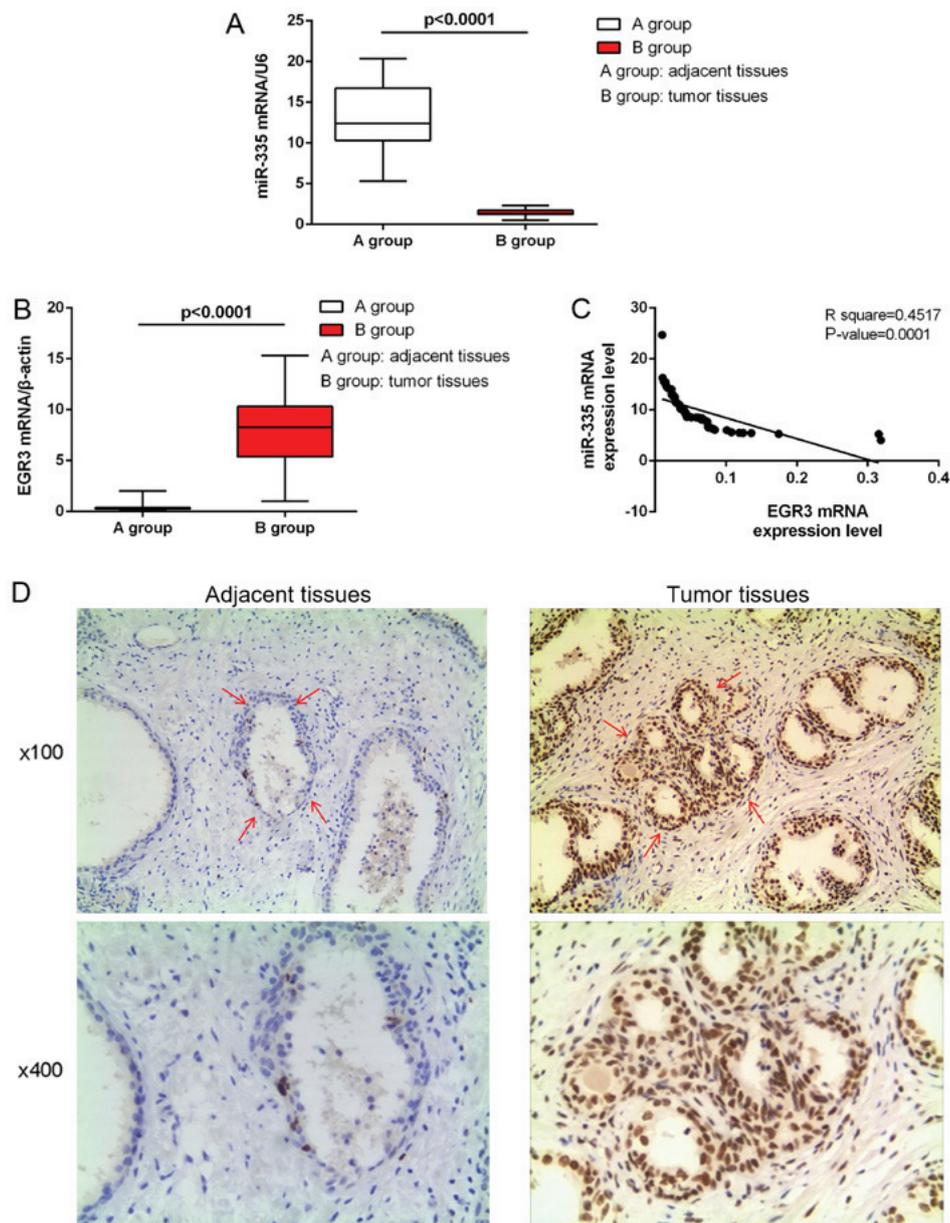


Figure 1. Expression of miR-335 and EGR3 in tissues of patients with PC. The expression of (A) miR-335 and (B) EGR3 in tumor tissues and match adjacent tissues of 53 patients were detected by reverse transcription-quantitative polymerase chain reaction. (C) Correlation between miR-335 expression and EGR3 mRNA expression. (D) EGR3 expression in prostate cancer tissues as determined by immunohistochemistry. Magnifications, x100 and 400. Comparison of EGR3 expression between tumor prostate tissue and adjacent normal prostate tissue are indicated by red arrows. EGR3, early growth response 3; miR, microRNA.

cages at room temperature ($22\pm 3^{\circ}\text{C}$) with constant humidity ($50\pm 10\%$) with a free access to food and water under a light/dark cycle (12 h). The animal experiments were approved by The Second Affiliated Hospital of Xi'an Jiaotong University Animal Ethics Committee (approval no. JT201805032M) and according to the Guidelines for the Care and Use of Laboratory Animals (34).

PC-3M and DU145 cells were transfected with miR-335 mimic and miR-NC as aforementioned. PC model was induced by a hypodermic injection of 5×10^6 non-transfected or transfected cells into lateral abdominal wall of mice. Tumor volume was measured every 6 days for 42 days. The mice were sacrificed 42 days after the cell transplantation and weighing.

IHC. IHC was performed to detect the EGR3 expression in 53 prostate cancer tissues and matched adjacent tissues and in the PC xenograft tissues of mice as previously described (35). In brief, tissue sections ($4\ \mu\text{m}$) were dewaxed in 4% paraformaldehyde for 10 min at room temperature and rehydrated with gradient ethanol (Sigma-Aldrich; Merck KGaA), followed by heat-induced antigen retrieval in 10 mM sodium citrate (pH 6.0) for 20 min. Then the sections were blocked in 10% normal goat serum (Vector Laboratories, Inc.) at room temperature for 10 min. Thereafter, the sections were incubated in 3% H_2O_2 at room temperature for 30 min to inactivate endogenous peroxidase. The sections were subsequently incubated with the anti-EGR3 (1:50, sc-390967, Santa Cruz Technology, Inc.) overnight at 4°C and hybridized

