

Up-regulation of miR-124 inhibits invasion and proliferation of prostate cancer cells through mediating JAK-STAT3 signaling pathway

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Abstract. – **OBJECTIVE:** Signal transducer and activator of transcription 3 (STAT3) is an important protein in Janus kinase (JAK)-STAT signaling pathway, and can facilitate expression of Bcl-2 and Cyclin D1 gene, thus playing a role in tumor pathogenesis. Bioinformatics analysis revealed targeted binding sites between microRNA-124 (miR-124) and 3'-UTR of STAT3 mRNA. This study aims to investigate the role of miR-124 in regulating STAT3 expression and proliferation, cycle, apoptosis and invasion of prostate cancer cells.

MATERIALS AND METHODS: Dual luciferase reporter gene assay demonstrated targeted regulation between miR-124 and STAT3. Expression of miR-124, STAT3, p-STAT3, Bcl-2 and Cyclin D1 were compared between normal human prostate epithelial cell RWPE-1 and prostate cancer cell DU145. *In vitro* cultured DU145 cells were treated with miR-124 mimic and/or si-STAT3, to compare expression of STAT3, phosphorylated STAT3 (p-STAT3), B-cell lymphoma-2 (Bcl-2) and Cyclin D1. Flow cytometry detected cell apoptosis, cell cycle, clonal formation and cell invasion. *In vivo* test malignant proliferation and cell invasion.

RESULTS: Targeted regulation was found between miR-124 and STAT3. Comparing to RWPE-1, DU145 cells had lower miR-124 expression, G0/G1 phase ratio, cell apoptosis, plus higher expression of STAT3, p-STAT3, Bcl-2 and Cyclin D1 ratio of S or G2/M phase. Transfection of miR-124 mimic and/or si-STAT3 remarkably decreased gene expression, weakened clonal formation, increased ratio of S and G2/M phase, increased apoptosis and increased G0/G1 ratio.

CONCLUSIONS: MiR-124 up-regulation significantly suppressed STAT3, pSTAT3 and down-regulated Bcl-2 and Cyclin D1 expression, weakens cell invasion or malignant proliferation potency, induces G0/G1 phase arrest, and facilitates cell apoptosis.

Keywords:

miR-124, STAT3, Cell apoptosis, Cell cycle, Invasion, Proliferation, Prostate cancer.

Introduction

Prostate cancer (PCa) is one common malignant tumor in male urinary-reproductive system, and is the sixth popular cancer in males¹. PCa frequently occurs in aged males, as more than 90% patients were between 60 and 80 years old. Geographic distribution of PCa patients showed specific patterns, as Western countries had significantly higher incidence than China². Janus kinase (JAK)-signal transducer and activator of transcription (STAT) signal transduction pathway is widely involved in regulating cell proliferation, apoptosis, migration and invasion, thus is closely correlated with tumor occurrence³. STAT3 is the most important member of STAT protein family. As one transcription factor, STAT3 can facilitate expression of genes involving in cell proliferation, cell cycle, apoptosis, invasion and degradation of extracellular matrix (ECM), and is thus closely correlated with occurrence, progression and distal metastasis of multiple tumors including pancreatic carcinoma⁴, colon cancer⁵ and breast cancer⁶. A previous study showed that anti-apoptotic factor B-cell lymphoma-2 (Bcl-2)^{7,8} and Cyclin D1^{9,10} were all targeted genes of STAT3 transcriptional factor. By enhancing gene transcription and expression, STAT3 participates in facilitating cell proliferation and cycle progression, and also in antagonizing cell apoptosis; thus, it is one STAT protein with most close correlation with human tumor pathogenesis. Recent studies^{11,12} showed the important role of enhanced STAT3 expression or functional activity in facilitating PCa pathogenesis. MicroRNA (miR) is one endogenous small RNA molecule in eukaryotic cells, and can regulate target gene expression via complementary binding on 3'-untranslated region (3'-UTR)

