

Protective effect of miR-146 against kidney injury in diabetic nephropathy rats through mediating the NF- κ B signaling pathway

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Abstract. – **OBJECTIVE:** To study the protective effect of micro ribonucleic acid (miR)-146 against kidney injury in diabetic nephropathy (DN) rats through the nuclear factor- κ B (NF- κ B) signaling pathway.

MATERIALS AND METHODS: In this experiment, 30 adult Sprague-Dawley rats with 5-6 weeks old and weighing 20-30 g were selected and randomly divided into control group (n=10), model group (n=10), and miR-146 Mimic group (n=10, DN rat model + miR-146 Mimic). The serum levels of creatinine (Cr) and blood urea nitrogen (BUN) in the three groups were determined using the full-automatic biochemical analyzer. The protein expression levels of phosphorylated-inhibitor of NF- κ B (p-I κ B), p-P65, p65, and Tubulin were detected *via* Western blotting. The messenger RNA (mRNA) of P65 was determined using quantitative real-time PCR (qPCR). Positive expression of p-I κ B in tissues was determined using immunohistochemistry. Moreover, the contents of inflammatory factors tumor necrosis factor- α (TNF- α), interleukin-1 β (IL-1 β) and IL-6 were detected using the enzyme-linked immunosorbent assay (ELISA) kits. Finally, the apoptosis was detected through Annexin V-fluorescein isothiocyanate (FITC) and propidium iodide (PI) dual-fluorescence labeling.

RESULTS: The serum levels of Cr and BUN were significantly higher in the model group than those in the control group ($p < 0.01$), while they were significantly lower in miR-146 Mimic group than those in the model group ($p < 0.05$). The levels of p-I κ B and p-P65/P65 significantly increased in the model group compared with those in the control group ($p < 0.01$), while they remarkably declined in the miR-146 Mimic group compared with those in the model group ($p < 0.05$). The results of qPCR showed that the mRNA level of P65 had no significant difference among the three groups ($p > 0.05$). The immunohistochemical assay showed that the positive expression of p-I κ B in tissues was consistent with those of the protein level as Western blotting revealed. The rats in the model group had

evidently increased levels of TNF- α , IL-1 β , and IL-6 compared with the control group ($p < 0.01$), while miR-146 Mimic group had evidently decreased levels of them compared with the model group ($p < 0.01$). Finally, apoptosis was enhanced in the model group compared with that in the control group, while it was remarkably inhibited in the miR-146 Mimic group.

CONCLUSION: MiR-146 can inhibit the NF- κ B signaling pathway, lower the levels of TNF- α , IL-1 β and IL-6, and reduce the apoptosis, thereby exerting protective effect against kidney injury in DN.

Diabetic nephropathy, MiR-146, NF- κ B signaling pathway.

Introduction

Diabetes mellitus often leads to microvascular injury, especially microvessels in kidneys, thus causing diabetic nephropathy (DN)¹. With the improvement of people's living standards in China, the incidence of diabetes mellitus also increases². The chronic development of diabetes mellitus is very harmful. Elevation of urinary protein, glomerular damage, and decline in the glomerular filtration rate are typical manifestations of diabetes mellitus and DN³. Therefore, it is extremely important to study the pathogenesis of DN and explore prevention approaches. Previous studies^{4,5} have shown that kidney injury in diabetic patients is mainly related to glucose metabolic disorders, oxidative stress, and inflammatory response. Nuclear factor- κ B (NF- κ B), an important nuclear transcription factor in cells, occupies a pivotal position in the inflammatory signaling pathway. When NF- κ B enters the nucleus, the transcription of a variety of cellular inflamma-

tory factors can be promoted, thereby mediating the intracellular inflammatory response through inflammatory factors⁶.

