The protective role of microRNA-140-5p in synovial injury of rats with knee osteoarthritis via inactivating the TLR4/Myd88/NF-κB signaling pathway

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ABSTRACT

Objective: Recently, many studies have revealed the effect of microRNAs (miRNAs) in knee osteoarthritis (KOA). This study aims to explore the role of miR-140-5p in protective effects and mechanisms of synovial injury of rats with KOA via regulating the TLR4/Myd88/NF-κB signaling pathway.

Methods: The models of KOA Wistar rats were established by operation of anterior cruciate ligament transection. Rats were injected with agomir NC or miR-140-5p agomir. MiR-140-5p expression in KOA synovial tissues and synoviocytes was evaluated by reverse transcription quantitative polymerase chain reaction (RT-qPCR). The synoviocytes were transfected with mimics NC sequence and miR-140-5p mimics sequence. The expression of TLR4/Myd88/NF-κB signaling pathway-related proteins was measured by RT-qPCR and western blot analysis. The proliferation and apoptosis of synoviocytes in rats with KOA were evaluated by a string of experiments. The expression levels of inflammatory factors in KOA synovial tissues and synoviocytes were detected.

Results: MiR-140-5p was down-regulated in KOA synovial tissues and synoviocytes. Upregulation of miR-140-5p could inhibit the inflammation reaction and the apoptosis of synoviocytes as well as promote proliferation of synoviocytes of rats with KOA. Furthermore, upregulated miR-140-5p could inactivate the TLR4/Myd88/NF-κB signaling pathway in rats with KOA.

Conclusion: This study suggests that upregulated miR-140-5p could protect synovial injury by restraining inflammation reaction and apoptosis of synoviocytes in KOA rats via TLR4/Myd88/NF-κB signaling pathway inactivation.

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Introduction

Knee osteoarthritis (KOA) is a kind of prevalent joint disease, in which over 250 million people were involved around the world [1]. A study which was based on population revealed that the all-cause mortality of patients with KOA or hip OA was 55% higher than ordinary people [2]. According to the data of a recent research, the prevalence of KOA in Chinese was higher than that in Caucasians [3]. The risk factors of KOA have been demonstrated in previous studies, such as meniscectomy [4], obesity, physically demanding work and traumatic knee injury [5]. However, the early examination of OA is still an unresolved issue, which is of great significance [6]. A non-coding small RNA is called microRNA (miRNA), which generally contains about 20 nucleotides and has the ability to modulate some target genes [7]. In recent years, several miRNAs have been confirmed in human diseases, and some extant studies have unraveled that miRNAs were implicated in the progression of KOA, such as miR-9 [8] and miR-29a [9]. Nevertheless, there is little known about the correlation between miR-140-5p and KOA.

As one of the miRNAs, miR-140-5p has been proved to be related to several human diseases, including breast cancer [10], pulmonary arterial hypertension [11] and multiple sclerosis [12]. Besides, several studies have provided evidence that miR-140-5p was involved in OA [13–15]. The toll-like receptor 4s (TLR4) is one of the TLRs, which is of great importance in detecting the invasion of pathogens, and nuclear factor kappa-B (NF-κB) is a common transcription factor that was related to immune and inflammation [16]. According to available literature, the TLR4/Myd88/NF-κB signaling pathway was implicated in experimental traumatic brain injury and was testified to have the ability to relieve inflammatory...
injury [17]. There was another research unraveled that the TLR4/Myd88/NF-κB signaling pathway was involved in osteoporosis [18]. However, the relation among miR-140-5p, the TLR4/Myd88/NF-κB signaling pathway and KOA has not been studied yet. Thus, this study was designed to determine the role of miR-140-5p in protective effects and mechanisms of synovial injury of rats with KOA through the TLR4/Myd88/NF-κB signaling pathway, and we speculated that upregulated miR-140-5p could protect the synovial injury by restraining inflammation reaction and apoptosis of synoviocytes in rats with KOA via the TLR4/ Myd88/NF-κB signaling pathway.

Materials and methods

Ethics statement

Animal experiments were strictly in accordance with the Guide to the Management and Use of Laboratory Animals issued by the National Institutes of Health. The protocol of animal experiments was approved by the Institutional Animal Care and Use Committee of Xi’an Honghui Hospital, Yanliang District.