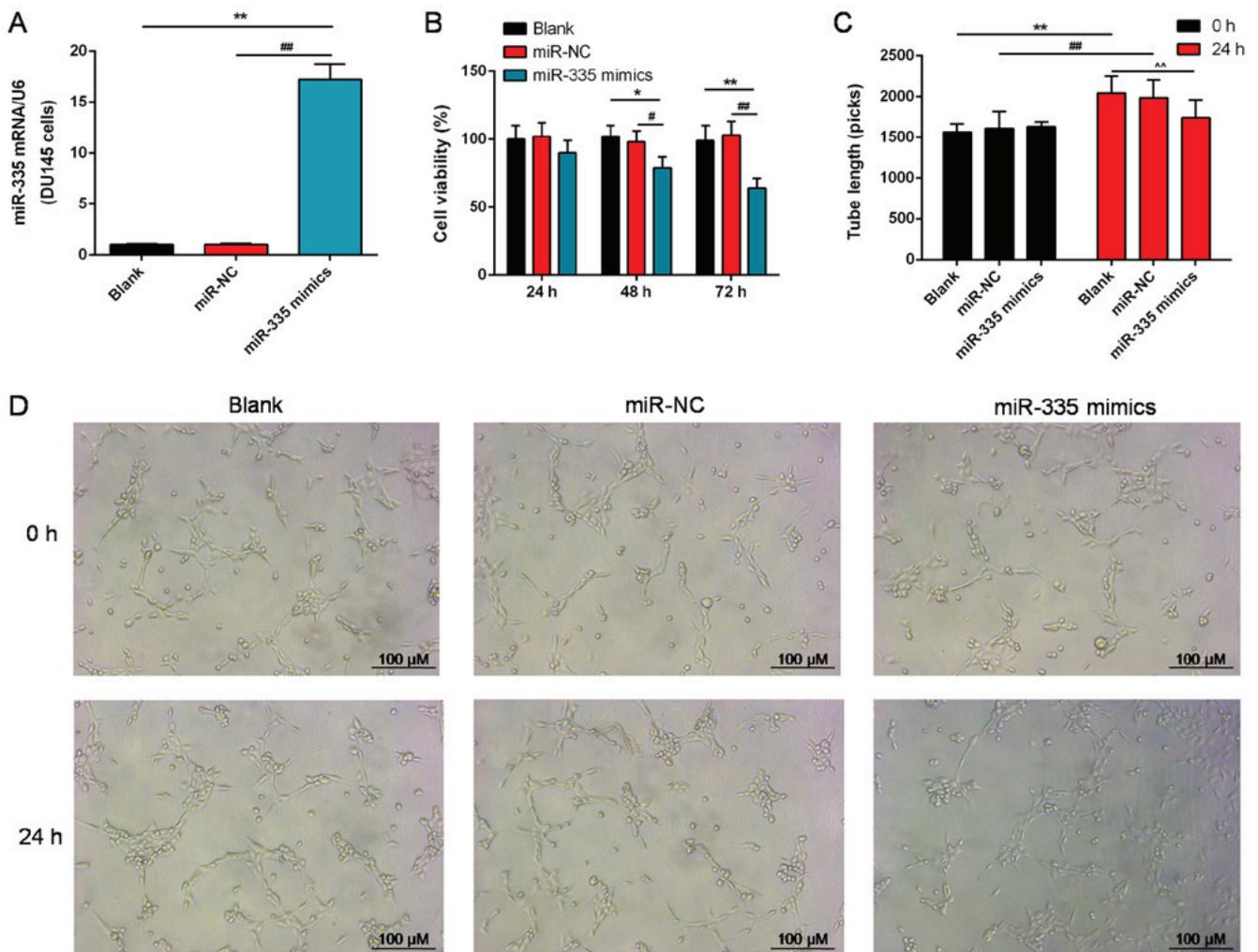


Figure 2. Overexpression of miR-335 inhibits viability and angiogenesis. (A) Elevated miR-335 expression indicated the successful transfection of miR-335 mimics. (B) Overexpression of miR-335 inhibited cell viability in a time-dependent manner. (C and D) The lengths and numbers of tubes were reduced following the overexpression of miR-335 in human umbilical vein endothelial cells. Scale bar=100 μm. *P<0.05, **P<0.01, #P<0.05, ##P<0.01, ^^P<0.01, n=3. Scale bar=100 μm. miR, microRNA; NC, negative control.

with a biotinylated horse anti-mouse IgG secondary antibody (1:200, BP-2000, Vector Laboratories, Inc.) for 1 h at room temperature. The cells were washed in TBST and incubated with avidin-biotin-peroxidase complex (1:1:100; Strept ABC complex/HRP; Dako; Agilent Technologies, Inc.) for 45 min in dark at room temperature. DAB was then added to incubate for 10 min at room temperature and sections were stained with hematoxylin for 30 sec at room temperature. The images were diagnosed under an Olympus microscope (Olympus Corporation; magnification, x400). Image-Pro Plus version 6.0 software (Media Cybernetics, Inc.) was used to evaluate the integrated optical density value of the tissue areas from five randomly selected fields.

Statistical analysis. All data were obtained from independent experiments and were presented as the mean ± standard deviation. P<0.05 was considered to indicate a statistically significant difference. A Student's t-test and one-way analysis of variance with a Dunnett's post-test were conducted using SPSS 17.0 software (SPSS, Inc. Chicago, IL, USA). Comparisons between tumor tissues and matched adjacent tissues were analyzed with

an independent samples t-test. Correlations between miR-335 expression and EGR3 mRNA were analyzed by Spearman's correlation analysis.

Results

Expression of miR-335 and EGR3 in tissues of patients with PC. Among the 53 pairs of tissues, the expression levels of miR-335 in the prostate tumor tissues were significantly reduced than in normal tissues (P<0.0001; Fig. 1A). On the contrary, EGR3 expression in tumor tissues was significantly elevated than in normal tissues (P<0.0001; Fig. 1B). The expression of miR-335 was negatively correlated with that of EGR3 in patients with PC (P=0.0001; Fig. 1C). Furthermore, upregulated EGR3 expression was observed in cancer tissues (Fig. 1D).

Effects of miR-335 on cell viability and angiogenesis. The notably reduced expression of miR-335 in DU145 cells was significantly enhanced following transfecting cells with miR-335 mimics compared with the controls (P<0.01; Fig. 2A). Additionally, miR-335 mimics significantly inhibited