of target gene mRNA to degrade mRNA or inhibit mRNA translation, modulating cell proliferation, differentiation and migration. The role of abnormal expression of function of mRNA in tumor onset has drawn increasing research focus¹³. Studies showed significantly decreased miR-124 expression in PCa tissues/cells^{14,15}, suggesting its potential role as tumor suppressor gene in PCa occurrence. Bioinformatics analysis showed the existence of complementary binding sites between miR-124 and 3'-UTR of STAT3. This study investigated the role of miR-124 in regulating STAT3 and downstream target genes Bcl-2 and Cyclin D, and in affecting proliferation, cycle, apoptosis and invasion of PCa cells.

Materials and Methods

Major Reagent and Materials

Human prostate cancer cell line DU145 and normal prostate epithelial cell line RWPE-1 were purchased from Shengbo Biomed (Zhanjiang, China). Dulbecco's Modified Eagle's medium (DMEM), Keratinocyte-serum-free media (SFM) and serum free medium, fetal bovine serum (FBS) and penicillin-streptomycin were purchased from Gibco (Rockville, MD, USA). Trizol and Lipofectamine 2000 were purchased from Invitrogen/Life Technologies (Carlsbad, CA, USA). QuantiTect SYBR Green RT-PCR Kit was purchased from Qiagen (Hilden, Germany). miR-124 mimic and miR-NC nucleotide fragments were designed and synthesized by Ruibo Bio-Tech (Shanghai, China). siRNA sequences and negative control sequence were synthesized by Gene Pharmakon (Milpitas, CA, USA). Mouse anti-STAT3 and p-STAT3 was purchased from Abcam (Cambridge, MA, USA). Rabbit anti-Bcl-2 and Cyclin D1 were purchased from GeneTex (Irvine, CA, USA). Transwell chamber was purchased from Greiner Bio-One (Frickenhausen, Germany). Matrigel was purchased from Bio-Rad Laboratories (Hercules, CA, USA). Dual-luciferase gene reporter plasmid pLUC-STAT3-wt and pLUC-STAT3-mut were purchased from Ambioin (Carlsbad, CA, USA). Dual-Luciferase Reporter Assay System was purchased from Promega (Madison, WI, USA). PI dye, cell apoptosis assay reagent and RIPA lysis buffer were purchased from Beyotime (Jiangsu, China).

Cell Culture

DU145 cells were cultured in Dulbecco's Modified Eagle Medium (DMEM) containing 10%

fetal bovine serum (FBS) and 1% penicillin-streptomycin, and were kept in a humidified incubator with 5% CO₂ at 37°C. RWPE-1 cells were kept in Keratinocyte-SFM medium containing 5 ng/mL EGF, 0.05 mg/mL bovine pituitary extract (BPE) in a humidified chamber with 5% CO₂ at 37°C. Cells at log-growth phase with satisfactory growth status were used for further experiments.

Luciferase Reporter Gene Construct

Full length or mutant fragment of 3'-UTR of STAT3 gene was sub-cloned into pLUC-luciferase vector, which was named pLUC-STAT3-wt. Luciferase reporter vector containing mutant form of 3'-UTR of STAT3 gene was also constructed as pLUC-STAT3-mut. Lipofectamine 2000 was used to transfect pLUC-STAT3-wt (or pLUC-STAT3-mut) and miR-124 mimic (or miR-124 inhibitor, or miR-NC) into HEK293T cells. After 48 h, dual luciferase activity assay kit was used to test dual luciferase activity. Nucleotide sequences were: miR-NC, 5'-ACUAC UGAGU CAGG UAGA-3'; miR-124 mimic, 5'-GGCAU UCAU GCGG CCUUA-3'; miR-124 inhibitor, 5'-UAAGG CAGC GGUGA AUGCC-3'.