Besides the NF- κ B pathway, many micro ribonucleic acids (miRNAs) are also related to the inflammation. MiRNAs, generally with 19-22 nt in length, are a kind of endogenous non-coding single stranded small RNAs, which can regulate the transcriptional expressions of the target genes⁷. They bind to the 3'-untranslated region (3'UTR) of mRNAs to directly degrade them or inhibit their translation, thereby regulating the gene expressions at the transcriptional level. Important roles of miRNAs in various biological processes have been identified, such as cell differentiation, proliferation, and apoptosis⁸⁻¹¹.

MiR-146 is a widely studied miRNA currently, and its vital function in regulating the innate immunity has been discovered¹². MiR-146 includes miR-146a located in the second exon on chromosome 5 and miR-146b located on chromosome 10¹³. As a multifunctional miRNA, miR-146 can be involved in a variety of physiological and pathological processes, such as inflammation, immunity, occurrence, and development of tumor, by regulating the expression of multiple genes¹⁴⁻¹⁶.

Materials and Methods

Animal Modeling and Grouping

Wild-type Sprague-Dawley (SD) rat aged 5-6 weeks old (Shanghai BRK Laboratory Animal Co., Ltd., Shanghai, China) were housed in the specific pathogen-free animal room under the temperature of 25°C, humidity 55% and 12/12 h light/dark cycle, and they had free access to food and water. After habituation for 1 week, SD rats were randomly divided into control group, model group, and miR-146 Mimic group. In the model group, the rats were fed with high-glucose high-fat diets for 6 weeks and intraperitoneally injected with 60 mg/kg streptozotocin solution for modeling. The blood glucose level >16.7 mmol/L indicated the successful establishment of the DN model. In the miR-146 Mimic group, the DN model was first established in the same way of the model group. Then, DN rats were administrated with miR-146 Mimic. In the control group, the same amount of normal saline as that in the model group was injected. All animal operations were performed strictly according to the Guidelines for the Care and Use of Laboratory Animals of the

National Institute. This study was approved by the Animal Ethics Committee of Jilin University Animal Center.

Immunohistochemistry of Kidney Tissues

Bilateral kidneys of rats were taken out under anesthesia and washed clean with normal saline. The right kidney was immediately fixed with 10% neutral formalin solution, routinely prepared into paraffin sections, dewaxed, immersed in methanol containing 3%-60% H₂O₂ at room temperature for 30 min, and washed with phosphate-buffered saline (PBS) for 3 times. Membrane permeabilization was conducted with 0.1% Triton X 100 in PBS for 20 min. They were incubated with normal goat serum at room temperature for 20 min, rabbit anti-mouse NF- κ B p65 monoclonal antibody (1:200) in a refrigerator at 4°C overnight, and biotinylated goat anti-rabbit IgG secondary antibody at 37°C for 1 h. After washing with PBS for 3 times, the sections were incubated with horseradish peroxidase (HRP)-labeled streptavidin antibody at 37°C for 30 min. After DAB staining in the dark at room temperature, hematoxylin counterstaining was conducted for 30 min. The sections were dehydrated in ethanol, transparent with xylene and embedded with neutral balsam. Finally, the sections were observed under an inverted fluorescence microscope.

The dark brown particles in kidney tissues indicated positive expression. The mean optical density (OD) value of immunohistochemistry-positive particles was determined using ImageJ professional image analysis system. The protein level of phosphorylated-inhibitor of NF- κ B (p-I κ B) was semi-quantitatively analyzed.

Detection of NF- κ B Signaling Pathway in Kidney Tissues using Western Blotting

The kidney tissues of rats were cut into pieces, homogenized, and added with lysis buffer, followed by centrifugation at 20000 g and 4°C for 30 min. The total protein concentration was measured using the bicinchoninic acid (BCA) protein assay kit (Pierce, Rockford, IL, USA). After sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE), the protein was transferred onto polyvinylidene difluoride (PVDF) membranes (IPVH00010, Millipore, Billerica, MA, USA). The membranes were incubated with primary antibodies p-I κ B, p-P65, P65, and Tubulin (CST, Danvers, MA, USA) at 4°C overnight. After being washed, the membranes were

incubated with HRP-conjugated secondary antibodies (CST, Danvers, MA, USA) for 1 h. Finally, the enhanced chemiluminescence (ECL) mixture was added to obtain images using the fluorescence development technique.