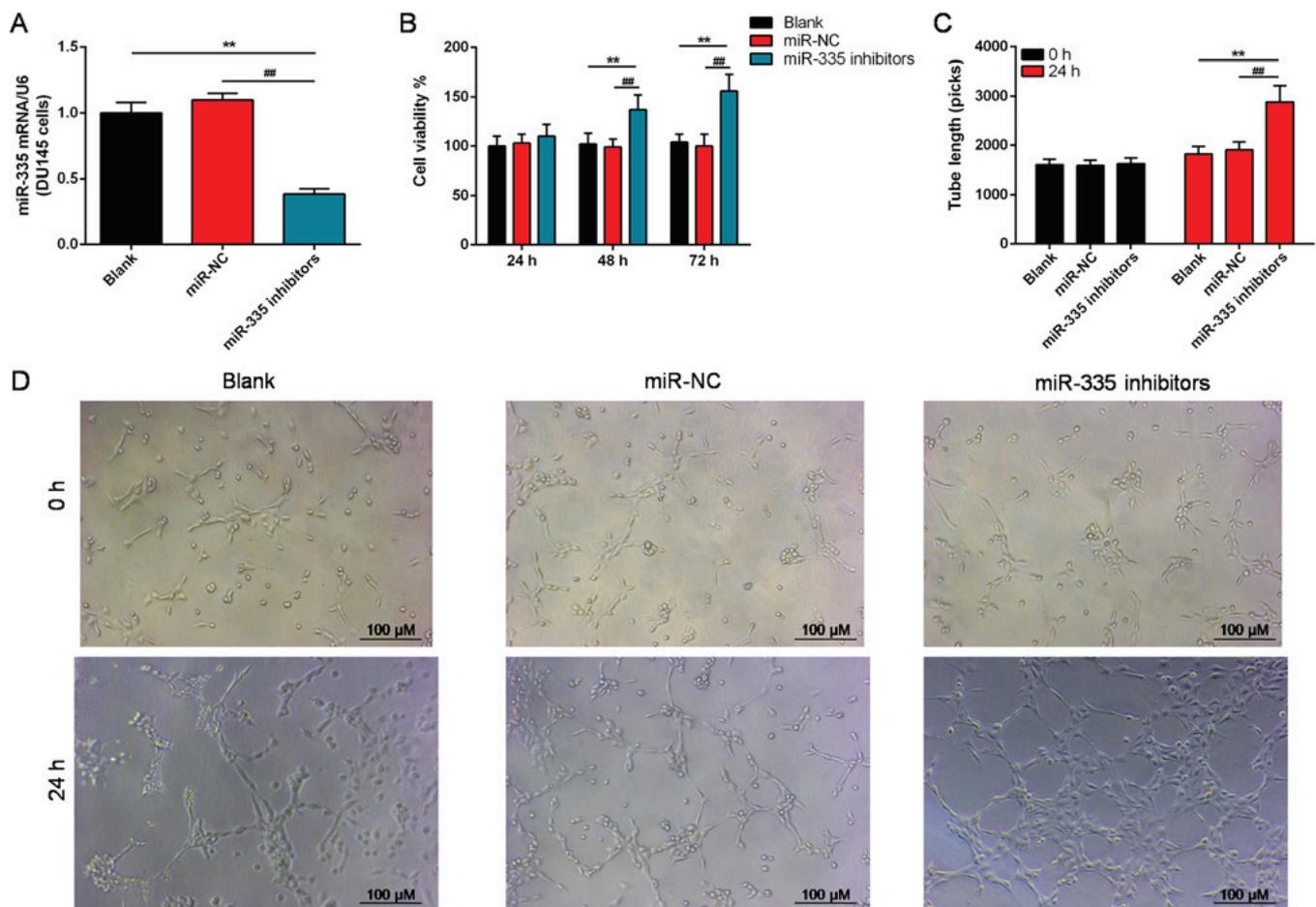


Figure 3. miR-335 knockdown promotes cell viability and angiogenesis. (A) miR-335 expression was significantly decreased following transfection with miR-335 inhibitors. (B) Silencing miR-335 in DU145 cells enhanced cell viability in a time-dependent manner. (C and D) miR-335 silencing increased the length and number of tubes in human umbilical vein endothelial cells. ** $P < 0.01$, ## $P < 0.01$, $n = 3$. miR, microRNA; NC, negative control.

cell viability after 48 h compared with the controls ($P < 0.01$; Fig. 2B). After 24 h of culture, the length of neoformative tubes of HUVECs was significantly suppressed by miR-335 mimics, compared with the blank and miR-NC groups ($P < 0.01$; Fig. 2C). In addition, the number of tubes formed in the miR-335 mimics group was markedly lower than in the blank and miR-NC groups (Fig. 2D).

To further explore the role of miR-335 in the PC, we also successfully silenced the miR-335 expression in DU145 cells ($P < 0.01$; Fig. 3A). After being cultured for > 48 h, the viability of DU145 cells was significantly enhanced in the miR-335 inhibitors group compared with the controls ($P < 0.01$; Fig. 3B). Additionally, the length of the tubes were significantly increased compared with the controls ($P < 0.01$; Fig. 3C and D) in miR-335 inhibitors group.

Role of miR-335 in the apoptosis of DU145 cells. miRNAs serve important roles in apoptosis; thus, the apoptosis of DU145 cells was analyzed in our study. A significant increase in the number of apoptotic cells was observed in the miR-335 mimics group, compared with the blank and miR-NC groups ($P < 0.01$; Fig. 4A). Whereas, the number of apoptotic cells in miR-335 inhibitors group exhibited no significant difference compared with the blank and miR-NC groups (Fig. 4B). The

number of apoptotic DU145 cells was significantly increased in the miR-335 mimics group, compared with that in cells transfected with miR-335 inhibitors ($P < 0.01$; Fig. 4C).

Furthermore, the number of caspase-3-positive cells markedly increased following miR-335 upregulation; a significant increase in activity of caspase-3 was observed in the miR-335 mimics group compared with the controls ($P < 0.01$; Fig. 5A and B). However, the proportion of caspase-3 positive cells in the miR-335 inhibitors group was notably less than that in blank group or miR-NC group; caspase-3 activity was significantly reduced following miR-335 knockdown ($P < 0.01$; Fig. 5C and D).

Regulation of miR-335 in the expression of inflammatory cytokines. The protein expression levels of several pro-inflammatory factors, including IL-6, IL-8 and IL-1 β were significantly suppressed in the miR-335 mimics group compared with blank and miR-NC groups ($P < 0.01$; Fig. 6A). Similarly, the mRNA expression levels of IL-6, IL-8 and IL-1 β in the miR-335 mimics group were significantly down-regulated than in blank and miR-NC groups ($P < 0.01$; Fig. 6B). The protein expression levels of IL-6, IL-8 and IL-1 β were significantly enhanced following transfection with miR-335 inhibitors compared with the control groups ($P < 0.01$; Fig. 6C);

