Abstract. The aim of the present study was to assess the expression of microRNA (miRNA)-4784 in the chondrocytes of early osteoarthritis (OA) and to determine the effect of double-stranded (ds)-miRNA-4784 transfection on chondrocyte function. Following the construction of an OA rabbit model, normal chondrocytes (normal control group), OA chondrocytes obtained 4 weeks after modeling (OA at week 4 group) and 8 weeks after modeling (OA at week 8 group) were used. The relative expression of miRNA-4784 in each group was detected using reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Western blotting was performed to measure the expression of type II collagen (Col2a1) and matrix metalloproteinase (MMP)-3 in each group with or without ds-miRNA-4784 transfection. The results revealed that the levels of miR-4784 in groups OA at week 4 and 8 were significantly lower than that of normal control group (P<0.05). It was also demonstrated that Col2a1 mRNA expression levels in groups OA at week 4 and 8 were 49 and 38% of that in the normal control group, respectively. Furthermore, MMP-3 mRNA expression levels increased by 3.12- and 3.95-fold in groups OA at week 4 and 8, respectively, compared with those in the normal control group (P<0.01). Following transfection with ds-miRNA-4784, Col2a1 mRNA expression levels increased by 63 and 126% compared with the levels prior to treatment in groups OA at week 4 and 8, respectively (P<0.01). The expression levels of MMP-3 mRNA in groups OA at week 4 and 8 decreased following transfection compared with the levels prior to treatment. Col2a1 and MMP-3 protein expression exhibited similar patterns to the mRNA expression. In summary, the results of the present study suggest that miRNA-4784 expression is significantly reduced in early stage OA chondrocytes. Transfection with ds-miRNA-4784 promotes the expression of Col2a1 and inhibits the MMP-3 expression in chondrocytes.

Introduction

Osteoarthritis (OA) is a disease characterized by progressive chondrocyte degeneration, synovial hyperplasia in joints, narrowing of the joint space and dysregulation of extracellular matrix metabolism (1). OA primarily affects the knee, hip and shoulder joints and the main symptoms include joint pain, swelling, joint deformity and limited mobility (2). The pathogenesis of OA is associated with aging, obesity, inflammation, immunity, genetics and many other factors (3). Cartilage degeneration is considered to be one of the primary pathological changes that cause OA, and this typically occurs due to metabolic disorders of extracellular matrix synthesis and degradation (4). Chondrocytes, as the primary cell type found in cartilage tissue, serve important roles in the maintenance of bone and joint structure and function (5,6). Previous studies have demonstrated that matrix metalloproteinases (MMPs) and collagen also serve an important role in the development of OA (7,8). MMPs are ion-active proteases that are ubiquitous in the cartilage and function to degenerate chondrocytes and degrade the extracellular matrix in the cartilage (9-11). MicroRNAs (miRNAs) are single-stranded non-coding RNAs that are associated with the development and progression of OA (12). It has been reported that miRNAs serve a role in the pathogenesis of OA regulating the expression of inflammatory mediators, vascular endothelial growth factor and nerve growth factor (12). Further studies have revealed that miRNAs may inhibit or promote the expression of MMPs and collagen, resulting in the degeneration of chondrocytes and cartilage extracellular matrix, eventually leading to OA (13,14).

Previous studies have also reported that miRNA-4784 expression is downregulated in certain types of cancer, including breast and liver cancer, and miRNA-4784 is involved in the development and progression of cancer by affecting the Akt signal pathway (12,13). To the best of our knowledge, no previous studies have assessed the expression and mechanism of action of miRNA-4784 in OA. The aim of the present study was to assess the expression of miRNA-4784 in OA chondrocytes and the effect of transfection with exogenous.
double-stranded (ds)-miRNA-4784 on the chondrocyte function. The results of the present study may provide a theoretical basis for a novel treatment of OA.

Materials and methods

Experimental reagents. Lipofectamine® 2000, Dulbecco's modified Eagle's medium (DMEM), PBS and penicillin were purchased from Gibco (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA). The miR-4784 mimic was purchased from GenePharma Co., Ltd. (Shanghai, China). Fetal bovine serum and western blotting reagents (cat no. YJ1012465) were purchased from Yiji (Shanghai, China) and radioimmuno-precipitation cell lysis solutions from Beyotime Institute of Biotechnology (Haimen, China). Cell culture plates, Col2a1 and MMP-3 rat anti-rabbit monoclonal antibodies were purchased from Pierce (cat nos. ab185430 and ab26111; Pierce; Abcam, Shanghai, China). The quantitative polymerase chain reaction (qPCR) kit and the reverse transcription (RT) kit were purchased from Fermentas (Thermo Fisher Scientific, Inc.).