Detection of mRNA Expression Level of P65 Via Quantitative Polymerase Chain Reaction (qPCR)

The mRNA was extracted from kidney tissues in each group using the TRIzol reagent (Invitrogen, Carlsbad, CA, USA), and reversely transcribed into complementary deoxyribose nucleic acid (cDNA) according to the instructions. 2 μ L of 5 \times PrimeScript RT Master Mix was added into 500 ng of RNA, and the total reaction system was 10 μ L. Then, PCR amplification was performed according to the instructions: 2 μ L of cDNA was added with 10 μ L of SYBR Premix Ex Taq II (Tli RNaseH Plus) (2 \times), 0.8 μ L of forward primers, 0.8 μ L of reverse primers, and 0.4 μ L of ROX Reference Dye II (50 \times), and deionized water was added finally till the total volume was 20 μ L. The mRNA expression level was calculated using the cycle threshold, with β -actin as an internal reference. The primer sequences were as follows: P65: F: 5'-CACCAAAGACCCACCTCACCT-3', R: 5'-CCGCATTCAAGTCATAGTCCC-3', β -actin: F: 5'-GCAGAAGGAGATTACTGCCCT-3', R: 5'-GCTGATCCACATCTGCTGC-3'.

Detection of Inflammatory Cytokines in Kidney Tissues

A total of 1-5 mg of kidney tissues were washed with PBS for 3 times, and then with 1-500 μ L of radioimmuno precipitation assay (RIPA) lysis buffer (Pierce and Warriner, Shanghai, China), and smashed using the homogenizer, followed by centrifugation at 3000 rpm and 4 $^{\circ}$ C for 10 min. Then, the supernatant was collected to detect the content of tumor necrosis factor- α (TNF- α), interleukin-1 β (IL-1 β), and IL-6 via enzyme-linked immunosorbent assay (ELISA), and quantified based on the protein concentration in the tissue supernatant.

Detection of Renal Function in Each Group

Serum levels of creatinine (Cr) and blood urea nitrogen (BUN) were detected for reflecting renal functions. 4 mL of blood was aseptically drawn from the caudal vein and centrifuged at 3000 g under low temperature for 10 min. The supernatant was collected and placed into the centrifuge

tube. Finally, the changes in serum indexes were detected using the full-automatic biochemical analyzer according to the instructions.

Detection of Apoptosis Via Flow Cytometry

The cells were suspended, and centrifuged at 1500 rpm for 5 min, and washed. The adherent cells were digested with trypsin containing ethylenediamine tetraacetic acid (EDTA) for an appropriate time, and the reaction was terminated with complete medium. Then, the cells were rinsed with PBS, and centrifuged at 1500 rpm for 5 min. 5×10^5 cells were collected, resuspended with 50 μ L of binding buffer, and incubated in 5 μ L of Annexin V-Light 650 and 10 μ L of propidium iodide (PI) at room temperature in dark place for 5-15 min. Flow cytometry was performed within 1 h, and the Annexin V-Light 650 fluorescence signal was detected through the FL4 channel, while the PI fluorescence signal was detected through the FL3 channel. The Annexin V-Light 650 single positive tube and PI single positive tube were detected simultaneously to determine the fluorescence compensation value and the position of the cross quadrant gate.

Statistical Analysis

GraphPad Prism 6.0 (La Jolla, CA, USA) was used for the statistical analysis of data. The data were expressed as ($\bar{x} \pm s$) and analyzed by *t*-test. $p < 0.05$ suggested the statistically significant difference.

Results

Expression of MiR-146 in Each Group

To observe the transfection efficiency of miR-146 Mimic, the expression level of miR-146 was detected. As shown in Figure 1, the expression of miR-146 was significantly upregulated in miR-146 Mimic group, while it significantly declined in the other two groups ($p < 0.05$), indicating that subsequent experiments can be performed.