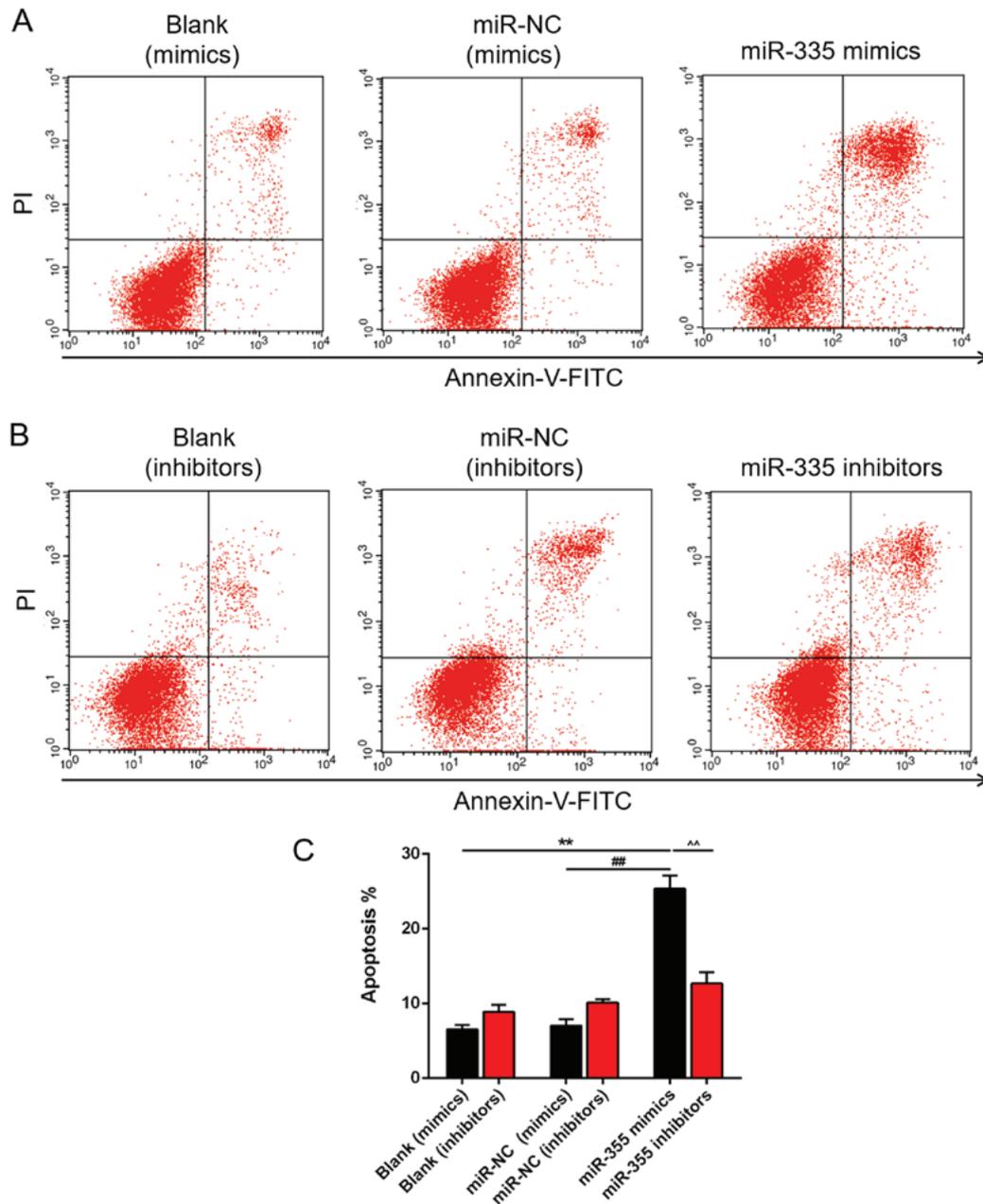


Figure 4. Effects of miR-335 on the apoptosis of DU145 cells. (A) The apoptotic cells in the blank, miR-NC and miR-335-mimics groups were determined by flow cytometry. (B) The apoptotic cells in the blank, miR-NC and miR-335-inhibitors groups were determined by flow cytometry. (C) Overexpression of miR-335 significantly promoted apoptosis, while miR-335 knockdown revealed no significant changes in apoptosis. **P<0.01, ##P<0.01, ^^P<0.01, n=3. miR, microRNA; NC, negative control.

a similar trend in mRNA expression was also for IL-6, IL-8 and IL-1 β (P<0.01; Fig. 6D).

EGR3 is a potential target of miR-335. Based on the bioinformatic prediction analysis, a miR-335-binding site was mapped to the 3'-UTR of EGR3 (Fig. 7A). The luciferase activity in cells transfected with a wild-type EGR3-3' UTR vector containing miR-335 mimics was significantly reduced compared with the mutated 3'-UTR. However, the luciferase activity was significantly enhanced in DU145 cells transfected with wild-type EGR3-3'-UTR vector containing miR-335 inhibitors (Fig. 7B). EGR3 expression in miR-335 mimics group was significantly suppressed at the protein and

mRNA levels compared with the controls (P<0.01; Fig. 7C and D). Conversely, the protein and mRNA expression levels of EGR3 were significantly increased in the cells transfected with miR-335 inhibitors compared with the controls (P<0.01; Fig. 7E and F).

Roles of EGR3 in the PC cells. To understand the effects of EGR3 on prostate tumors, we detected the angiogenesis and apoptosis of the DU145 cells. EGR3 expression was significantly knocked down at the transcription and translation levels in cells following transfection with siEGR3 (P<0.01; Fig. 8A and B). Cell viability was significantly decreased following EGR3 knockdown in a time-dependent manner

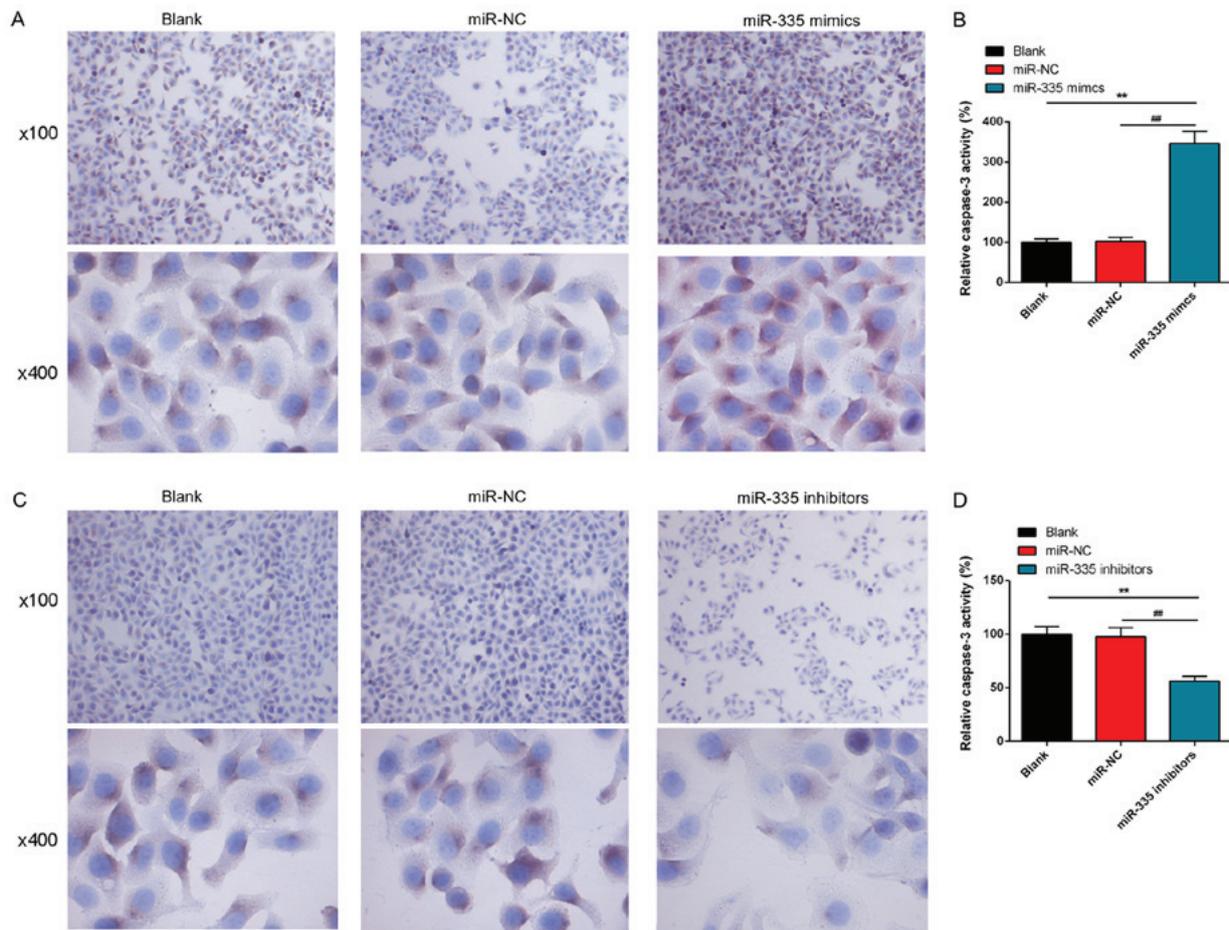


Figure 5. Effects of miR-335 on caspase-3 activity in DU145 cells. (A) Caspase-3-positive cells in the blank, miR-NC and miR-335-mimics groups were analyzed by immunocytochemistry. (B) The caspase-3 activity was significantly increased following miR-335 overexpression. (C) The caspase-3-positive cells in the blank, miR-NC and miR-335-inhibitors groups. (D) The caspase-3 activity was significantly reduced following downregulation of miR-335. **P<0.01, ##P<0.01, n=3. miR, microRNA; NC, negative control.