Study subjects

A total number of 55 healthy Wistar rats (aging 3 weeks, weighing 160–180 g) were obtained from Hunan SJA Laboratory Animal Co., Ltd. (Hunan, China). The rats were adaptively fed for 1 week for the following experiments, the temperature of feeding environment was 18–26°C, relative humidity was 40–70%, the noise was below 85 dB, ammonia concentration was below 22 PPM, and ventilated at 8–12 times/h.

Establishment of animal models and grouping

The Wistar rats were grouped into control group (15 rats) and experimental group (40 rats), the models of experimental group were established by operation of anterior cruciate ligament transection (ACLT). After weighed before the operation, the rats were injected by 0.1 mL/kg anesthetic, placed on the operating table, their patellae were rightly touched and inner 5 mm from patellae was conducted with longitudinal incision until the knee joint cavities were exposed, the tissues of rats in the control group were layer sutured, then the rats were raised in cages after they were completely awake. Rats in the experimental group were conducted with patella dislocation and flexion of knees, the anterior cruciate ligaments were divided and the medial meniscuses were excised, then the joint cavities were closed and the incisions were layer sutured. The rats were disinfected and raised in cages after awaking, without fixation of the limbs.

The 30 rats of successful models were grouped into three groups: the ACLT group (10 rats), the ACLT + agomir negative control (NC) group (10 rats) and the ACLT + miR-140-5p agomir group (10 rats), and 10 rats were set in the control group. Rats in the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group were injected in the knee joint cavities, respectively, with 0.5 mL agomir NC and miR-140-5p agomir (200 nM, purchased from Guangzhou RiboBio Co., Ltd., Guangdong, China) after 3 days, 1 week, 2 weeks, 3 weeks and 4 weeks, respectively. After 4 weeks from the operation, 10 rats were adopted from each group and euthanized, the knee joints were collected, part of which were conducted with X-Ray imaging analysis, and the other part were made into frozen sections and paraffin sections, with their total RNA and total proteins were extracted for histology and molecular biological detection.

Joint imaging assessment

The rats were weighed and injected with 1% pentobarbital sodium solution (Beijing Mairuida Technology Co., Ltd., Beijing, China), after muscular tension and conjunctival reflex were disappeared, the rats were fixed on the plates with their limbs stretched, then conducted with X-Ray photographs, the iconography data were collected.

Preparation and observation of frozen sections

The synovial tissues were fixed in 4% paraformaldehyde at 4°C overnight and rinsed by phosphate-buffered saline (PBS) 3 times, 15 min/time. The tissues were placed in 30% sucrose solution at 4°C overnight, embedded and sectioned by a freezing
microtome (Thermo Fisher Scientific Inc., Waltham, MA, USA) at a thickness of 35 μm, paved on slides treated with polylysine, and sealed by neutral resins. The agomir NC and miR-140-5p agomir injected in the rats were observed by a fluorescence microscope (Olympus Optical Co., Ltd, Tokyo, Japan).

**Hematoxylin-eosin (HE) staining**

The specimens were fixed by 10% formaldehyde, then embedded by paraffin and sectioned into 4 μm sections. The roasted sections successively dewaxed in xylene I and xylene II for 10 min; the dewaxed tissues were successively soaked in absolute ethanol I, absolute ethanol II, 95% ethanol, 80% ethanol and 70% ethanol, each for 2 min, and rinsed by PBS for 2 times, 5 min/time. Next, the tissues were stained with hematoxylin for 3 min, washed by tap water for 3 min, developed by 1% hydrochloric alcohol for 2 s, washed by tap water for 2 min, then successively soaked in 50%, 70% and 80% ethanol for 2 min and immersed in eosin for 5 s, and washed by tap water for 3 min. Afterward, the tissue sections were successively soaked in 90% ethanol, absolute ethanol I and absolute ethanol II for 3 min, and successively immersed in xylene I and xylene II for 5 min; sealed by neutral resins and conducted with microscopic examination. Quantitative analysis was carried on by Mankin score method.

**Electron microscopic observation**

The synovial tissues were fixed in 40 g/L glutaraldehyde for 1 h and washed by 0.1 mol/L PBS (pH 7.4) for 3 times, 5 min for each time, fixed by 10 g/L osmium tetroxide for 1.5 h and rinsed by 0.1 mol/L PBS (pH 7.4) for 3 times, 5 min for each time. Subsequently, the tissues were conducted with dehydration by gradient ethanol, soaked in the mixed liquor of acetone and equal Epon812 for 3 h, embedded by Epon812, and polymerized at 60°C for 48 h. The tissues were sectioned and stained by 40 g/L uranyl acetate for 20 min and 27 g/L lead nitrate for 20 min, then observed under an electron microscope (JEOL, Tokyo, Japan).