Transfection and Grouping

In *in vitro* cultured DU145 cells were divided into five groups: miR-NC transfection group, miR-124 mimic transfection group, si-NC transfection group, si-STAT3 group, and miR-124 mimic + si-STAT3 transfection group. 72 h after transfection, cells were collected for gene and protein expression assay. Nucleotide sequences for transfection were: si-STAT3 sense, 5'-CAUCU GC-CUA GAUCG GCUA-3'; si-STAT3 anti-sense: 5'-UAGCC GAUCU AGGCA GAUG-3'; si-NC sense: 5'-UUCUC CGAAC GUGUC ACGU-3'; si-NC anti-sense, 5'-ACGUG ACACG UUCGG AGAA-3'.

qRT-PCR for Gene Expression

Trizol reagent kit was used to extract RNA following manual instruction. QuantiTect SYBR Green RT-PCR Kit was used to test gene expression by one-step qRT-PCR. In a 20 µL qRT-PCR system, there were 10 µL 2×QuantiTect SYBR Green RT-PCR Master Mix, 1.0 µL of forward and reverse primer (0.5 µM/L), 2 µg template RNA, 0.5 µL QuantiTect RT Mix, and ddH₂O. Primer sequences used were: miR-124_F: 5'-CGGTA AGGCA CGCGG TGA-3'; miR-124_R: 5'-AGTGC GA-ACT GTGGC GAT-3'; U6P_F: 5'-ATTGG AAC-GA TACAG AGAAG ATT-3'; U6P_R: 5'-GGAAC

GCTTC ACGAA TTTG-3'; STAT3_F: 5'-ATCAC GCCTT CTACA GACTG C-3'; STAT3_R: 5'-CA-TCC TGGAG ATTCT CTACC ACT-3'; Bcl-2_F: 5'-GGTGG GGTCA TGTGT GTGG-3'; Bcl-2_R: 5'-CGGTT CAGGT ACTCA GTCAT CC-3'; CyclinD1_F: 5'-CAATG ACCCC GCACG ATT-TC-3'; CyclinD1_R: 5'-CATGG AGGGC GGATT GGAA-3'; β-actin_F: 5'-GAACC CTAAG GC-CAA C-3'; β-actin_R: 5'-TGTCA CGCAC GATTT CC-3'. PCR conditions were: 95°C pre-denature for 15 min, followed by 40 cycles each containing 94°C denature for 15 s, 60°C annealing for 30 s, and 72°C elongation for 30 s. Gene expression was examined on ABI ViiA™ 7 fluorescent quantitative polymerase chain reaction (PCR) cycler.

Western Blot

Radioimmunoprecipitation assay (RIPA) lysis buffer was used to extract protein. A total of 40 μg samples was separated in sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) and was then transferred to polyvinylidene fluoride (PVDF) membrane, which was blocked in 5% de-fatted milk powder at room temperature incubation. Primary antibody (STAT3 at 1:300, p-S-STAT3 at 1:100, Bcl-2 at 1:200, Cyclin D1 at 1:200, and β-actin at 1:800) was added for 4°C overnight incubation. After phosphate buffered saline tween-20 (PBST) rinsing, horse radish peroxidase (HRP) conjugated secondary antibody (1:5000 dilution) was added for 60 min of incubation. The membrane was rinsed in phosphate buffered saline tween-20 (PBST) and incubated using Enhanced Chemiluminescence (ECL, Amersham Biosciences, Little Chalfont, UK) method. After dark exposure and development, the film was scanned and analyzed.

Clonal Formation Assay for Malignant Growth Potency

Cells from all transfected groups were inoculated into 6 cm diameter culture dishes at 100 cells density. Cells were incubated for 14-21 weeks. After that, cells were fixed in paraformaldehyde for 15 min and stained with Giemsa dye. Clones were counted under 100× magnification microscope. Clonal formation rate = (clone number/inoculate cell number) × 100%.