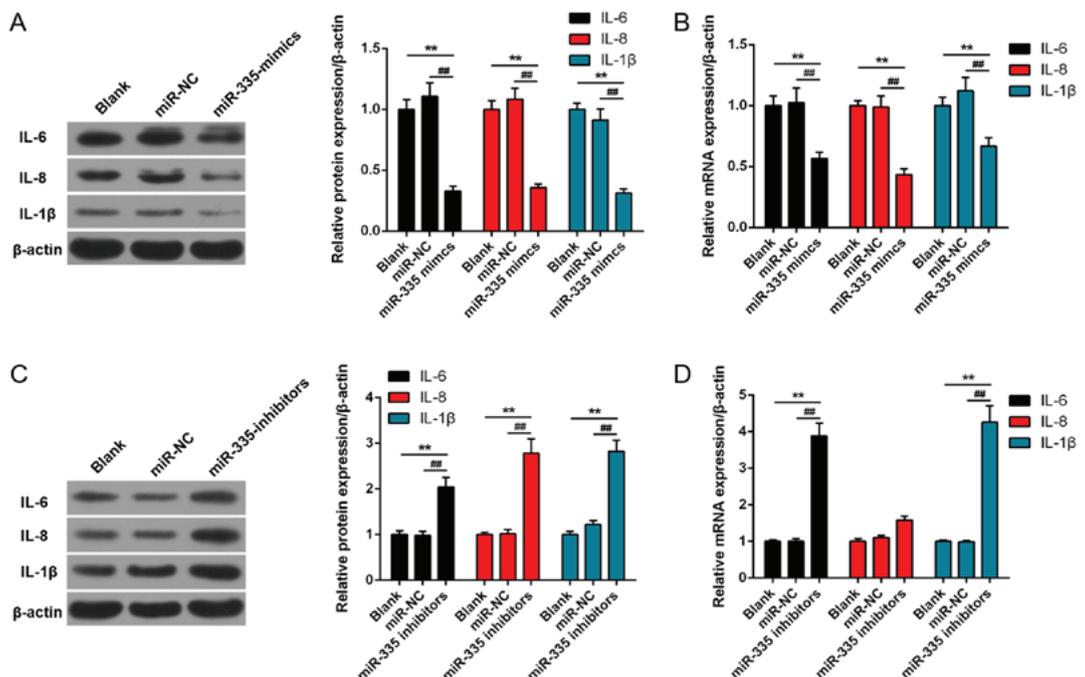


Figure 6. Effects of miR-335 on the expression of inflammatory cytokines in DU145 cells. (A and B) The protein and mRNA levels of IL-6, IL-8 and IL-1β were significantly reduced following miR-335 overexpression. (C and D) The protein and mRNA expression levels of IL-6, IL-8 and IL-1β were significantly enhanced following transfection with miR-335 inhibitors. **P<0.01, ##P<0.01, n=3. IL, interleukin; miR, microRNA; NC, negative control.

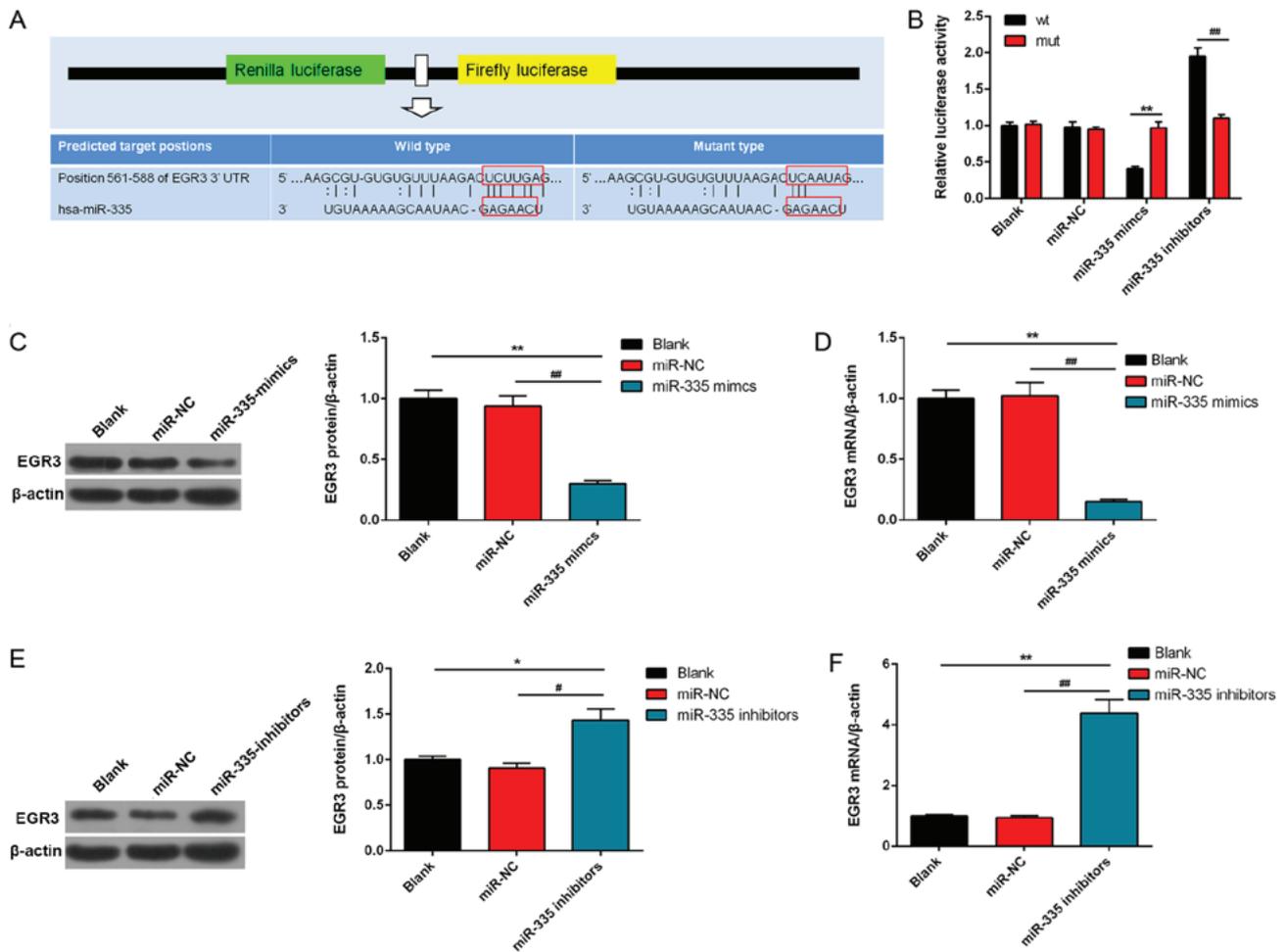


Figure 7. EGR3 is a potential target gene of miR-335. (A) EGR3 had a miR-335-binding site mapped to its 3'-UTR. (B) The effects of miR-335 mimics and miR-335 inhibitors on the luciferase activities of DU145 cells transfected with EGR3-wt 3'-UTR vector or EGR3-mut 3'-UTR. (C and D) EGR3 expression was significantly reduced in the miR-335 mimics group at protein and mRNA levels. (E and F) Protein and mRNA expression levels of EGR3 were significantly enhanced in the miR-335 inhibitors group. * $P < 0.05$, ** $P < 0.01$, # $P < 0.05$, ## $P < 0.01$, $n = 3$. EGR3, early growth response 3; miR, microRNA; mut, mutant; NC, negative control; 3'-UTR, 3'-untranslated region; wt, wild-type.