Biochemical Indexes in Kidney Injury in Each Group

The renal function indexes in the three groups were detected using the conventional biochemical analyzer. As shown in Table I, the serum levels of Cr and BUN significantly decreased in the miR-

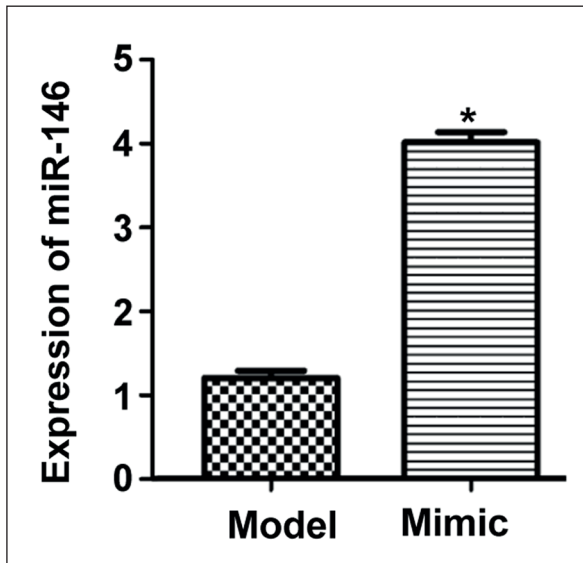


Figure 1. Expression of miR-146 in each group. Mimic: miR-146 Mimic group, * $p < 0.05$ vs. control group and model group.

146 Mimic group, while they were elevated in the model group ($p < 0.05$). It is suggested that the renal functions in DN rats were improved by miR-146.

Effect of MiR-146 on NF-κB Signaling Pathway

The protein was extracted from kidney tissue and detected via Western blotting. It was found that the levels of p-IκB and P65/P50 were remarkably upregulated in the model group compared with those in the control group ($p < 0.01$), while they significantly declined in the miR-146 Mimic group compared with those in the model group ($p < 0.05$). The results of qPCR showed that the mRNA level of miR-146 had no significant difference among the three groups (N.S.), and the results of an immunohistochemical assay for the positive expression of p-IκB in tissues were consistent with the protein level as Western blotting

revealed (Figure 2). The above findings demonstrated that the NF-κB signaling pathway could be activated in the model group by increasing the phosphorylation levels of IκB and P65. DN-induced activation of the NF-κB signaling pathway could be inhibited in the miR-146 Mimic group.

Effects of MiR-146 on Expression Levels of Inflammatory Factors TNF-α/IL-1β/IL-6

The expression levels of TNF-α/IL-1β/IL-6 in kidney tissues were further detected using the ELISA kits. The model group had evidently increased the levels of TNF-α/IL-1β and IL-6 compared with the control group ($p < 0.01$), while miR-146 Mimic group had evidently decreased their levels compared with the model group ($p < 0.01$) (Figure 3). The above results suggested that the expression levels of downstream cytokines TNF-α/IL-1β/IL-6 were markedly elevated in the model group after activation of the NF-κB signaling pathway. The overexpression of miR-146 suppressed the NF-κB signaling pathway and expressed the levels of these cytokines.

Effect of MiR-146 on Apoptosis of Kidney Tissues

After Annexin V-FITC and PI dual-fluorescence labeling for kidney tissues, apoptosis was detected using flow cytometry. It was found that the apoptosis was enhanced in the model group compared with that in the control group, while it was remarkably inhibited in the miR-146 Mimic group (Figure 4).

Discussion

MiRNAs are endogenous, non-coding, single-stranded, small RNA molecules generally with 19-22 nt in length^{17,18}, which can regulate the target gene expressions at the transcriptional level⁷. The maturation of miRNAs involves

Table 1. Changes of Content of Cr and BUN.

