



Downregulation of miR-182-5p inhibits the proliferation and invasion of triple-negative breast cancer cells through regulating TLR4/NF- κ B pathway activity by targeting FBXW7

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Background: To investigate the role of miR-182-5p in the proliferation and invasion of triple-negative breast cancer (TNBC) cells, as well as the underlying mechanism.

Methods: qRT-PCR was used to detect the level of miR-182-5p in tissues and cells. CCK-8 assay, flow cytometry, and Transwell assay were performed to detect cell proliferation, apoptosis rate, and invasion, respectively. Pearson correlation analysis was used to determine the correlation between miR-182-5p and its predicted target gene FBXW7, and the target of miR-182-5p was identified by dual-luciferase reporter gene assay. ELISA assay was used to examine the levels of cytokines in the culture supernatant of tumor cells. Western blot analysis was used to detect the expressions of FBXW7, TLR4, and NF- κ B pathways in tumor cells.

Results: In MDA-MB-231 and BT-549 cells, downregulation of miR-182-5p significantly inhibited cell proliferation and invasion and promoted tumor cell apoptosis. Pearson correlation analysis showed that miR-182-5p had a negative correlation with FBXW7. Dual-luciferase reporter gene assay showed that miR-182-5p could directly target FBXW7. Further studies showed that FBXW7 overexpression significantly inhibited cell proliferation and invasion and increased the apoptosis rate. Downregulation of miR-182-5p significantly reduced the levels of TNF- α , IL-1 β , IL-6, and IL-18 in the culture supernatant, and decreased the activity of TLR4/NF- κ B pathway in tumor cells, while downregulation of FBXW7 significantly inhibited the effect of miR-182-5p on tumor cells.

Conclusions: Downregulation of miR-182-5p inhibits TLR4/NF- κ B pathway activity by increasing FBXW7 expression, thereby suppressing the proliferation and invasion of TNBC cells.

Keywords: miR-182-5p; triple-negative breast cancer (TNBC); proliferation; invasion; TLR4/NF- κ B pathway

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Introduction

Breast cancer (BC), the most common cancer among women in the world, can be divided into many molecular subtypes, including Luminal A, Luminal B, human epidermal growth factor receptor 2 (HER2)-overexpressing, normal breast-like and basal-like or triple-negative breast

cancer (TNBC). TNBC is a BC subtype that does not express the estrogen receptor (ER), progesterone receptor (PR), and HER2. TNBC can be further classified into six subtypes, including immunomodulatory, mesenchymal, mesenchymal stem-like, luminal androgen receptor, basal-like 1, and basal-like 2. The total incidence rate of TNBC

is about 15–20%. Compared with hormone receptor and HER2 positive BC, TNBC has a highly invasive clinical process, with earlier onset age, more significant metastatic potential, poorer clinical prognosis, higher recurrence rate, and lower survival rate (1,2). The traditional treatment of TNBC mainly includes adjuvant therapy, surgical treatment and radiotherapy. Adjuvant therapy is a key strategy for cancer treatment, which can avoid the risk of metastasis accompanied by rapid progression and tumor recurrence. However, chemoresistance is a major problem in adjuvant treatment of cancer, with a failure rate of up to 90% in the treatment of metastatic cancer (3). In addition, TNBC is not sensitive to endocrine therapy or molecular targeted therapy because of its special molecular phenotype. Therefore, systemic chemotherapy is still the main means of treatment, but conventional postoperative adjuvant radiotherapy and chemotherapy has poor efficacy, residual metastases will eventually lead to tumor recurrence, and more side effects (4). However, there is no approved targeted therapy strategy for TNBC. Systemic chemotherapy is still the primary method of treatment, and the therapeutic effect of chemotherapy often lasts for a short time, causing the treatment of TNBC to be a clinical challenge (4). Therefore, the development of more effective treatment strategies is of great significance for the treatment of TNBC.

Increasing evidence showed that inflammation contributes to the occurrence of tumors. Inflammatory cells can promote tumor formation, release survival factors, promote angiogenesis and lymphangiogenesis, stimulate DNA damage, reshape the extracellular matrix to promote invasion, coat tumor cells to provide receptors that transmit cells through lymphatics and capillaries, and evade host defense mechanisms (5). Toll-like receptor 4 (TLR4), one of the essential receptors on the surface of immune cells, is often expressed in tumor cells and participates in the progression, invasion, and drug resistance of BC (6,7). TLR4 activates myeloid differentiation factor 88 (MyD88) after receiving tumor antigen information, promotes the nuclear transfer of nuclear factor- κ B (NF- κ B), and activates gene transcription, inducing the production of pro-inflammatory cytokines and causing inflammatory reactions (8,9). According to research reports, NF- κ B pathway may be a key regulator in the progress of TNBC (10), which can participate in the growth, apoptosis, differentiation and inflammatory response of tumor cells by promoting the transcription of tumor regulatory factors, including cytokines, chemokines, cell adhesion molecules, and

inducible pro-inflammatory enzymes, and ultimately affect the invasion and progress of tumor (11-14).

microRNAs (miRNAs), a type of small non-coding RNA composed of 18-24 single nucleotides, participates in the tumorigenesis process of various cancers, including BC (15). For instance, miR-185-5p inhibits invasion, migration, and epithelial-mesenchymal transition by regulating the specific receptor for advanced glycation end products in BC. MiR-19b can be used as a prognostic biomarker for BC and can promote cell proliferation, migration, and invasion by mediating phosphatidylinositol 3-kinase and serine-threonine kinase pathways (16). MiR-182, a miRNA with dual functions, is found on human chromosome 7q32.2 in the miR-183/96/182 cluster. It has been proved that miR-182 is up-regulated in TNBC, melanoma, endometrium, colon, and prostate and plays a tumor inhibitory role in lung cancer and gastric cancer (17-19). Also, studies have revealed that miR-182-5p promotes the invasion and proliferation of prostate cancer cells by mediating forkhead protein F2, reversal inducing cysteine-rich Kazakh motif protein, and transferring inhibitor 1 to promote the invasion and proliferation of prostate cancer cells (20). Inhibition of miR-182-5p can induce apoptosis and antiproliferation by regulating the expression of Caspase 9 in BC (18). Recent studies have shown that down-regulation of miR-182-5p can inhibit the proliferation and invasion of breast cancer by targeting the phosphatase tensin homologous gene (PTEN) deleted on chromosome 10 (21). However, the role and potential molecular mechanism of miR-182-5p in TNBC are still unclear.