Transwell Assay for Cell Invasion Potency

Transwell assay was employed to test cell invasion potency. In brief, 1 × 10⁵ cells were inoculated into the upper chamber containing Matrigel and serum free Dulbecco's Modified Eagle Medium (DMEM). DMEM containing 10% fetal bovine

serum (FBS) was added to the bottom chamber. After 48 h, un-penetrated cells were removed. Chambers were then fixed in methanol and stained with 0.1% crystal violet. Cell number was counted under five randomly selected high-magnification fields.

Cell Apoptosis Assay

Cells were collected and digested with trypsin, and were re-suspended in binding buffer. 5 μL Annexin V-fluorescein isothiocyanate (FITC) and 5 μL propidium iodide staining buffer were sequentially added. Flow cytometry was used to test cell apoptosis.

PI Staining for Cell Cycle

Cells were digested with trypsin and rinsed in phosphate buffered saline (PBS). After 70% ethanol fixation overnight and phosphate buffered saline (PBS) washing, PI was added for staining in dark at 37°C for 30 min. Flow cytometry was used to detect cell cycle.

Statistical Analysis

SPSS16.0 software (SPSS Inc., Chicago, IL, USA) was used for data analysis. Measurement data were presented as mean ± standard deviation. Student *t*-test was used to compare measurement data between groups. Statistical significance was defined when *p* < 0.05.

Results

miR-124 Targeted and Inhibited STAT3 Expression

Bioinformatics analysis showed the existence of complementary binding sites between miR-124 and 3'-UTR of STAT3 mRNA (Figure 1A). Dual luciferase gene reporter assay showed that transfection of miR-124 mimic and miR-124 inhibitor remarkably decreased or increased relative luciferase activity in HEK293T cells, respectively (Figure 1B), suggesting that miR-124 could target 3'-UTR of STAT3 mRNA and inhibited its expression. qRT-PCR results showed that transfection of miR-124 mimic and miR-124 inhibitor remarkably decreased and potentiated STAT3 mRNA expression in DU145 cells, respectively (Figure 1C).

MiR-124 Down-Regulation and STAT3 up-Regulation in DU145 Cells

qRT-PCR results showed that, compared to RWPE-1 cells, DU145 cells had significant-