**Terminal deoxynucleotidyl transferase-mediated dUTP nick end-labeling (TUNEL) staining**

The paraffin-embedded sections were dewaxed and dehydrated, incubated by pepsase (0.25–0.5% hydrochloric acid solution) at 37°C for 25 min, added with 50 μL TUNEL reactive mixed solution (Beijing Zhongshan GoldenBridge Biotechnology Co. Ltd., Beijing, China) and incubated in a wet box at 37°C for 60 min. Subsequently, the sections were added with 50 μL peroxidase (POD) (Beijing Zhongshan GoldenBridge Biotechnology Co. Ltd., Beijing, China) and incubated in a wet box at 37°C for 30 min, then added with diaminobenzidine (DAB) agent and the color was observed by a microscope. The developing was stopped, and the sections were placed in hematoxylin for 2 min and washed for 2 min, dipped by 95% ethanol, absolute ethanol I-II for 3–5 min, xylene I-II for 3–5 min, and sealed in neutral resins. The results were analyzed under a light microscope. Fields of view in each group were randomly adopted under a high power lens (× 400) and the number of apoptotic cells, as well as the total cells, were counted (graded by double-blinded method).

**Enzyme-linked immunosorbent assay (ELISA)**

Blood from arteria cruralis of anesthetic rats was placed for 1 h and centrifuged, then the serum samples were collected, the synovial tissues were ground into homogenate and centrifuged, the supernatant of each group was collected and sub-packaged in aseptic Eppendorf (EP) tubes. Eight standard substances were prepared according to the construction of ELISA kit of IL-1β and TNF-α (Raybiotech, GA, USA), and the 8th well was the blank control group. The standard substances and samples (100 μL) were, respectively, added into 96-well plates, incubated at 37°C for 2 h, added with 100 μL primary antibody and incubated for 1 h. Each well was added with 100 μL secondary antibody, then each well was added with 100 μL chromogenic reagent and incubated for 30 min. Every well was added with 50 μL stop solution to stop the reaction. The absorbance value and concentration of each well were measured, and the standard curve was graphed.
Reverse transcription quantitative polymerase chain reaction (RT-qPCR)

The total RNA was extracted from specimens and cells by Trizol kit (Invitrogen, Carlsbad, CA, USA). The primers were designed by Takara Biotechnology Co., Ltd. (Osaka, Japan) (Table 1). The RNA was reversely transcripted into cDNA according to the internal reference of TLR4, Myd88, NF-κB, β-actin, U6, and TNF-α. The relative transcriptional levels of miRNA were calculated by 2^(-ΔΔCt) method [19].

Western blot analysis

The total proteins of synovial tissues and cells were extracted. The protein concentration of each sample was evaluated and adjusted by deionized water, insuring the loading quantities were in accordance. The samples and loading buffer were mixed up and boiled at 100°C for 5 min, electrophoresis separation was conducted, the proteins were transferred onto the nitrocellulose membrane, which was then sealed by 5% nonfat dry milk at 4°C overnight. The proteins were added with primary antibodies: TLR4 (Abcam, Cambridge, UK, 2 μg/mL), Myd88 (diluted at 1: 1000, Cell Signaling Technology, MA, USA), NF-κB p65 (Abcam, Cambridge, UK, 0.5 μg/mL), Bcl-2 and Bax (diluted at 1: 500, both from Proteintech, CHI, USA), β-actin (diluted at 1:1000, Santa Cruz Biotechnology, Santa Cruz, CA, USA), incubated overnight and washed by PBS for 3 times, 5 min/time. The proteins were added with IgG secondary antibody (1: 1000, Wuhan Boster Biological Technology Co., Ltd., Hubei, China) which was marked by horseradish peroxidase (HRP), and incubated at 37°C for 1 h. The membrane was soaked into enhanced chemiluminescent (ECL) reaction reagent (Pierce, Rockford, IL, USA) for 1 min. The results were observed after exposure, development and fixation. β-actin was taken as internal reference, and the protein maker was obtained from Piercenet (#854,785), the Western blot image was analyzed by ImageJ2x software (National Institutes of Health, MD, USA).