In this study, we hypothesized that inhibition of miR-182-5p could take part in the regulation of TNBC cell proliferation and invasion by downregulating TLR4/NF- κ B inflammatory pathway activity. For this purpose, we evaluated the effect of miR-182-5p down-regulation on TNBC cell proliferation, apoptosis, and invasion as well as its possible mechanism by performing miR-182-5p loss of function experiment.

We present the following article in accordance with the MDAR reporting checklist (available at <http://dx.doi.org/10.21037/atm-20-5192>).

Methods

Materials

The tissue samples and adjacent normal tissues of 50 patients with TNBC treated in State Key Laboratory

Table 1 qRT-PCR primer sequences

Gene	Primer	Sequence
<i>miR-182-5p</i>	F	5'-TGCGGTTTG GCAATGGTAGAAC-3'
	R	5'- CCA GTGCAGGGTCCGAG GT-3'
<i>FBXW7</i>	F	5'-ACTGGAAGTGAC TCTGGGA-3'
	R	5'-TACTGGGGCTAGGCAAACAA-3'
<i>U6</i>	F	5'-TTATGGGTCCTAGCCTGAC-3'
	R	5'-CACTATTGCGGGCTGC-3'
<i>GAPDH</i>	F	5'-ACAGCAACAGGGTGGTGGAC-3'
	R	5'-TTTGAGGGTGCAGCGAACTT-3'

of Oncology in South China, Sun Yat-sen University Cancer Center from January 2017 to December 2019 were collected, and the basic information of patients including age, gender, blood pressure, tumor stage, whether there was distal metastasis and whether there was lymph metastasis was recorded. Normal human breast cells MCF10A, TNBC cells MDA-MB-231, MDA-MB-453, MDA-MB-361, and BT-549 were purchased from Shanghai Cell Bank of the Chinese Academy of Sciences (Shanghai, China). This study was approved by the Ethics Committee of Sun Yat-sen University Cancer Center (No. RDDA2020001590).

Cell culture and transfection

Cells were cultured in DMEM medium supplemented with 10% fetal bovine serum (FBS), 100 mg/mL penicillin, and 100 U/mL streptomycin in a constant temperature incubator at 37 °C, 5% CO₂, saturation humidity. Cell transfection: MDA-MB-231 and BT-549 cells were selected for miR-182-5p loss of function experiment. Experiment 1: cells were divided into negative control group (NC) and miR-182-5p loss of function group (miR-182-5p inhibitor). The medium was replaced with a serum-free medium that did not have penicillin and streptomycin during transfection. When cell culture reached about 70% confluence, miR-182-5p inhibitor NC or miR-182-5p inhibitor are transfected into MDA-MB-231 or BT-549 cells respectively in line with the description of Lipofectamine™ reagent. Experiment 2: the cells were divided into two groups: negative control group (pc-NC) and FBXW overexpression group (FBXW7). When the fusion degree of cell culture reached about 70%, the overexpression plasmids of pc-NC and pcDNA3.1-FBXW7

were transfected into MDA-MB-231 or BT-549 cells in the same way as above. Experiment 3: cells were divided into empty-NC group (pc-NC), FBXW7 overexpression group (FBXW7), and miR-182-5p inhibitor + FBXW7 interference group (miR-182-5p inhibitor + si-FBXW7). When the fusion degree of cell culture reached about 70%, miR-182-5p inhibitor NC or miR-182-5p inhibitor and si-FBXW7 plasmids were transfected into MDA-MB-231 or BT-549 cells in the same way. The above plasmids of miR-182-5p inhibitor/NC, pcDNA3.1, pc-NC and si-FBXW7/si-NC plasmids were designed and completed by Sangon Biotech (Shanghai) Co., Ltd. (Shanghai, China). After transfection for 6–8 hours, the culture medium was replaced by normal complete medium.

qRT-PCR to detect miR-182-5p and FBXW7 expression

Trizol kit (Invitrogen, CA, USA) was employed to extract total RNA in cells or tissues. The concentration of RNA was determined by NanoDrop 2000 spectrophotometer (Thermo Fisher Scientific, USA). MiRNA was reverse transcribed into cDNA utilizing TaqMan miRNA reverse transcription kit (Thermo Fisher Scientific, USA). The expression level of miR-182-5p was determined by TaqMan miRNA detection kit (Thermo Fisher Scientific, USA). PrimeScript™ RT (Takara, Japan) was used for reverse transcription to synthesize cDNA of mRNA. PCR amplification was performed by SYBR® Green Real-time PCR-Master Mix (Toyobo, Japan) to detect the level of FBXW7. The expression of miR-182-5p and FBXW7 were normalized with U6 and GAPDH as internal parameters, respectively, and the relative expression was calculated using the 2^{-ΔΔCt} method. QRT-PCR reaction conditions: pre-denaturation at 95 °C for 10 min, then 45 cycles of incubation at 95 °C for 15 s and 60 °C for 60 s. The experiment was repeated three times. The primer sequences of the target gene were shown in Table 1.

Detection of cell proliferation, apoptosis, and invasion

Cell counting kit-8 (CCK-8; Dojindo, Japan) was adopted to detect cell proliferation. Stably transfected cells were collected and seeded into 96-well plates at a density of 1×10⁴ cells per well. After culturing for 24, 48, or 72 hours respectively, cells were treated with 10 μL CCK-8 reagent for 2 hours. The absorbance was measured at 450 nm by enzyme immunoassays (Bio-Rad Laboratories, CA, USA), and the cell proliferation ability was determined.

Flow cytometry detected the apoptosis level. Cells in the logarithmic phase were collected and separated by trypsin to prepare cell suspension. Then, the cells were fixed overnight with cold ethanol at 4 °C, centrifuged for 5 minutes, and incubated with 5 µL of annexin V- FITC. After 3 min, the cells were supplemented with 10 µL propidium iodide (PI), incubated at 37 °C for 15 min, centrifuged, and resuspended in 0.5 mL precooled buffer. Flow cytometry (BD Biosciences, NJ, USA) was used to observe the apoptosis.