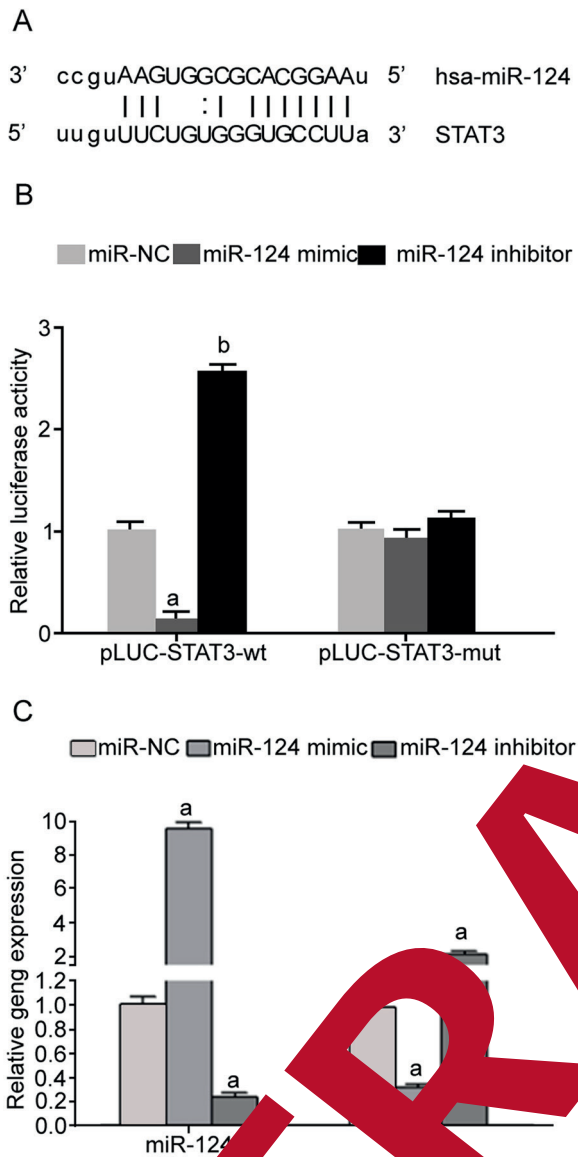


Figure 1. miR-124 up-regulated and inhibited STAT3 expression. (A) Functional site between miR-124 and 3'-UTR of STAT3 mRNA; (B) Dual luciferase gene reporter assay; (C) qRT-PCR for DU145 cell gene expression. a, $p < 0.05$ comparing between miR-124 mimic and miR-NC; b, $p < 0.05$ comparing between miR-124 inhibitor and miR-NC.

ly decreased miR-124 expression, plus lower expression of STAT3, Bcl-2, Cyclin D1 mRNA (Figure 2A). Flow cytometry results showed significantly higher protein expression of STAT3, p-STAT3, Bcl-2 and Cyclin D1 in DU145 cells compared to RWPE-1 cells (Figure 2B). Cell cycle test showed significantly higher S phase and G2/M phase ratio in DU145 cells than RWPE-1 cells, whilst G0/G1 phase ratio was lower in DU145 cells (Figure 2C). Flow cyto-

metry results showed significantly lower basal apoptotic rate of DU145 cells compared to RWPE-1 cells (Figure 2D).

MiR-124 up-Regulation Induced STAT3 Expression, Cell Invasion and Proliferation, Induced cell Apoptosis and Cycle Arrest

Transfection of miR-124 mimic and miR-124 inhibitor significantly increased expression of STAT3, p-STAT3, Bcl-2 and Cyclin D1 in DU145 cells (Figure 3A), weakened clonal formation ability (Figure 3B), decreased cell invasion potency (Figure 3C) and potentially induced cell apoptosis (Figure 3D) or G1 phase arrest (Figure 3E).

Discussion

The JAK-STAT signal transduction pathway can respond to multiple extracellular growth factors and cytokine stimulation. Under the existence of activator ligand, the JAK-STAT signal pathway, member receptor may undergo dimerization, which can further phosphorylate and activate JAK kinase to phosphorylate receptor tyrosine, facilitating the recruitment of STAT onto tyrosine phosphorylation site of receptor complex via SH2 domain. Under this scenario, JAK kinase can phosphorylate and activate STAT protein with spatial proximity, separating it from receptor complex to form dimer, which is transported from cytoplasm to nucleus, where it can facilitate transcription and expression of genes related to cell proliferation, cyclin, and apoptotic regulation¹⁶. STAT3 is the most important member of STAT protein family (STAT1, STAT2, STAT3, STAT4, STAT5a, STAT5b and STAT6). Previous studies attributed anti-apoptotic factor B-cell lymphoma-2 (Bcl-2)^{7,8} and Cyclin D1^{9,10} as important target genes under regulation by STAT3 transcriptional factor. By enhancing gene transcription and expression, STAT3 participates in facilitating cell proliferation and cycle progression, and modulating/antagonizing cell apoptosis, makes it one STAT protein with closest correlation with human tumor pathogenesis. A previous study¹⁷ showed elevated STAT3 expression in PCa patient tumor tissues, indicating its tumor-facilitating role in PCa. Reports^{14,15} also indicated significantly lower miR-124 expression in PCa tumor tissues/cells, indicating its possible role as tumor suppressor gene in PCa pathogenesis. Bioinformatics analysis showed complementary binding sites between miR-124 and 3'-UTR of STAT3. This work thus investigated if miR-124