Cell isolation, culture and appraisal

The synovial tissues separated from normal rats and rats with KOA were soaked in PBS containing double-antibody for 5 min, washed by PBS for 3 times and placed in the culture dishes. The tissues were cut into small pieces and trypsinized by 6 mL collagenase II (4 mg/mL) containing 10% FBS for 3 h, centrifuged with the supernatant discarded, then added with 2 mL Dulbecco’s modified Eagle medium (DMEM) culture solution and centrifuged with the supernatant discarded, added with 4 mL DMEM solution and mixed up, then added with 1 mL PBS (20% final concentration) and incubated with the adherent cells discarded. The cells were trypsinized and passaged by 0.25% trypsin after the primary cells were formed into monolayer, cells in the logarithmic growth phase were adopted for the experiment. The growth of the cells was observed under an inverted microscope. Purity identification of synoviocytes: the third passage cells were seeded on the 24-well plates (1 × 10⁵ cells/well), the medium was

Table 1. Primer sequence.

| Gene          | Primer sequence (5’-3’)                           |
|---------------|---------------------------------------------------|
| miR-140-5p    | F: 5'-AACAACGCGGCGCCGCGCCGGA- 3'                 |
| U6            | R: 5'-CTCACTGGTGTCGGGAA- 3'                       |
| TLR4          | F: 5'-CTGGCTCCGCGGAGCACA- 3'                      |
| Myd88         | R: 5'-AACGCTCAGAATGTGGGA- 3'                      |
| NF-κB         | F: 5'-ACAAAACGGGGAAATTTTCTG- 3'                  |
| Bcl-2         | R: 5'-TGCGGACACACACAACTTTAGC- 3'                 |
| Bax           | F: 5'-TTGCCAGGGGCAATTTGAG- 3'                    |
| IL-1β         | R: 5'-ACACGTGAGTGGCAACTTTG- 3'                   |
| TNF-α         | F: 5'-TGACTGACATCCGTCCTTTTCAAG- 3'               |
| β-actin       | R: 5'-TCACAGATCAAATGCGGTT- 3'                    |

Note: F, forward; R, reverse; miR-140-5p, microRNA-140-5p; TLR4, toll-like receptor 4; NF-κB, nuclear factor kappa-B; IL-1β, interleukin (IL)-1β; TNF-α, tumor necrosis factor-α.
discarded after the cells covered the wells, and the cells were fixed with goat blood for 1 h, incubated by vascular cell adhesion molecule (VCAM)-1 (1:500, Abcam Inc., Cambridge, UK) at 4°C overnight, then the cells were rinsed by PBS for 3 times, incubated by fluorescence secondary antibody for 1 h, stained by 4',6-diamidino-2-phenylindole 2 hci (DAPI) for 10 min and observed under a fluorescence microscope, the results were preserved.

**Cell grouping and transfection**

The synoviocytes were grouped into the normal group (normal synoviocytes without any treatment), the ACLT group (KOA synoviocytes without any transfection), the ACLT + mimics NC group (KA synoviocytes transfected with mimics NC), the ACLT + miR-140-5p mimics group (KOA synoviocytes transfected with miR-140-5p mimics). The cells were transfected with miR-140-5p mimics and mimics NC (Shanghai GenePharma Co., Ltd., Shanghai, China) according to the instructions of Lipofectamine 2000 kit (Invitrogen Inc., Carlsbad, CA, USA), then the transfected cell was incubated at 37°C and 5% CO₂. The culture solution was changed after 24 h; the transfection efficiency was observed under an inverted fluorescence microscope.

**Colony formation assay**

The fourth passage monolayer cells of each group in the logarithmic growth phase were trypsinized into single cell by 0.25% trypsin with the single cell rate was above 95%, the cell concentration was adjusted using DMEM containing 10% FBS. The cells were seeded into 6-well plates at 100 cells/well and incubated in 5% CO₂ at 37°C for 2–3 weeks. The colony formation rate was calculated as the number of colonies/the number of seeded cells × 100%.

**3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2′-h-tetrazolium bromide (MTT) assay**

The synoviocytes in each group were detached by trypsin and the cell concentration was adjusted to 1 × 10⁵ cells/mL by DMEM, then the cells were seeded onto 96-well plates at 200 μL/well, and incubated at 37°C and 5% CO₂ in an incubator for 24 h, the medium was discarded after the cells became adherent. Next, each well was appended with 20 μL MTT solution (5 μg/mL, Sigma-Aldrich Chemical Company, St Louis, MO, USA) for the 4-h incubation. With supernatant discarded, each well was supplemented with dimethyl sulfoxide (Sigma-Aldrich Chemical Company, St Louis, MO, USA) at 200 μL/well. After the crystals were fully dissolved, the optical density (OD₄₉₀) value was analyzed by a microplate reader (Thermo Fisher Scientific Inc., Waltham, MA, USA).