A Transwell experiment was used to detect cell invasion ability. MDA-MB-231 and BT-549 cells were resuspended in FBS-free DMEM and inoculated into the upper cavity of precoated matrix gel (Becton-Dickinson, CA, USA) for cell invasion analysis. DMEM medium holding 10% FBS was added to the lower chamber. After incubation for 24 hours, the remaining cells on the upper chamber were wiped off with cotton swabs. Cells infiltrated into the basal surface of the membrane were fixed with 4% paraformaldehyde, stained with 0.1% crystal violet after 10 min, and finally, the cells in 5 randomly selected fields were counted under a microscope (Olympus, Japan).

Double luciferase reporter gene experiment

The binding sites of miR-182-5p and FBXW7 were predicted by using the bioinformatics website (<http://www.targetscan.org>). FBXW7 3'-UTR promoter region sequence containing miR-182-5p binding site was synthesized, and FBXW7-WT and its corresponding mutation binding site FBXW7-MUT were constructed. MDA-MB-231 and BT549 cells in the logarithmic growth phase were inoculated onto 96-well plates. When cell confluence reached 70%, mimics NC or miR-182-5p mimics, FBXW7-WT and FBXW7-MUT were co-transfected into MDA-MB-231 and MCF-7 cells. Forty-eight hours after transfection, cell lysis was performed, and luciferase activity was measured by luciferase detection kit.

ELISA to detect the levels of TNF- α , IL-18, IL-1 β , and IL-6

The levels of TNF- α , IL-18, IL-1 β , and IL-6 in the supernatant of cell culture were determined by ELISA kits following the instructions. Streptavidin-HRP and tetramethylbenzidine sulfonate were used as color development indicators. ELISA microplate reader DR-200Bs (Jinan Laibao Laboratory Equipment Co., Ltd., Jinan, China) is used to read the values at 450 nm wavelengths, and the concentration level of corresponding

cytokines is calculated through standard curves.

Western blot to detect the protein expression of TLR4, MyD88, NF- κ B, and I- κ B

RIPA buffer (Sigma-Aldrich, MO, USA) containing protease inhibitor was used to split and extract total protein on ice, followed by centrifugation at 4 °C for 15 min after 40 min. The protein concentration was then measured using a BCA protein assay kit (Beyotime Biotechnology, Shanghai, China). SDS-PAGE separated the protein in 12% separation gel, and 4% concentrated gel and transferred to PVDF membrane. PVDF membrane carrying transfer protein is sealed in TBST buffer solution containing 0.1% Tween 20 and 5% bovine serum albumin (BSA) for 2 hours at normal temperature. Subsequently, the designated primary antibody (rabbit anti-FBXW7 antibody (1:1,000), rabbit anti-TLR4 antibody (1:1,000), rabbit anti-MyD88 antibody (1:1,000), rabbit anti-p-P65 antibody (1:1,000), rabbit anti-P65 antibody (1:1,000), rabbit anti-p-I-kappa b antibody (1:1,000), rabbit anti-I-kappa b antibody (1:1,000), and mouse anti-GAPDH antibody (1:5,000)) were incubated overnight at 4 °C. Then, the membrane was incubated with the corresponding secondary antibody for 2 hours. After enhanced chemiluminescence and exposure color development, the gray values of protein bands were analyzed by ImageJ software.

Statistical analysis

SPSS 22.0 software was used to process the data, and the results were expressed by the mean \pm standard deviation (SD) of n experiments. *T*-test was used to compare the two groups. Multiple comparisons between groups were conducted by one-way analysis of variance (ANOVA), followed by LSD (least significant difference) post hoc test, with $P < 0.05$ showing significant difference.

Results

MiR-182-5p is highly expressed in TNBC

To determine the relationship between miR-182-5p and TNBC, we detected the expression level of miR-182-5p in TNBC tissues and cells by qRT-PCR. The results revealed that the expression level of miR-182-5p in the tumor group was significantly higher than that in the normal group (*Figure 1A*). Whereas, the expression level of miR-182-5p

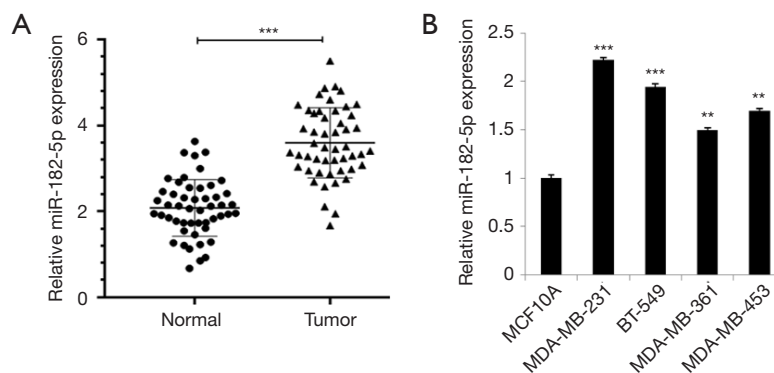


Figure 1 The expression of miR-182-5p in TNBC tissues and cells. (A) The expression of miR-182-5p in TNBC tissues. (B) The expression of miR-182-5p in TNBC cells. **, $P < 0.01$; ***, $P < 0.001$. TNBC, triple-negative breast cancer.

in TNBC cells was significantly higher than that in normal MCF10A cells, and the expression level was highest in MDA-MB-231 and BT-549 cells (Figure 1B). The above results suggested that miR-182-5p may contribute to the progress of TNBC.

Downregulation of miR-182-5p expression inhibits TNBC cell proliferation and invasion and promotes cell apoptosis

To further determine the role of miR-182-5p in TNBC, we evaluated the effect of miR-182-5p silencing on TNBC cell proliferation, apoptosis, and invasion through function loss experiment. According to the results in Figure 1B, MDA-MB-231 and BT-549 cells with the highest expression level of miR-182-5p were selected as the research objects of the follow-up experiment. CCK8 results manifested that downregulation of miR-182-5p expression significantly inhibited the cell proliferation ability of MDA-MB-231 (Figure 2A) and BT-549 cells (Figure 2B), and the inhibition became more evident with the prolongation of action time. Flow cytometry results evinced that downregulation of miR-182-5p expression significantly increased the apoptosis rate of MDA-MB-231 (Figure 2C) and BT-549 cells (Figure 2D). Transwell results verified that the downregulation of miR-182-5p expression significantly inhibited the invasion rate of MDA-MB-231 (Figure 2E) and BT-549 cells (Figure 2F). The above results suggested that miR-182-5p may contribute to TNBC cell proliferation and invasion and inhibit TNBC cell apoptosis.