Group	Cr (μmol/L)	BUN (mmol/L)
Control group	21.01 ± 3.11	7.23 ± 1.21
Model group	91.25 ± 2.34 [#]	31.32 ± 3.14 [#]
miR-146 Mimic group	36.66 ± 5.32*	14.24 ± 1.33*

Note: The content of Cr and BUN is decreased significantly in miR-146 Mimic group, while it is the opposite in model group ($p < 0.05$). * $p < 0.05$ vs. model group, [#] $p < 0.05$ vs. control group.

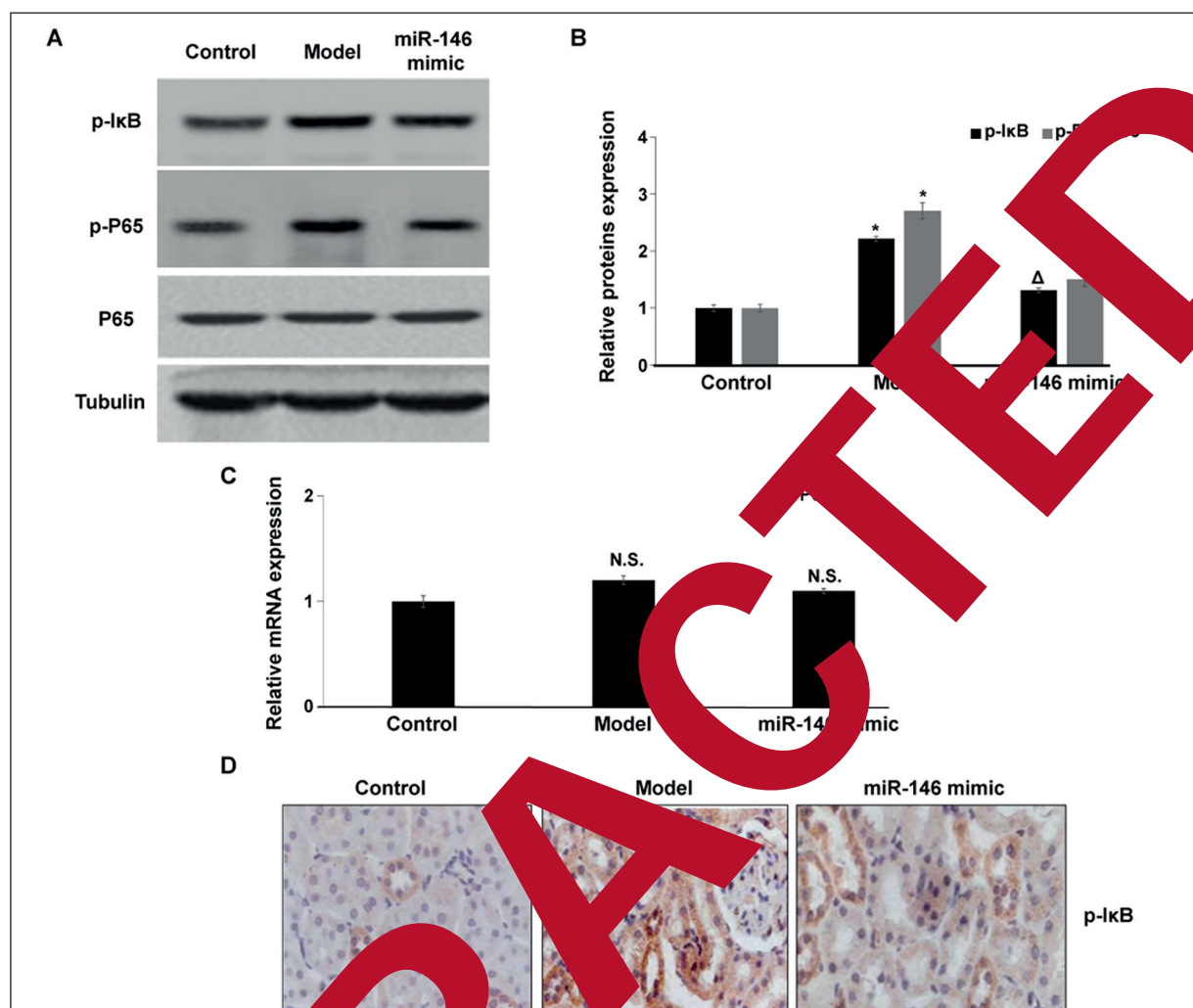


Figure 2. Effect of miR-146 on NF-κB signaling pathway. **A**, Protein levels of p-IκB, p-P65, P65 and Tubulin detected via Western blotting. **B**, Quantification of **A**. **C**, The mRNA level of P65 detected via qPCR. **D**, Positive expression of P-IκB in kidney tissues detected via immunohistochemistry (magnification ×40). * $p < 0.01$: model group vs. control group, $^{\Delta}p < 0.05$: miR-146 Mimic group vs. model group.

two processes. First, Drosha and DGCR8 proteins jointly process the nuclear pri-miRNA molecules, forming pre-miRNAs with 70 nt in length. Under the action of nuclear export protein Exportin5/RanGTP, pre-miRNAs are frequently transported from the nucleus to the cytoplasm, where they are further processed into mature miRNA molecules with about 21 nt in length under the assistance of Dicer. As a result, mature miRNA molecules form the RNA-induced silencing complex through the target gene mRNA, thereby inhibiting or degrading the expression of the target genes¹⁹.