**Hoechst 33,342 staining**

The cells of each group were added with 2 mL DMEM containing 10% FBS (both from Gibco, Carlsbad, CA, USA), and added with Hoechst 33,342 dye (5 μg/mL, Biolab technology Co., Ltd., Beijing, China), incubated at 37°C without light exposure for 90 min, then added with 1 mL DMEM containing 10% FBS and photographed under a fluorescence microscope.

**Flow cytometry**

The cell apoptosis was measured by AnnexinV-APC/propidium iodide (PI) double staining: cells in each group were centrifuged at 1000 rpm and 4°C for 5 min and collected. Binding buffer was 4 times diluted by deionized water (4 mL binding buffer+12 mL deionized water). Subsequently, the cells were washed by precooled PBS twice, and centrifuged at 1000 rpm for 5 min, with supernatant discarded, the cells were resuspended by 250 μL binding buffer, and the cell concentration was adjusted to 1 × 10⁶ cells/well; cell suspension (5 mL) was added into 5 mL flow tubes, then was added with 5 μL Annexin V-APC and 5 μL PI solution (both from BD Biosciences, Franklin Lakes, NJ, USA) and incubated without light exposure for 15 min. PBS (400 μL) was added into the reaction tubes, which was then injected in flow cytometry, the results were analyzed by a computer.
Statistical analysis

All data analyses were conducted using SPSS 21.0 software (IBM-SPSS, Inc, Chicago, IL, USA). The measurement data conforming to the normal distribution were expressed as mean ± standard deviation. One-way analysis of variance (ANOVA) was used for comparisons among multiple groups. Tukey’s multiple comparisons test was used for pairwise comparisons after the ANOVA. *p value < 0.05 was indicative of a statistically significant difference.

Results

MiR-140-5p was down-regulated in synovial tissues of rats with KOA

According to the results of RT-qPCR (Figure 1a), in comparison to the control group, the expression of miR-140-5p was declined in the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group (all P < 0.05). miR-140-5p expression in the ACLT + miR-140-5p agomir group was elevated contrasted to the ACLT group (P < 0.05). There was no evident difference in the miR-140-5p expression between the ACLT group and the ACLT + agomir NC group (P > 0.05).

The outcomes of fluorescence microscopy observation of frozen sections implied that there were many green fluorescence could be observed in the synovial tissues of the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group (Figure 1b), indicating that agomir NC and miR-140-5p agomir have been successfully injected into the rats.

Upregulated miR-140-5p inhibits KOA development in synovial tissues of rats

The images of rats’ knees under the X-Ray are shown in Figure 2a: there was no obvious osteoporosis in the control group, the joint space was obvious and no bone destruction could be found. Compared with the control group, rats in the ACLT group and the ACLT + agomir NC group were with local hyperostosis of knee joint, changes of alignment and narrowing of joint space, which suggested an evident KOA. Relative to the ACLT group, rats in the ACLT + miR-140-5p group were with local sclerotin reduction and clear joint space of knees, indicating a decline of KOA level.

The outcomes of HE staining unraveled that (Figure 2b): under a light microscope, there were regular cellular morphology of joint synoviocytes and neat arrangement in the control group; while there were disorder arrangement, partial necrosis and exfoliation, tissue edema and interstitial osteoporosis of synoviocytes in the ACLT group and the ACLT + agomir NC group. With complete tissue structures, the synoviocytes edges were slightly irregular in the ACLT + miR-140-5p agomir group. Mankin score illustrated that in contrast to the control group, the Mankin scores of the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group were observably heightened.

Figure 1. MiR-140-5p is down-regulated in synovial tissues of rats with KOA. a, the expression of miR-140-5p in synovial tissues was detected by RT-qPCR; b, the results of fluorescence microscopy observation (400 ×), * P < 0.05 vs the control group, # P < 0.05 vs the ACLT group. N = 10, the measurement data conforming to the normal distribution were expressed as mean ± standard deviation. ANOVA was used for comparisons among multiple groups. Tukey’s multiple comparisons test was used for pairwise comparisons after ANOVA.
The Mankin score of the ACLT + miR-140-5p agomir group was apparently reduced, which was relative to the ACLT group ($P < 0.05$). No evident difference could be found in the Mankin score between the ACLT group and the ACLT + agomir NC group ($P > 0.05$).