MiR-182-5p targets FBXW7

The target of miR-182-5p was predicted through

microRNA.org and Targetscan online tools. FBXW7 scored the highest among all potential targets was found. We first detected the expression level of FBXW7 in TNBC tissue. QRT-PCR results identified that the expression level of FBXW7 in TNBC tissues was significantly lower than that in normal tissues (Figure 3A). Pearson correlation analysis suggested that the expression of miR-182-5p and FBXW7 in TNBC tissue was significantly negatively correlated (Figure 3B). Moreover, we constructed FBXW7 3'-UTR-WT and FBXW7 3'-UTR-MUT luciferase reporter plasmids (Figure 3C). After co-transfecting FBXW7 3'-UTR-WT or 3'-UTR-MUT and miR-182-5p mimics or NC into TNBC cells, we found that miR-182-5p mimic significantly reduced luciferase activity in the cells through double luciferin reporter gene experiments (Figure 3D). Further Western Blot tests indicated that downregulation of miR-182-5p expression in TNBC cells markedly increased FBXW expression (Figure 3E). The above results confirmed that miR-182-5p could target binding and negatively regulate FBXW7.

Overexpression of FBXW7 inhibits proliferation and invasion of TNBC cells and promotes apoptosis

We further discussed the effects of FBXW7 on the proliferation, apoptosis, and invasion of TNBC cells. CCK8 results suggested that overexpression of FBXW7 significantly inhibited the cell proliferation of MDA-MB-231 (Figure 4A) and BT-549 cells (Figure 4B), and the inhibition became more evident with the prolongation of the action time. Flow cytometry indicated that overexpression of FBXW7 remarkably increased the apoptosis rate of MDA-MB-231 (Figure 4C) and BT-

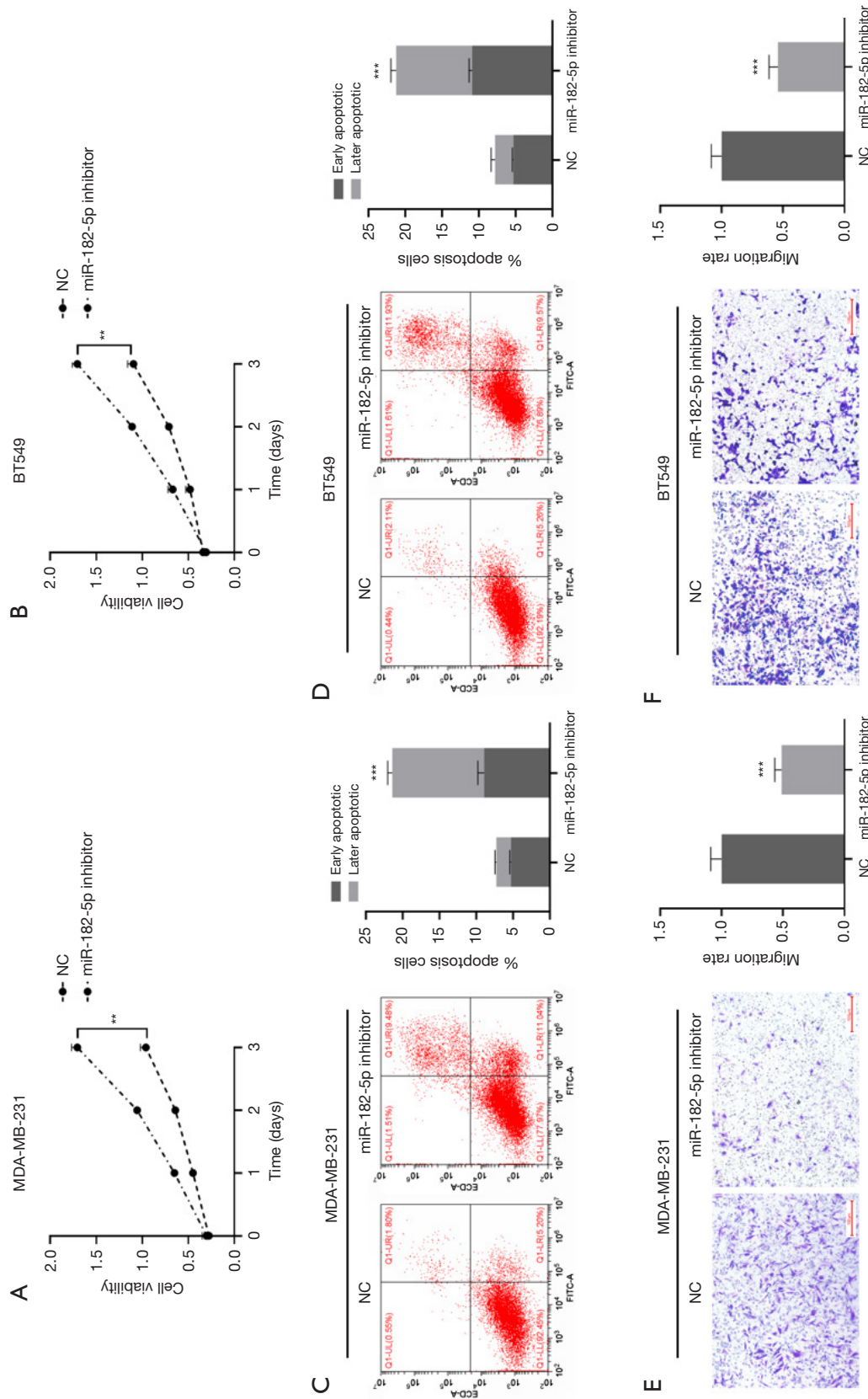


Figure 2 Effect of miR-182-5p on cell proliferation, apoptosis, and invasion. (A,B) Cell proliferation was examined by CCK8 assay in MDA-MB-231 and BT-549 cells. (C,D) The apoptosis rate was measured by flow cytometry in MDA-MB-231 and BT-549 cells. (E,F) Cell invasion was determined by Transwell assay in MDA-MB-231 and BT-549 cells. **, P<0.01; ***, P<0.001.