MiR-146 is a widely studied miRNA currently, and it has been found to play an important role

in regulating the cellular innate immunity. The binding sequences between promoter regions of NF-κB and miR-146 exist. LPS/TNF-α could stimulate the upregulation of miR-146, and subsequently, the expression levels of two target genes IRAK1 and TRAF6 are downregulated²⁰, thereby inhibiting the immunoinflammatory process. Therefore, it is believed that miR-146 regulates the inflammatory signaling pathway through negative feedback. In addition, miR-146 is involved in pathophysiological processes, such as autoimmune diseases, rheumatoid arthritis, inflammation, and breast cancer^{21,22}.

NF-κB, an important nuclear transcription factor, occupies a pivotal position in the inflam-

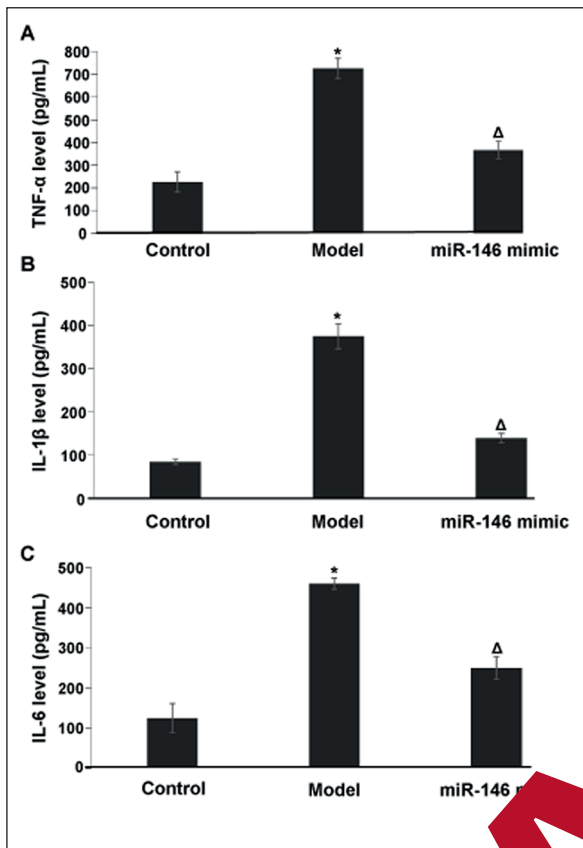


Figure 3. Effects of miR-146 on expression levels of inflammatory factors TNF- α /IL-1 β /IL-6. **A**, TNF- α level detected. **B**, IL-1 β level detected. **C**, IL-6 level detected. * p <0.01: model group vs. control group; ^Δ p <0.05: miR-146 Mimic group vs. model group.