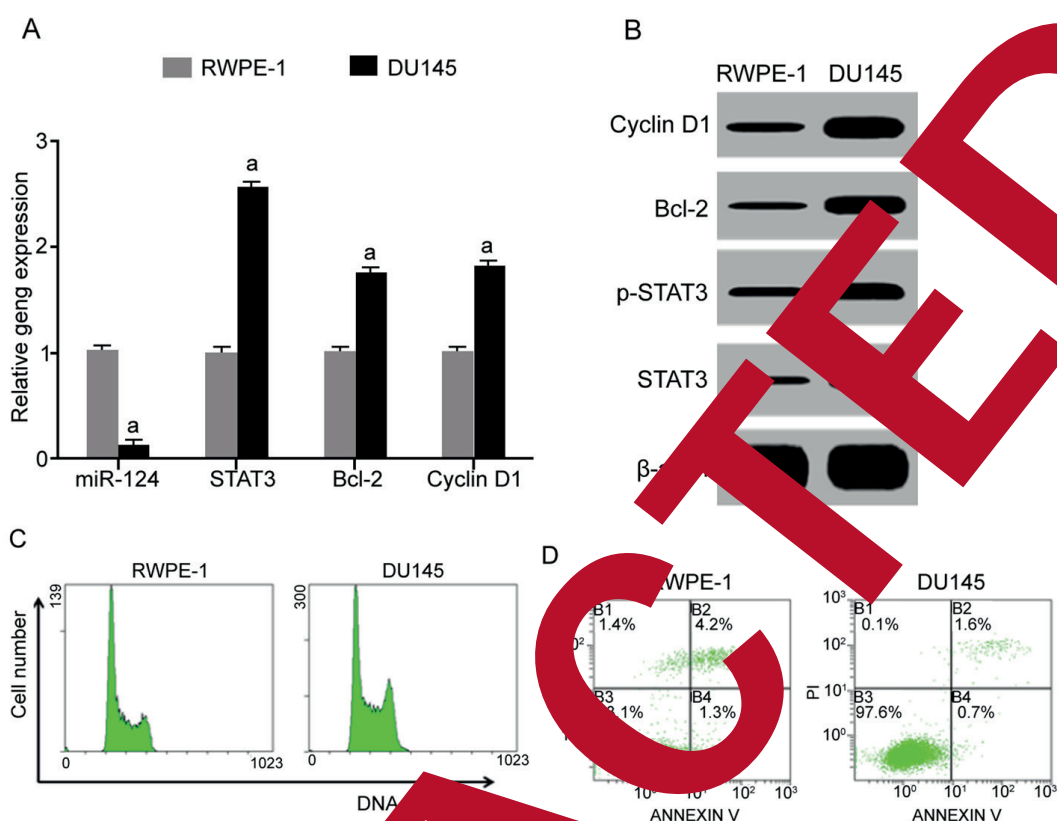


Figure 2. Higher miR-124 and lower STAT3 expression in DU145 cells. (A) RT-PCR for gene expression; (B) Western blot for protein expression; (C) Flow cytometry for cell cycle; (D) Flow cytometry for cell apoptosis. a, $p < 0.05$ comparing to RWPE-1 cells.

played a role in mediating STAT3 expression and affecting proliferation, cycle, apoptosis and invasion of PCa cells. Dual luciferase reporter assay showed that transfection of miR-124 significantly depressed relative luciferase activity, whilst miR-124 inhibitor elevated luciferase activity, indicating a targeted regulatory correlation between miR-124 and STAT3. Transfection of miR-124 mimic and miR-124 inhibitor remarkably increased and decreased STAT3 mRNA expression in DU145 cells, respectively, further demonstrating targeted regulation between miR-124 and STAT3. S/G1 phase ratio of DU145 cells was significantly lower than that of RWPE-1 cells, whilst the proportion of S phase and G2/M phase were significantly increased, indicating cell cycle progression. Flow cytometry results showed that comparing to RWPE-1 cells, DU145 cells had significantly lower cell cycle rate. Further assay found decreased miR-124 expression in DU145 cells, whilst expression of STAT3, p-STAT3, Bcl-2 and Cyclin D1 were elevated. These findings showed possible role of miR-124 down-regulation up-regulating