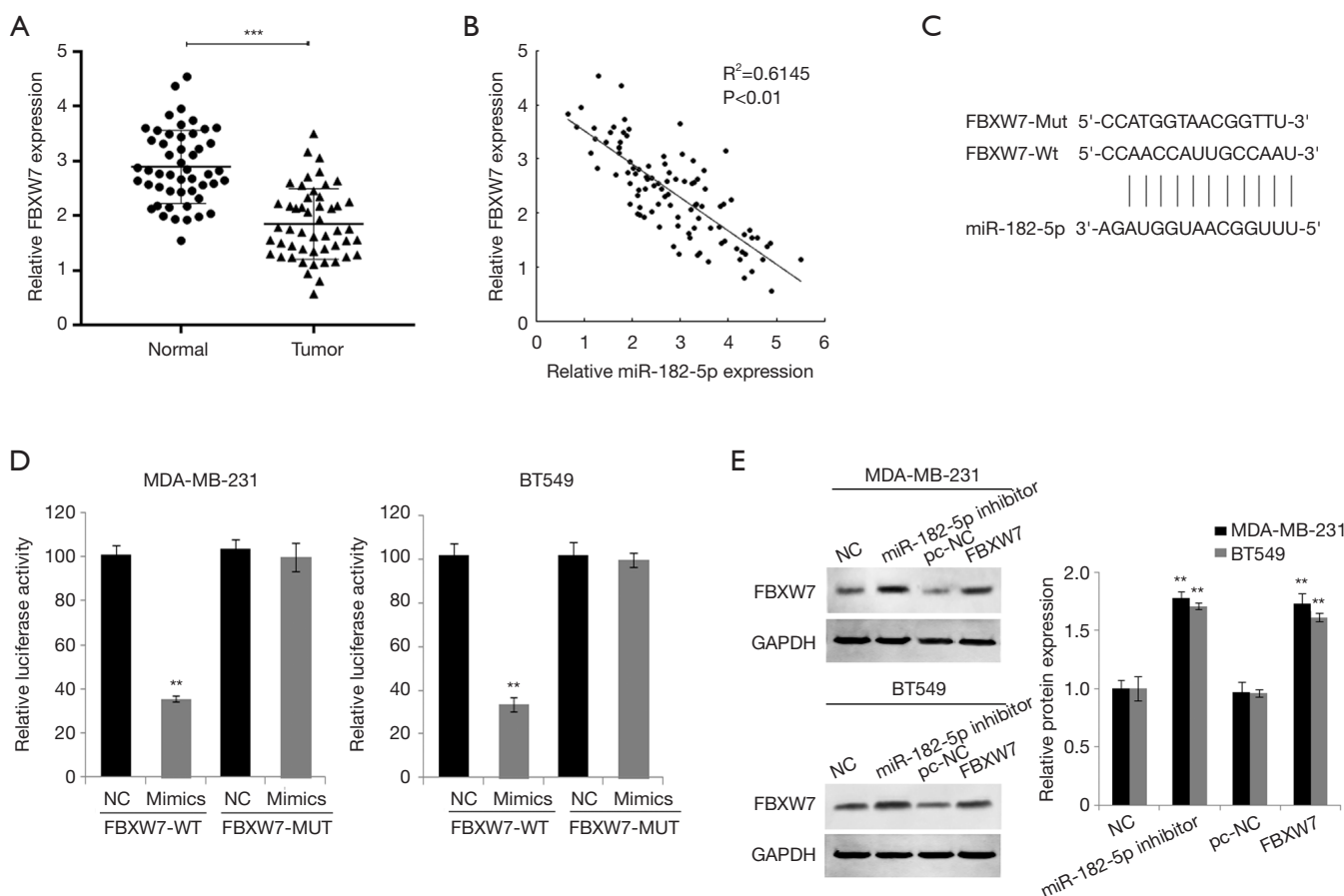


Figure 3 MiR-182-5p targets and negatively regulates FBXW7. (A) The expression of FBXW7 in TNBC tissues was examined by qRT-PCR. (B) Pearson correlation analysis was performed to identify the correlation between miR-182-5p and FBXW in TNBC tissues. (C) The schematic diagram reflected the complementary binding sites between miR-182-5p and FBXW. (D) Dual-luciferase reporter gene assay showed the luciferase activity in TNBC cells. (E) The expression of FBXW7 in TNBC cells was measured by Western blot assay. **, $P < 0.01$; ***, $P < 0.001$. TNBC, triple-negative breast cancer.

549 cells (Figure 4D). Transwell results confirmed that overexpression of FBXW7 notably inhibited the invasion rate of MDA-MB-231 (Figure 4E) and BT-549 (Figure 4F) cells. The above results suggested that FBXW7 may inhibit TNBC cell proliferation, invasion, and apoptosis.

Downregulation of miR-182-5p inhibits TNBC cell proliferation and invasion and promotes cell apoptosis via FBXW7

After co-transfecting TNBC cells with miR-182-5p inhibitor and si-FBXW7, we found that the down-regulation of FBXW7 expression reversed the inhibition of miR-182-5p expression on cell proliferation and invasion, and reversed the promotion of miR-182-5p expression down-regulation

on cell apoptosis (Figure 5A,B,C,D). The results suggested that miR-182-5p may play a regulatory role in tumor proliferation, apoptosis, and invasion by targeting FBXW7.