inflammatory signaling pathway. When NF- κ B enters the nucleus, the transcription of a variety of cellular inflammatory factors can be promoted, thereby mediating the cellular inflammatory

response through inflammatory factors^{23,24}. Under the resting state, NF- κ B binds to I κ B, and its subunits P65 and P50 form the dimer in the cytoplasm. After exogenous stimuli, such as TNF- α , and ROS, I κ B is phosphorylated through signal transduction, and further degraded by the ubiquitinated proteasome, thereby releasing the NF- κ B-I κ B complex. The nuclear NF- κ B is activated, and the exposed P50 protein binds to specific sequences of target genes in the nucleus thereafter, leading to inflammatory responses by initiating the activity of the inflammatory factors²⁵⁻²⁶. As a result, acute kidney injury can be caused due to the NF- κ B-activated inflammatory response.

In the present study, SD rats were divided into control group, model group, and miR-146 Mimic group. First, the serum levels of kidney injury indicators Cr and BUN were detected. The serum levels of Cr and BUN significantly increased in the model group, while they remarkably declined in the miR-146 Mimic group compared with those in the model group, suggesting that the kidney injuries in DN rats could be alleviated by miR-146. Then, the effect of miR-146 on NF- κ B signaling pathway was analyzed in the three groups. The results showed that the levels of p-I κ B and p-P65 upregulated and the NF- κ B signaling pathway was activated in the model group, which were inhibited in the miR-146 Mimic group. However, the above treatments had no significant effect on the mRNA level of P65. In addition, p-I κ B level showed a similar trend. Furthermore, the effects of miR-146 on the expression levels of the downstream cytokines of NF- κ B signaling pathway were detected using ELISA. The results revealed that the model group had evidently increased levels of TNF- α ,

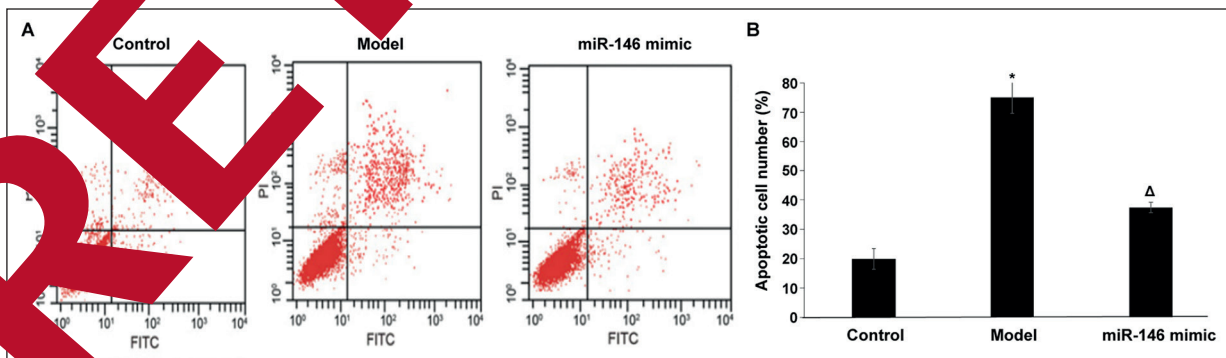


Figure 4. Effect of miR-146 on apoptosis of kidney tissues. **A**, Apoptosis determined using Annexin V-FITC and PI dual-fluorescence labeling and flow cytometry. **B**, Quantification of (A). * p <0.01: model group vs. control group, ^Δ p <0.05: miR-146 Mimic group vs. model group.

IL-1 β and IL-6, while miR-146 Mimic group had evidently decreased levels compared with the model group. Finally, apoptosis was enhanced in the model group compared with that in the control group, while it was remarkably inhibited in the miR-146 Mimic group.

Conclusions

In summary, miR-146 can inhibit the NF- κ B signaling pathway, lower the levels of TNF- α , IL-1 β , and IL-6, and reduce the apoptosis, thereby exerting a protective effect against kidney injury in DN.

Conflict of Interest

The Authors declare that they have no conflict of interests.

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