STAT3 and p-STAT3, facilitating downstream gene Bcl-2 and Cyclin D1 expression, and facilitating proliferation of PCa cells for suppressing their apoptosis. Chu et al¹⁸ showed hyper-methylation of miR-124 gene promoter region in PCa cells, leading to suppression of miR-124 expression. Shi et al¹⁵ found that comparing to prostate epithelium RWPE-1, PCa cell lines including 22Rv1, LNCaP, LAPC-4, cds2 and C4-2B, all had decreased miR-124 expression. Moreover, miR-124 expression level in PCa tissues was also lower than benign prostate tissue hyperplasia¹⁵. Falzarano et al¹⁹ observed significantly elevated miR-124 expression in PCa patients after treatment, indicating that miR-124 expression was one important mechanism governing PCa pathogenesis. In this study, miR-124 was down-regulated in prostate cells, indicating its role in PCa pathogenesis, as consistent with Chu et al¹⁸, Shi et al¹⁵, and Falzarano et al¹⁹. Abdulghani et al¹⁷ showed significantly elevated STAT3 expression in PCa tumor tissues, with higher expression level in those with bone or lung metastasis, indicating the correlation between

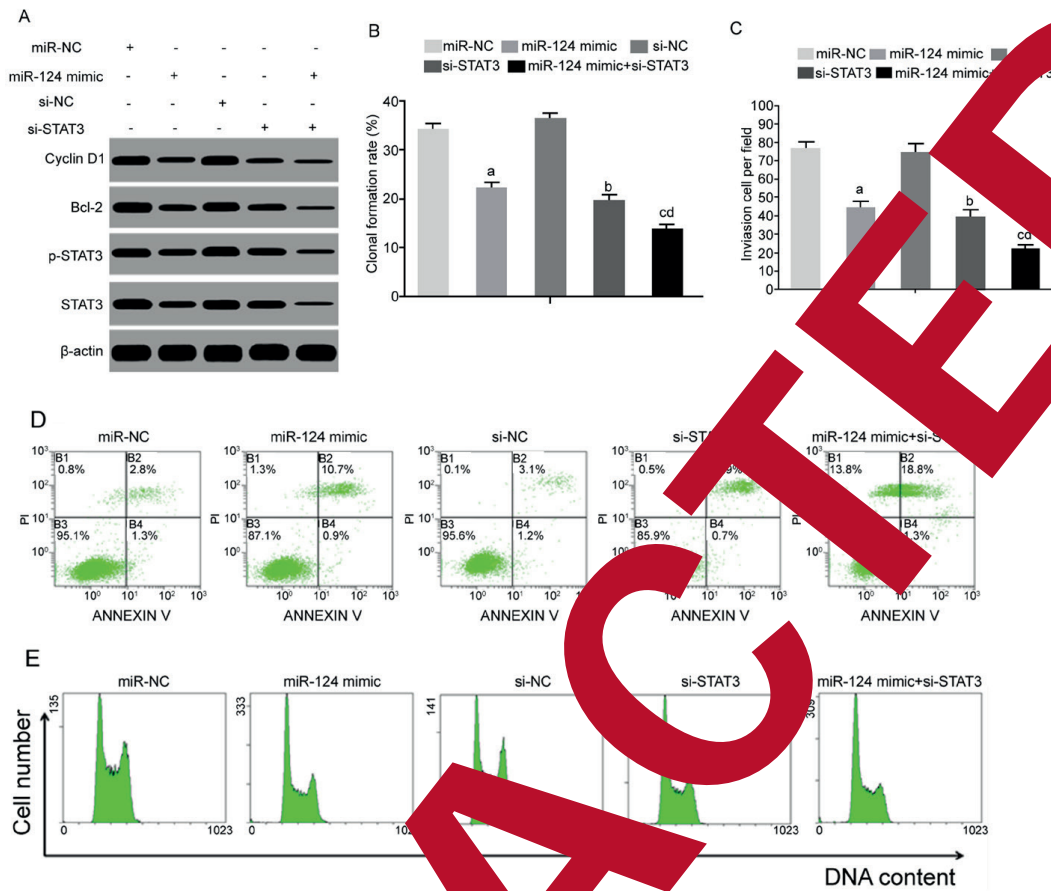


Figure 3. Elevated miR-124 expression inhibited STAT3 expression, cell invasion or proliferation, and induced cell apoptosis or cycle arrest. (A) Western blot for protein expression; (B) Clonal formation assay; (C) Transwell assay for cell invasion potency; (D) Flow cytometry for cell apoptosis; (E) Flow cytometry for cell cycle. a, $p < 0.05$ comparing between miR-124 and miR-NC group; b, $p < 0.05$ comparing between si-STAT3 and si-NC group; c, $p < 0.05$ comparing between miR-124 mimic + si-STAT3 and miR-NC group; d, $p < 0.05$ comparing between miR-124 mimic + si-STAT3 and si-NC group.

en STAT3 expression and both on and distal metastasis potency in Ca. In this study, STAT3 expression was significantly higher than normal human prostate epithelium, indicating the role of STAT3 up-regulation in PCa cell genesis, as similar with Abdulghani et al¹⁷. Further researches found that transfection of miR-124 mimic and/or si-STAT3 significantly down-regulated STAT3, p-STAT3 and downstream Bcl-2 and Cyclin D1 expression in DU145 cells, thus weakening cell proliferation and cell cycle progression, inducing cell cycle arrest at G0/G1 phase, and facilitating cell apoptosis. Shi et al¹⁵ found that over-expression of miR-124 remarkably weakened proliferation potency of PCa cells. Chu et al²⁴ showed that miR-124 up-regulation may suppress *in vitro* proliferation and invasion of PCa cells via targeted inhibition on androgen receptor