The outcomes of electron microscope observation suggested that Figure 2c there were complete cytomembrane, less cytoplasm, many collagen tissues and no obvious vascular proliferation of the rats' joint synoviocytes in the control group; while there were cellular atrophy and apparent accumulation of dense fibrous substances of synoviocytes in the ACLT group and the ACLT + agomir NC group ($P > 0.05$).

According to the outcomes of RT-qPCR and Western blot analysis (Figure 3b-d), the expression of Bcl-2 in the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group was decreased, while the expression of Bax was heightened in contrast to the control group (all $P < 0.05$). The expression of Bcl-2 in the ACLT + miR-140-5p agomir group was enhanced and the expression of Bax was reduced relative to the ACLT group (all $P < 0.05$).

### Upregulated miR-140-5p suppresses synoviocytes apoptosis in synovial tissues of rats with KOA

The results of TUNEL staining suggested that Figure 3a the synoviocytes apoptosis in synovial tissues was accelerated in the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group, which was compared with the control group ($P < 0.05$). In comparison to the ACLT group, the synoviocytes apoptosis in synovial tissues in the ACLT + miR-140-5p agomir group was attenuated ($P < 0.05$).

### Upregulated miR-140-5p represses inflammation reaction of rats with KOA

Results of ELISA and RT-qPCR revealed that Figure 4a-b the expression levels of IL-1β and...
TNF-α of rats’ serum and synovial tissues in the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group were elevated compared with the control group (all \( P < 0.05 \)). The expression levels of IL-1β and TNF-α in the ACLT + miR-140-5p agomir group were considerably reduced relative to the ACLT group (all \( P < 0.05 \)).

**Figure 3.** Upregulated miR-140-5p suppresses synoviocytes apoptosis in synovial tissues of rats with KOA. a, results of TUNEL staining (400 ×); b, the mRNA expression of Bax and Bcl-2 was detected by RT-qPCR; c, protein bands of Bax and Bcl-2; d, the statistical results of expression of Bax and Bcl-2 in the synovial tissues in each group by Western blot analysis, * \( P < 0.05 \) vs the control group, # \( P < 0.05 \) vs the ACLT group. N = 10, * \( P < 0.05 \) vs the control group, # \( P < 0.05 \) vs the ACLT group. The measurement data conforming to the normal distribution were expressed as mean ± standard deviation. ANOVA was used for comparisons among multiple groups. Tukey’s multiple comparisons test was used for pairwise comparisons after ANOVA.

**Figure 4.** Upregulated miR-140-5p represses inflammation reaction of rats with KOA. a, the expression of IL-1β and TNF-α in the serum and synovial tissues of each group was detected by ELISA; b, the expression of IL-1β and TNF-α of synovial tissues was detected by RT-qPCR; * \( P < 0.05 \) vs the control group, # \( P < 0.05 \) vs the ACLT group. N = 10, * \( P < 0.05 \) vs the control group, # \( P < 0.05 \) vs the ACLT group. The measurement data conforming to the normal distribution were expressed as mean ± standard deviation. ANOVA was used for comparisons among multiple groups. Tukey’s multiple comparisons test was used for pairwise comparisons after ANOVA.

Upregulated miR-140-5p inhibits activation of TLR4/Myd88/NF-κB pathway in synovial tissues of rats with KOA

According to the outcomes of RT-qPCR and western blot analysis (Figure 5a-c), the expression of TLR4, Myd88 and NF-κB of rats’ synovial tissues in the ACLT group, the ACLT + agomir NC group and the ACLT + miR-140-5p agomir group was elevated,
which was compared with the control group (all $P < 0.05$). The expression of TLR4, Myd88 and NF-κB of rats’ synovial tissues in the ACLT + miR-140-5p agomir group was lowered relative to the ACLT group (all $P < 0.05$).

**The cultured cells were rats’ synovioblasts**

As shown in Figure 6a, synovioblasts became adherent and extended growth protuberances after 24 h, spindle dendritic growth of cells and thin or short protoplasmic protuberances could be observed under the microscope, which were of characteristics of fibroblasts; cells were clustered after 72-h culture; cells were connected into a whole piece under a microscope after 120-h culture.