compared with the controls ($P < 0.01$; Fig. 8C). In addition, the number of regenerative vessels were notably reduced in siEGR3 group, and the tube length was significantly decreased compared with the controls ($P < 0.01$; Fig. 8D and E). Following transfection with siEGR3, the number of apoptotic cells was significantly elevated compared with the control cells ($P < 0.01$; Fig. 8F and G). Furthermore, the number of caspase-3 positive cells markedly increased following transfection with siEGR3 (Fig. 9A). Correspondingly, caspase-3 activity was significantly promoted in the siEGR3 group compared with the controls ($P < 0.01$; Fig. 9B). Additionally, the protein and mRNA expression levels of IL-6, IL-8 and IL-1 β were significantly suppressed in the siEGR3 group compared with the controls ($P < 0.01$; Fig. 9C and D). The viability of cells was significantly increased after miR-335 knockdown compared with cells transfected with miR-335 inhibitors and siEGR3, of which the viability was notably higher than those in non-transfected cells, and cells transfected with miR-NC or siNC ($P < 0.05$; Fig. 9E).

miR-335 inhibits the formation of PC solid tumor in vivo. In order to explore the effects of miR-335 on the tumorigenicity of PC, nude mice were transplanted with PC-3M or DU145

cells to establish PC models *in vivo*. The results revealed that the mRNA expression levels of miR-335 were significantly upregulated in PC-3M and DU145 cells compared with the controls ($P < 0.01$; Fig. 10A). The tumor weight and volume exhibited no significant differences between the blank and miR-NC groups; however, when mice were transplanted with PC-3M + miR-335 mimic or DU145 + miR-335 mimic, the weight and volume of tumors were significantly reduced ($P < 0.01$ and $P < 0.05$, respectively; Fig. 10B-E). IHC staining demonstrated that EGR3 was positively expression in the blank and NC groups; tissues expressing miR-335 mimic exhibited significantly downregulated EGR3 protein expression ($P < 0.01$; Fig. 10F and G). The results indicated that miR-335 inhibited the formation of PC solid tumor possibly by decreasing the expression of EGR3.

Discussion

miRNAs can regulate numerous genes at the post-transcriptional level, and the aberrant expression of miRNAs is closely associated with the development of neoplasms in humans (36). Recently, a substantial number of cancer-specific miRNAs

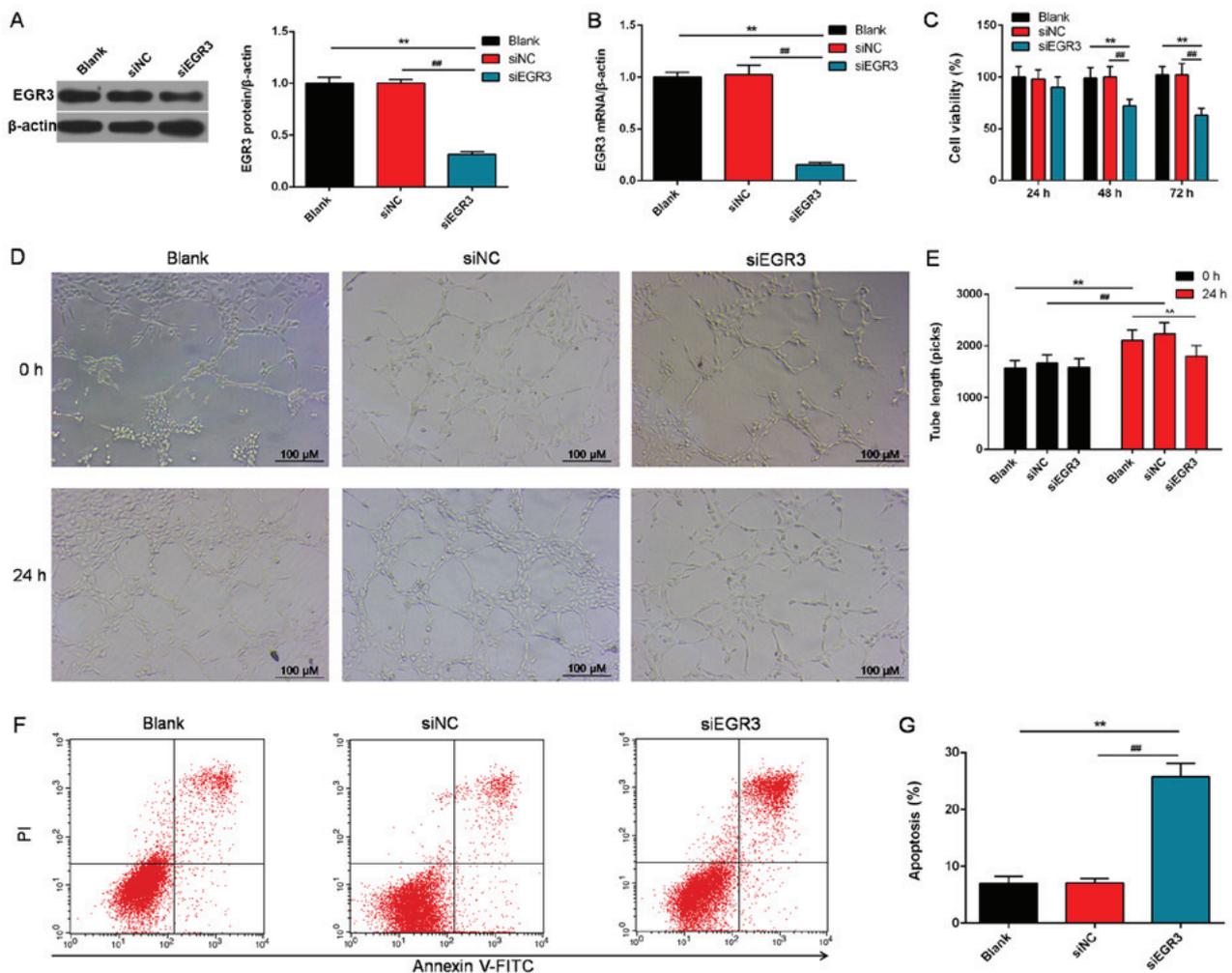


Figure 8. Role of EGR3 in cell viability, angiogenesis and apoptosis. (A and B) The expression of EGR3 was successfully knocked down at protein and mRNA levels following transfection with siEGR3. (C) Cell viability was significantly inhibited by the knockdown EGR3. (D and E) The length and number of tubes of HUVECs were reduced in siEGR3 group. (F and G) Apoptosis was significantly promoted by silencing EGR3 expression. Scale bar=100 μ m. ** P <0.01, ## P <0.01, ^ P <0.01, n=3. EGR3, early growth response 3; NC, negative control; si, small interfering RNA.

have been identified as tumor biomarkers in various types of cancer (37,38). Several miRNAs, including miR-1, miR-133a, miR-143 and miR-145 were reported to inhibit tumor development by targeting oncogenic genes in PC (39-41). EGR3, a member of zinc finger transcription factors, was demonstrated to be involved in angiogenesis, inflammation and neurodevelopment (42-44). Furthermore, it was revealed that EGR3 is involved in the occurrence and development of numerous types of cancers, and that EGR3 may be regarded as a tumor suppressor in various cancers (30,35,45). In the study, we demonstrated that miR-335 expression was downregulated, while EGR3 expression was enhanced in PC tissues. Additionally, miR-335 suppressed cell viability, angiogenesis and inflammation, and promoted the apoptosis of PC cells. *In vivo* miR-335 inhibited the formation of PC solid tumor. We also reported that EGR3 could be a potential target of miR-335 and inhibition of EGR3 inhibited the tumorigenesis of a xenograft PC *in vivo*.