expression, and weakening the tumor formation potency in recipient animals. Kang et al²⁰ revealed that miR-124 could target and inhibit PACE4 expression to exert anti-cancer effects for antagonizing PCa cell proliferation. Qin et al²¹ showed that miR-124 up-regulation can inhibit PCa cell motility, migration and invasion via targeted inhibition on Slug gene expression to weaken the epithelial-mesenchymal transition (EMT) process. Shi et al²² found that over-expression of miR-124 may suppress PCa cell proliferation, and potentiate their drug sensitivity towards enzalutamide induction. Moreover, intravenous injection of miR-124 significantly inhibited *in vivo* growth of PCa tissues and facilitated tumor cell apoptosis²³. Abdulghani et al¹⁷ demonstrated that over-expression of STAT3 significantly enhanced *in vitro* DU145 cell motility and distal metastasis *in vivo*. After

activation of STAT3 by JAK kinase inhibitor, PCa cell had significantly weakened cell motility and migration potency. All these studies revealed the role of miR-124 up-regulation in weakening malignant biological features of PCa cells, as supported by our results. In this study, STAT3 down-regulation weakened proliferation or invasion of PCa cells, as consistent with Abdulghani et al¹⁷. This work revealed the role of miR-124 down-regulation in inducing STAT3 up-expression and in facilitating PCa occurrence, whilst miR-124 up-regulation could weaken PCa proliferation, invasion or apoptosis resistance via targeted inhibition of STAT3 expression. All these results have not been reported before.

Conclusions

MiR-124 up-regulation significantly decreases STAT3, p-STAT3 and downstream Bcl-2 or Cyclin D1 expression in DU145 cells, whose invasion and malignant proliferation potency are weakened, along with induction of cell cycle arrest at G0/G1 phase to facilitate cell apoptosis.

Acknowledgments

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Conflict of interest

The authors declare no conflict of interest.

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