Downregulation of miR-182-5p inhibits TLR4/NF- κ B pathway activity in TNBC cells via FBXW7

FBXW7 gene is a p53-dependent tumor suppressor gene that takes part in the regulation of inflammation. Some studies have demonstrated that FBXW7 can inhibit inflammatory signal activity by downregulating the expression of C/EBP δ and its target gene TLR4 (3). In this study, we first detected the level of inflammatory cytokines in TNBC cell culture supernatant. ELISA results showed that the downregulation of miR-182-5p expression

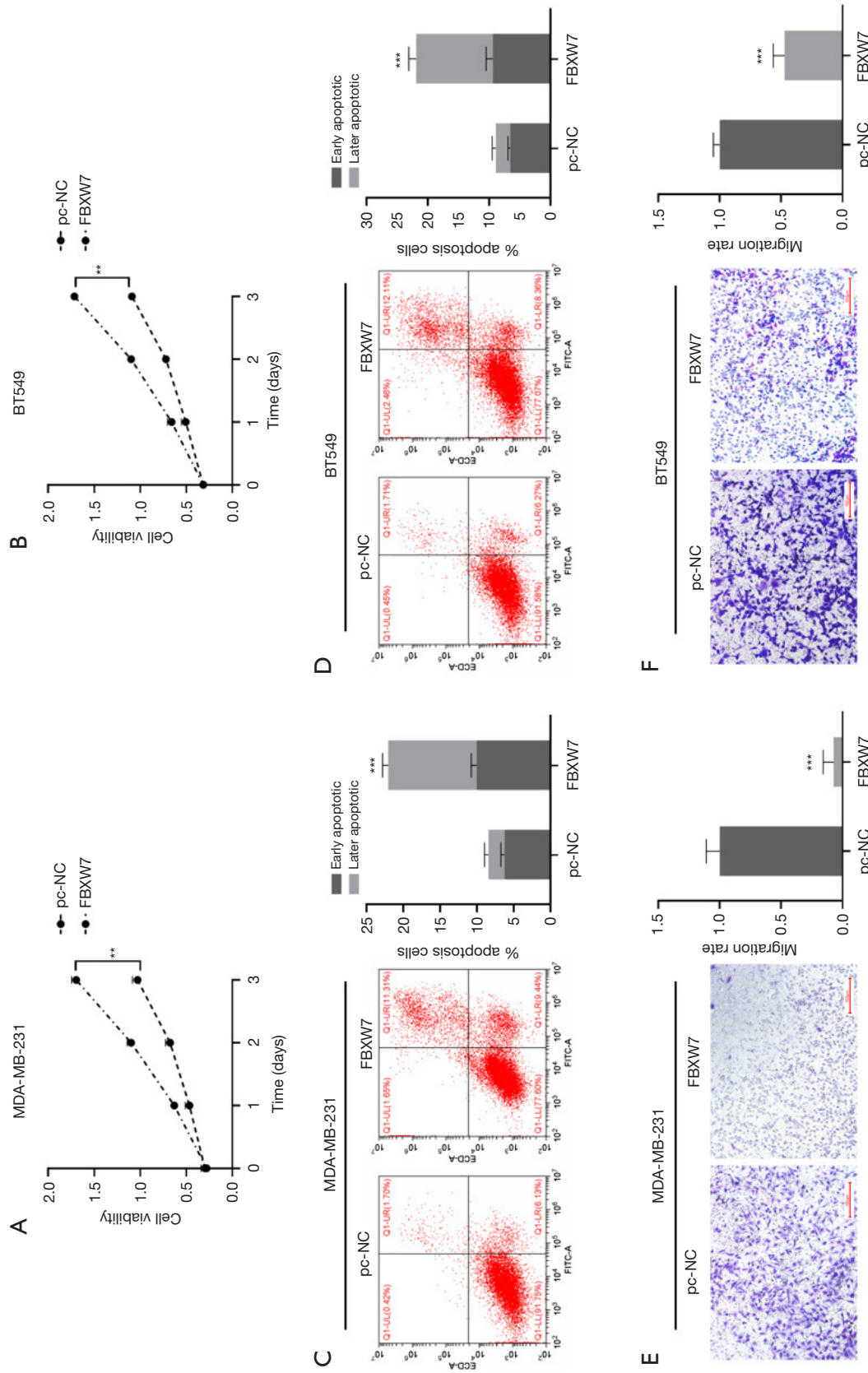


Figure 4 Effect of FBXW7 on the cell proliferation, apoptosis, and invasion. (A,B) Cell proliferation was examined by CCK8 assay in MDA-MB-231 and BT-549 cells. (C,D) The apoptosis rate was examined by flow cytometry in MDA-MB-231 and BT-549 cells. (E,F) Cell invasion was examined by Transwell assay in MDA-MB-231 and BT-549 cells. **, P<0.01; ***, P<0.001.