Downregulated miR-335 expression has been reported in many types of cancer, such as BCa, acute myeloid leukemia, gastric cancer and adrenocortical carcinomas (16,46-48). In the present study, the miR-335 expression was significantly

reduced in the tissues of patients with PC, indicating that miR-335 may serve a vital role in tumorigenesis (49). The malignant transformation of normal cells is a multilevel and intricate process that is associated with hereditary and epigenetic changes (50). Oncogenes and tumor-suppressor genes could be closely related to proliferation and/or apoptosis; transformed cells may possess anti-apoptosis, invasive, and/or metastatic properties, thus leading to tumorigenesis (51). In addition, miR-335 can function as a tumor suppressor by inhibiting proliferation and inducing apoptosis in various cancer types. For instance, miR-335 expression is reduced in BCa tissues, blood and cells; and miR-335 regulates the viability and apoptosis of cells by targeting the upstream genes of BRCA1 in BCa (23,24). Furthermore, elevated miR-335 expression inhibits the proliferation and mammosphere formation of BCa stem-like cells (52). Overexpression of miR-335 can also reduce cell viability by inhibiting rho-associated, coiled-coil-containing 1 expression in hepatocellular carcinoma cells (53), and can promote apoptosis by targeting specificity protein 1 and BCL-w in lung cancer (54). Furthermore, miR-335 can also suppress cell viability and induce apoptosis by targeting members of

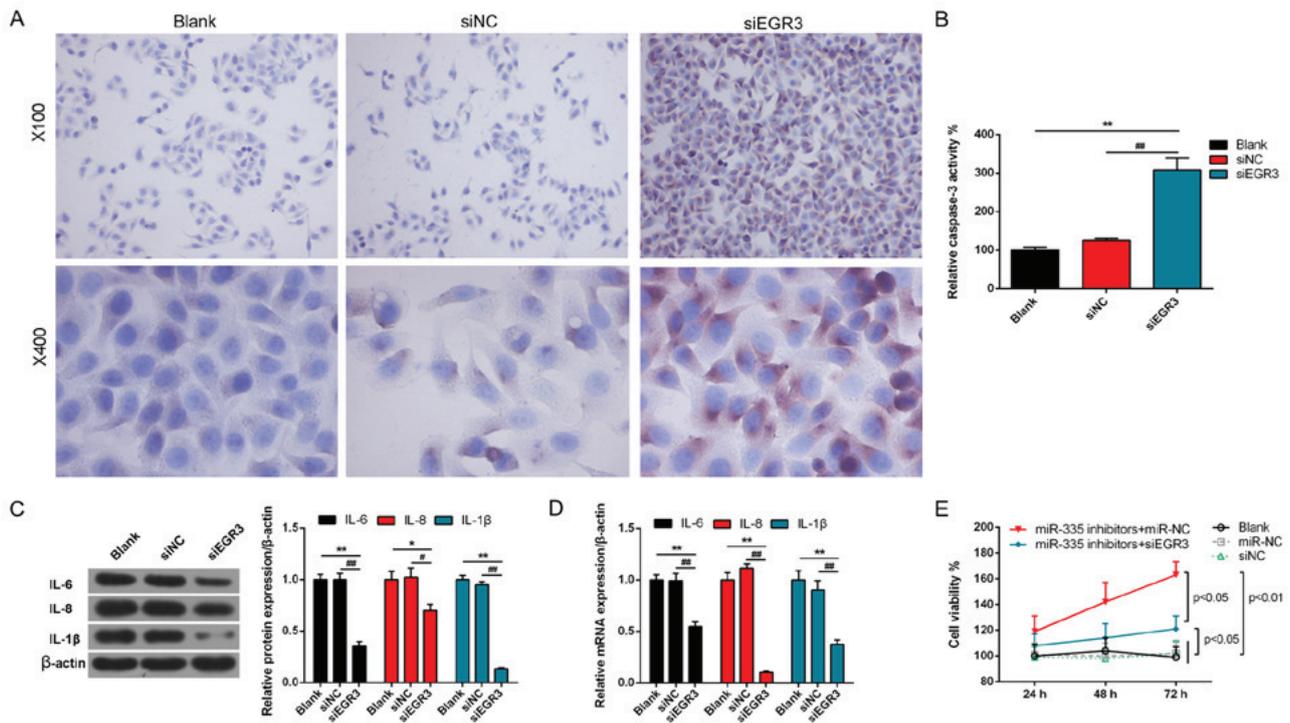


Figure 9. miR-335 exhibits anticancer effects by regulating EGR3 in DU145 cells. (A and B) Caspase-3 activity was significantly enhanced by silencing EGR3 expression. Magnifications, x200 and 400. (C and D) The protein and mRNA expression levels of IL-6, IL-8 and IL-1 β were significantly inhibited by downregulating EGR3. (E) miR-335 inhibitors suppressed the viability of cells following the downregulation of EGR3. *P<0.05, **P<0.01, #P<0.05, ##P<0.01, n=3. EGR3, early growth response 3; IL, interleukin; miR, microRNA; NC, negative control; si, small interfering RNA.

the inhibitors of apoptosis protein family IAP; miR-335 can regulate the cell cycle and the apoptotic pathways in gastric cancer (55). miR-335 can initiate the p53 pathway to inhibit cell proliferation by targeting the 3'-UTR of retinoblastoma 1, and increased proliferation caused by impaired p53 signaling has been linked to increased miR-335 levels, suggesting that the regulation of cell viability by miR-335 may be dependent on p53 (56). Apoptosis is relevant to various pathological processes, and the cells which evade apoptosis are involved in tumorigenesis (25). miR-335 was reported as a proapoptotic miRNA (57); low miR-335 expression can result in the proliferation and migration of mesenchymal stem cells (58). Additionally, inflammation is one of the indicators of cancer and exerts a crucial effect on tumor development (59,60). A previous study indicated that miR-335 may be involved in the IL-6/JNK2/STAT3 pathway (61). We observed that overexpression of miR-335 could inhibit the viability and angiogenesis, and promote the apoptosis of PC cells and reduce the expression of inflammatory cytokines. In addition, we also silenced the expression of miR-335, and found that the inhibition of miR-335 could exert opposing effects on than miR-335 overexpression. These results confirmed that miR-335 could function as a potential tumor suppressor of PC (62).

miR-335 regulates numerous target genes involved in the progression of cancers (14). In the present study, EGR3 was identified as a candidate target gene of miR-335 based on bioinformatics analysis. Furthermore, the dual-luciferase assay also suggested that EGR3 was a potential target of miR-335 and was negatively regulated by miR-335 in PC cells. A previous study reported reduced EGR3 expression in gastric cancer tissues, which was associated with adverse prognosis (35);

however, EGR3 overexpression increased cell invasion in BCa (63), suggesting that EGR3 may serve complicated roles in a tumor-specific manner. Of note, we demonstrated that EGR3 expression was enhanced in PC tissues, which was inversely correlated with the expression of miR-335. This suggested that EGR3 may serve a potential role in PC. Furthermore, EGR3 can mediate multiple biological processes (64). EGR3 was revealed to be involved in angiogenesis, inflammation and cancer (32,65). It has been reported that IL-6 can result in enhanced proliferation and epithelial-mesenchymal transition of PC cells (66). In addition, IL-8 silencing increased apoptosis, induced cell cycle arrest and improve chemotherapeutic outcomes of PC cells (67). In the present study, we reduced the expression of EGR3 in DU145 cells, and observed that the knockdown of EGR3 inhibited the viability and angiogenesis, promoted apoptosis and decreased the expression of inflammatory cytokines in these cells. The results were related to suppressed cell viability, as reported with miR-335 overexpression. Furthermore, we synchronously inhibited the expression of miR-335 and EGR3, and found that the viability of cells was reduced, indicating that miR-335 inhibited cell viability by targeting the EGR3.