Purity appraisal of synovioblasts (Figure 6b): VCAM-1 was the symbolic protein of synovioblasts, which was mainly expressed in the cytoplasm. The third passage cells were constructed with VCAM-1 immunofluorescence staining, more than 98% of fibroblasts expressed VCAM-1 positively and all distributed in the cytoplasm, suggesting that the cultured cells were rats’ synovioblasts with extremely high purity, which were met the demands of following experiments.

**MiR-140-5p was down-regulated in synoviocytes of rats with KOA**

According to the results of RT-qPCR (Figure 6c), in comparison to the normal group, the expression of miR-140-5p was declined in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group (all $P < 0.05$). MiR-140-5p expression in the ACLT + miR-140-5p mimics group was elevated contrasted to the ACLT group ($P < 0.05$). There was no evident difference in the miR-140-5p expression between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).

**Upregulated miR-140-5p induces proliferation of synoviocytes of KOA rats**

The colony formation ability of synoviocytes in each group was detected by colony formation assay (Figure 7a-b), outcomes of which unraveled that the colony formation ability of synoviocytes in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was evidently reduced in comparison to that of the normal group (all $P < 0.05$), the colony formation ability of synoviocytes in the ACLT + miR-140-5p mimics group was considerably heightened relative to the ACLT group ($P < 0.05$). There was no obvious difference in colony formation ability...
between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).

The proliferation ability of synoviocytes in each group was measured by MTT assay (Figure 7c), the results suggested that the proliferation ability of synoviocytes in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was obviously lowered in contrast to that of the
normal group, the proliferation ability of synoviocytes in the ACLT + miR-140-5p mimics group was apparently advanced contrasted to the ACLT group ($P < 0.05$). There was no evident difference in proliferation ability between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).

**Upregulated miR-140-5p inhibits apoptosis of synoviocytes of rats with KOA**

Cell apoptosis was evaluated using Hoechst 33,342 staining (Figure 8a), the results of which implied that the number of apoptotic cells in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was observably enhanced in comparison to the normal group (all $P < 0.05$); while the number of apoptotic cells in the ACLT + miR-140-5p mimics group was decreased relative to the ACLT group ($P < 0.05$). The outcomes of AnnexinV-APC/PI double staining (Figure 8b) suggested that the apoptosis rate of synoviocytes in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was evidently elevated compared with the normal group (all $P < 0.05$), while the apoptosis rate of synoviocytes in the ACLT + miR-140-5p mimics group was lowered relative to the ACLT group ($P < 0.05$). There was no apparent difference in the apoptosis rate of synoviocytes between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).

The results of RT-qPCR and western blot analysis (Figure 8c-e) indicated that the expression of Bcl-2 in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was decreased, while the expression of Bax was heightened in contrast to the normal group (all $P < 0.05$). The expression of Bcl-2 in the ACLT + miR-140-5p mimics group was enhanced and the expression of Bax was reduced relative to the ACLT group (both $P < 0.05$). There was no obvious difference in the expression of Bcl-2 and Bax between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).
Upregulated miR-140-5p represses inflammation reaction of synoviocytes in rats with KOA

Results of ELISA and RT-qPCR revealed that (Figure 9a-c) the expression levels of IL-1β and TNF-α of synoviocytes in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was elevated compared with the normal group (all \( P < 0.05 \)). The expression levels of IL-1β and TNF-α in the ACLT + miR-140-5p mimics group was considerably reduced relative to the ACLT group (both \( P < 0.05 \)). There was no evident difference in the expression levels of IL-1β and TNF-α between the ACLT group and the ACLT + mimics NC group (\( P > 0.05 \)).

Upregulated miR-140-5p represses activation of TLR4/Myd88/NF-κB pathway in synoviocytes of rats with KOA

According to the outcomes of RT-qPCR and western blot analysis (Figure 10a-c), the expression of TLR4, Myd88 and NF-κB of synoviocytes in the ACLT group, the ACLT + mimics NC group and the ACLT + miR-140-5p mimics group was elevated, which was compared with the normal group (all \( P < 0.05 \)); while the expression of TLR4, Myd88 and NF-κB of synoviocytes in the ACLT + miR-140-5p mimics group was lowered relative to the ACLT group (all \( P < 0.05 \)). No apparent difference could be found in the expression of TLR4, Myd88 and NF-κB of synoviocytes.
between the ACLT group and the ACLT + mimics NC group ($P > 0.05$).