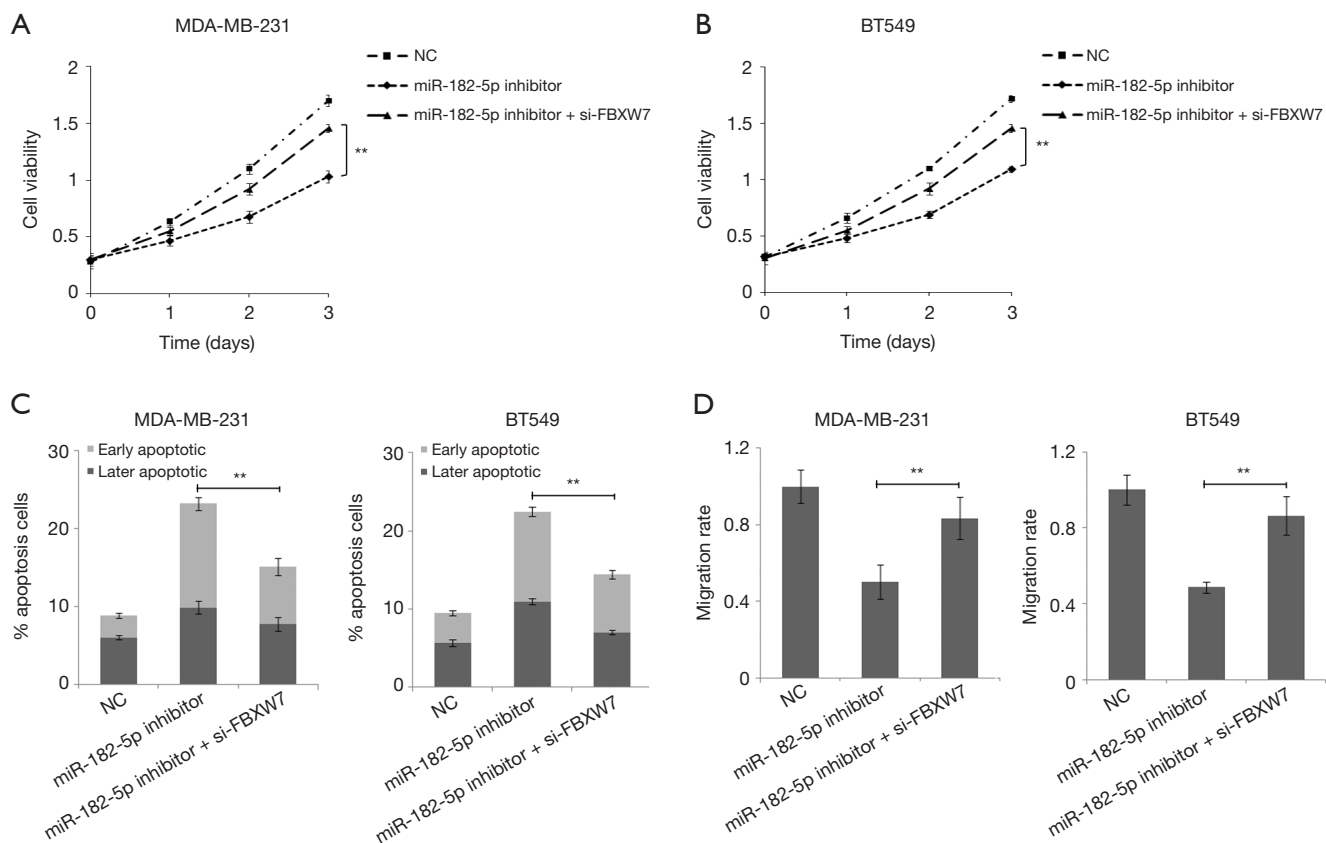


Figure 5 MiR-182-5p regulates TNBC cell proliferation, apoptosis, and invasion via FBXW7. (A,B) Cell proliferation was examined by CCK8 assay in MDA-MB-231 and BT-549 cells. (C) The apoptosis rate was examined by flow cytometry in MDA-MB-231 and BT-549 cells. (D) Cell invasion was examined by Transwell assay in MDA-MB-231 and BT-549 cells. **, $P < 0.01$.

significantly reduced the levels of TNF- α , IL-1 β , IL-6, and IL-18 in the supernatant of MDA-MB-231 and BT-549 cell cultures while inhibiting the expression of FBXW7 reversed the effect of miR-182-5p (Figure 6A,B). Further analysis by Western blot revealed that downregulation of miR-182-5p remarkably reduced the expression of TLR4, MyD488, p-p65, and p-I- κ B in MDA-MB-231 and BT-549 cells, while inhibition of FBXW7 significantly reversed the effect of miR-182-5p (Figure 6C,D). The above results indicated that the downregulation of miR-182-5p expression might inhibit the release of inflammatory factors and the activity of inflammatory pathways by up-regulating the expression of FBXW7, thus inhibiting the proliferation and invasion of TNBC cells and promoting cell apoptosis.

Discussion

TNBC, characterized by its highly aggressive and metastatic

features, has higher recurrence and mortality than other types of BC due to insufficient effective treatment (22). MiRNA, as a new biomarker, plays a significant role in regulating the invasion, proliferation, metastasis, apoptosis, and chemotherapy of BC (23). In this study, we hypothesized that miR-182-5p participates in the regulation of TNBC cell proliferation and invasion and discussed the target and potential mechanism of miR-182-5p. We found that miR-182-5p was overexpressed in TNBC tissues and cells. Down-regulation of miR-182-5p expression can inhibit the proliferation and invasion of TNBC cells and promote cell apoptosis. These results suggested that miR-182-5p may be a potential regulator in TNBC.

Through bioinformatics analysis, we predicted and verified the direct target FBXW7 of miR-182-5p, and the expression of the two in TNBC tissue was negatively correlated. *FBXW7* is a human tumor suppressor gene. It is reported that its total mutation rate in human tissues