In this study, there were some limitations. For instance, more cell lines are required to validate the presented data; more groups could also be established to demonstrate the targeting of miR-335 to EGR3. We aim to conduct more detailed and comprehensive experiments in the future, using certain groups: miR-335 mimics + EGR3 [wild-type (wt)], miR-335 mimics + EGR3 (mutant), miR-335 inhibitors + EGR3 (wt), miR-335 inhibitor + EGR3 (mutant) and another cell line.

In conclusion, miR-335 decreased cell viability and angiogenesis, suppressed inflammatory-marker expression

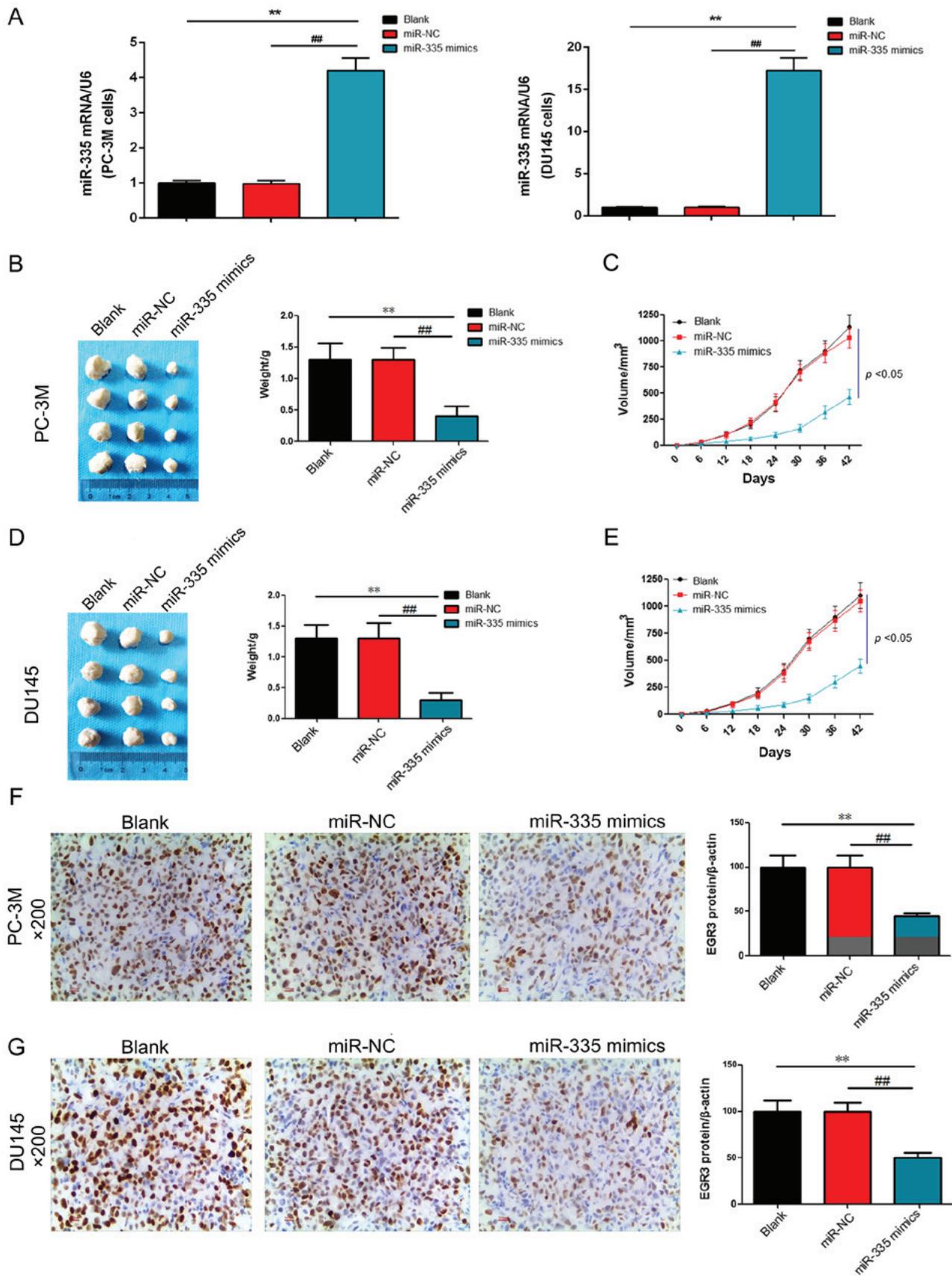


Figure 10. miR-335 inhibits the formation of PC solid tumor by regulating EGR3 expression *in vivo*. A xenograft mouse model of prostate cancer was generated by a hypodermic injection of 5×10^6 cells into the lateral abdominal wall of mice. PC-3M-derived tumors were collected 42 days after administration. (A) miR-335 expression was successfully upregulated in the xenograft tumors of in PC-3M and DU145 cells expressing miR-335 mimics. (B) Tumor weight and (C) volume in PC-3M cells were quantified. DU145-derived tumors were collected 42 days after the administration of cells. (D) Tumor weight and (E) volume in DU145 cells were quantified. Immunohistochemistry staining was performed to analyze EGR3 protein expression in (F) PC-3M-derived tumors tissues and (G) DU145-derived tumors tissues. Magnification, x200. β -actin served as an internal control. Data were expressed as mean \pm standard deviation from three independent experiments. ** $P < 0.05$, ## $P < 0.05$. EGR3, early growth response 3; miR, microRNA; NC, negative control.

and promoted apoptosis by targeting the EGR3 in PC cells. These findings suggested that miR-335 is closely associated with the initiation and progression of cancer, and may act as a potential biomarker in the treatment of patients with PC.

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Availability of data and materials

The analyzed data sets generated during the study are available from the corresponding author on reasonable request.

Authors' contributions

PZ and XY made substantial contributions to conception and design of the present study. LW and DZ conducted data acquisition, data analysis and interpretation. PZ drafted the article and critically revised it for important intellectual content. All authors approved for the final version to be published. BW and QL agreed to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of the work are appropriately investigated and resolved. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The present study was approved by the Ethics Committee of The Second Affiliated Hospital of Xi'an Jiaotong University

Patient consent for publication

Not applicable.

Competing interests

The authors declare no conflicts of interest.

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