**Discussion**

OA is the commonest chronic joint disease which usually accompanied by arthralgia as well as disability [20]. It has been testified that miRNAs, which were characterized as small non-coding RNAs, played a role of leading molecules in the RNA silencing [21]. Moreover, there were several recent studies have unraveled that miR-140-5p was implicated in some human diseases, such as ovarian cancer [22] and non-small cell lung cancer [23]. Nevertheless, there is little known about miR-140-5p and the TLR4/Myd88/NF-κB signaling pathway in KOA. Hence, this study was focused on the impacts of miR-140-5p as well as the TLR4/Myd88/NF-κB signaling pathway on KOA, and we have found from the results of this research that upregulation of miR-140-5p could protect synovial injury by inhibiting inflammation reaction and apoptosis of synoviocytes with KOA through the TLR4/Myd88/NF-κB signaling pathway inactivation.

One of the findings of our study illustrated that miR-140-5p was poorly expressed in KOA synovial tissues and synoviocytes. Similar to this result, Hao Yang et al. have assessed in their study that miR-140-5p performed a poor expression in hepatocellular carcinoma tissues [24]. As for miR-140-5p expression in cells, it has been demonstrated that the expression...
of miR-140-5p was down-regulated in the CD4+ T cells of patients with multiple sclerosis [12]. Another essential result of this study is that upregulation of miR-140-5p could evidently inhibit the expression of inflammation-related factors, such as IL-1β and TNF-α in KOA synovial tissues and synoviocytes, indicating an attenuation of inflammation in the rats with KOA. It is consistent with this result that a recent study has provided evidence to prove that the overexpression of miR-140-5p was able to relieve lipopolysaccharide-induced human intervertebral disc inflammation and degeneration by restraining the expression of inflammation cytokines, such as TLR4, TNF-α, IL-1β and IL-6 [25], which has contributed to the role of miR-140-5p in inflammation of joint diseases. What’s more, we have also found that the upregulation of miR-140-5p was able to inhibit the activation of the TLR4/Myd88/NF-kB signaling pathway in KOA synovial tissues and synoviocytes of rats. Similarly, the relation between upregulated miR-140-5p and the TLR4/Myd88/NF-kB signaling pathway has been identified in a present study that overexpression of miR-140-5p could suppress the TLR4/Myd88/NF-kB signaling pathway [26].

The biological function of miR-140-5p has been unraveled in this research that the upregulation of miR-140-5p could promote the proliferation of synoviocytes of KOA rats. In line with this outcome, Xin et al. have revealed that overexpressed miR-140-5p could elevate the proliferation of human dental pulp stem cells [27]. The effects of miR-140-5p have been unearthed in another existing literature, in which the authors illustrated that miR-140-5p could suppress the proliferation of gastric cancer by modulating YES1 [28]. Besides, we have demonstrated in this study that the upregulation of miR-140-5p has the capacity of inhibiting the apoptosis of synoviocytes in rats with KOA. Similar to this result, a recent research has identified that the overexpression of miR-140-5p could decelerate the apoptosis of human primary chondrocytes by regulating fucosyltransferase 1 [14], which has also confirmed the role of miR-140-5p in OA. Except for that, Zhang et al. have also reported in their study that the overexpression of miR-140-5p could induce the cell apoptosis in colorectal cancer [29]. All of these studies have contributed to proving the mechanism and function of miR-140-5p and the TLR4/Myd88/NF-kB signaling pathway in human diseases.

To sum up, our study provides evidence that the upregulation of miR-140-5p could alleviate the inflammation reaction of rats with KOA. Furthermore, the upregulation of miR-140-5p has the ability to suppress the apoptosis and induce the proliferation of synoviocytes, resulting in a protective impact on the synovium of KOA rats. These outcomes would pave a new way of KOA therapy. However, more efforts such as enlarge the experimental specimens are needed to be carried on to further clarify the function mechanisms of miR-140-5p in the development of KOA.

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Availability of data and material

Not applicable

Authors’ contributions

Guarantor of integrity of the entire study: Xiaoqiang Huang
Study design: Feng Qiao
Experimental studies: Peng Xue
Manuscript editing: Feng Qiao, Peng Xue
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Consent for publication

Not applicable

Disclosure statement

No potential conflict of interest was reported by the authors.

Ethical statement

Animal experiments were strictly in accordance with the Guide to the Management and Use of Laboratory Animals issued by the National Institutes of Health. The protocol of animal experiments was approved by the Institutional Animal Care and Use Committee of Xi’an Honghui Hospital, Yanliang District.

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