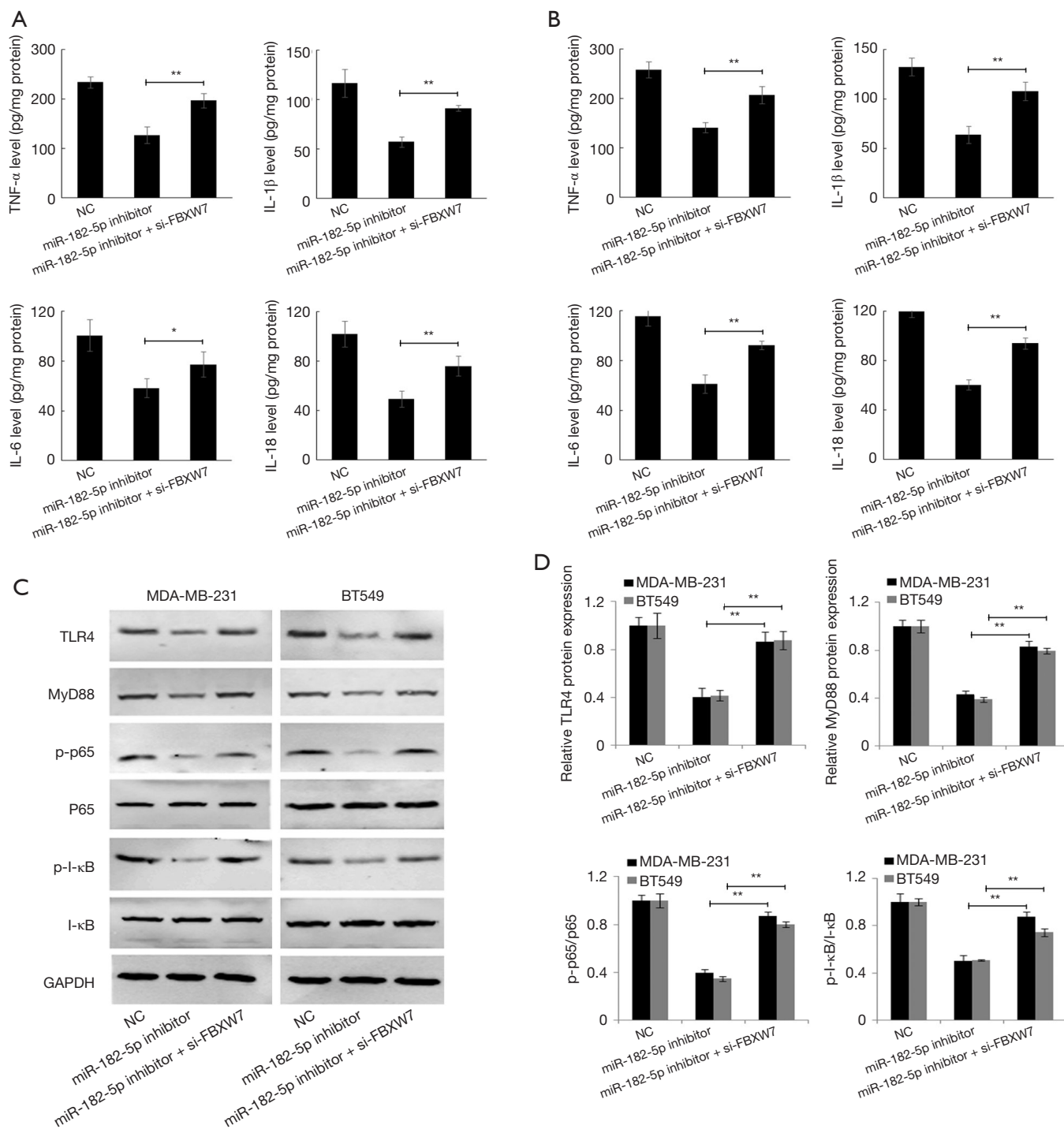


Figure 6 miR-182-5p regulates TLR4/NF-κB pathway activity in TNBC cells via FBXW7. (A,B) The levels of TNF-α, IL-1 β, IL-6, and IL-18 in MDA-MB-231 and BT-549 cells were evaluated by ELISA assay. (C) The expression of the TLR4/NF-κB pathway in MDA-MB-231 and BT-549 cells was examined by Western blot. (D) The quantitative diagram of the expression of the TLR4/NF-κB pathway. *, P<0.05; **, P<0.01. TNBC, triple-negative breast cancer.

including bile duct, blood, bone, brain, breast, colon, endometrium, stomach, lung, ovary, pancreas, and prostate is 6%, and it can take part in the regulation of genetic instability or growth disorder of tumors by affecting ubiquitination and cycle of various tumor proteins (24,25). Studies have confirmed down-regulation of FBXW7 expression promotes the proliferation of BC cells and inhibits cell apoptosis (26), whereas over-expression of FBXW7 can inhibit the proliferation of BC cells and promote cell apoptosis by targeting MTDH (27). In TNBC, FBXW7 loss of function can promote tumor growth and metastasis by stabilizing interferon receptor signals (28). In this study, we found that overexpression of FBXW7 significantly inhibited the proliferation and invasion of TNBC cells and promoted cell apoptosis. This result was consistent with earlier reports. Our further research showed that inhibiting FBXW7 expression can reverse the effect of miR-182-5p on TNBC cell proliferation, apoptosis, and invasion. The above results suggested that miR-182-5p regulates the proliferation, apoptosis, and invasion of TNBC cells through targeted negative regulation of FBXW7. However, the mechanism of miR-182-5p regulating TNBC cells via FBXW7 needs further research.

According to research, *FBXW7* can inhibit inflammatory signal activity on the one hand by downregulating the expression of C/EBP δ and its target gene TLR4. However, it is also inhibited by C/EBP δ and promotes tumor metastasis (3,29). It is suggested that FBXW7 is strongly associated with inflammatory signals in the regulation of tumor progression. Therefore, we have detected the level of inflammatory signals in TNBC cells. The results indicated that the downregulation of miR-182-5p expression significantly reduces the release level of inflammatory factors in TNBC cells, including TNF- α , IL-1 β , IL-6, and IL-18, while inhibiting the expression of FBXW7 reverses the reduction of these inflammatory factors, indicating that miR-182-5p may regulate the proliferation, apoptosis, and invasion of TNBC cells by regulating the level of inflammatory signals. In this process, FBXW7 is the critical mediator of miR-182-5p regulation. A TLR4/NF- κ B signaling pathway is widely reported taking part in the regulation of tumor progression and treatment through inflammatory reaction. For example, TLR4 can activate IL-4 receptor-related kinase 4 (IRAK4) and NF- κ B by combining with paclitaxel or lipopolysaccharide, promote the production of IL-6, IL-8, vascular endothelial growth factor, and monocyte

chemotactic protein-1, resist tumor cell apoptosis induced by chemotherapy drugs (30), and inhibit IL-6 and IL-8, thereby effectively reducing the chemotherapy drug resistance of BC mediated by TLR4 (7). In this study, we further detected the activity level of the TLR4/NF- κ B pathway in TNBC cells. The results reflected that the downregulation of miR-182-5p significantly reduced the expression of TLR4, MyD488, p-p65, and p-I- κ B in cells, while inhibition of FBXW7 significantly reversed the effect of miR-182-5p. This result supplied further evidence for the earlier conclusion.

In conclusion, this study proved that miR-182-5p participates in the regulation of TNBC cell proliferation, apoptosis, and invasion. The downregulation of miR-182-5p inhibits the production of inflammatory factors and activation of inflammatory signals in TNBC cells by targeting FBXW7, thereby inhibiting the proliferation and invasion of TNBC cells and promoting cell apoptosis. This study will supply new therapeutic targets and the theoretical basis for TNBC treatment. Nevertheless, the role of MiR-182-5p in TNBC tumor growth and metastasis needs further study.

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Footnote

Reporting Checklist: The authors have completed the MDAR reporting checklist. Available at <http://dx.doi.org/10.21037/atm-20-5192>

Data Sharing Statement: Available at <http://dx.doi.org/10.21037/atm-20-5192>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/atm-20-5192>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. This study was approved by the Ethics Committee of Sun Yat-sen University Cancer Center (No. RDDA2020001590).

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