Experimental animals and OA model establishment. A total of 40 New Zealand rabbits (age, 6 months; 20 male and 20 female; weight, 2.0-3.0 kg) were purchased from the West China Center of Medical Sciences, Sichuan University (Chengdu, China). The rabbits were raised with controlled temperature and light cycles (24°C and 12/12 light cycles) and free access to water and food. The humidity was 60±10% for 2 weeks prior to surgery. The following method was used to establish the adult rabbit OA model: pentobarbital sodium (3%; 30 mg/kg) was injected into the ear vein for anesthesia, and then the rabbits were fixed on an operating table in the supine position. The right knee was shaved for anesthesia, and then the rabbits were fixed on an operating table in the supine position. The right knee was shaved and disinfected with iodophor (Abcam). The medial patella meniscus was opened and the patella was everted. The knee joint was then buckled and the anterior segment ligament and collateral ligament were cut with an ophthalmic clip. The surgical site was rinsed, the joint capsule was closed and the skin was disinfected. The right limb remained unfixed and rabbits were kept in a separate case, and had free access to water and food until they were sacrificed. A daily intramuscular injection of 80×10⁶ U of penicillin was administered for 1 week. Right knee tissues were collected at 4 and 8 weeks following modeling to serve as groups OA at week 4 and 8, respectively (each, n=10). The left knee of rabbit without treatment was set as control group (n=10). The present study was approved by the Ethics Committee of the 174th Hospital of Chinese PLA (Chenggong Hospital Affiliated to Medical College of Xiamen University, Xiamen, China).

Cell culture. Rabbits were sacrificed at week 4 and 8 following surgery in groups OA at week 4 and 8, respectively. Aseptic cartilage specimens were obtained under sterile conditions. Cartilage tissues with a thickness of 1-2 mm were scraped using a surgical blade and digested with 0.25% trypsin for 30 min at 37°C. Tissues were then digested with 0.2% collagen II protease for 2 h at 37°C. This procedure was repeated 2-3 times to collect chondrocytes, which were subsequently cultured in DMEM supplemented with 10% fetal bovine serum and 200 U/ml penicillin at 37°C and 5% CO₂.

Detection of miRNA expression using RT-qPCR. U6 snRNA was used as an endogenous control to quantify the expression of miRNA-4784. The expression of Col2a1 and MMP-3 was quantified using β-actin as an endogenous control before and after transfection. Total RNA was extracted from OA chondrocytes using TRIzol (Thermo Fisher Scientific, Inc.) and was reverse transcribed into cDNA using the RT kit according to the manufacturer's instructions (37°C for 15 min, 95°C for 5 min). PCR thermocycling conditions were as follows: 94°C for 2 min, followed by 50 cycles of 94°C for 30 sec, 55°C for 30 sec and 72°C for 30 sec. Cq values were processed by 2ΔΔCq method and the relative expression of each gene was normalized to their corresponding endogenous controls (15). U6 forward, 5'-TGC GGG TCG TGG CTT CCG CAG CAG-3' and reverse, 5'-CCA GTG CAG GTG CCG AGG-3'; β-actin forward, 5'-GCT GCCGTGTTGCCCTCTAGG-3' and reverse, 5'-ACGCAGGATGGCATGGAGGGA-3'; Col2a1 forward, 5'-TCC TAA GGG TGC CAA TGG TGA-3' and reverse, 5'-AGG ACC AAC TTT GCA TAA CGG ATC-3'; MMP-3 forward, 5'-ATTCCATGAGCCAGGCTTC-3' and reverse, 5'-CATTTGGGTCAAACTCCAACGTG-3'.

Detection of Col2a1 and MMP-3 protein expression using western blotting (cat. no. YJ1012465; Yiji). Cultured chondrocytes were rinsed with PBS and lysed using cell lysis solution for 30 min. Cell lysate was then transferred into a tube and centrifuged at 9,800 g for 20 min at 4°C to collect cell supernatant. Total protein concentration was determined using a BCA assay. Protein samples (~90 µg per lane) were separated by 10% SDS-PAGE and transferred onto polyvinylidene membranes. The membranes were then blocked using 5% bovine serum albumin (Abcam) at 24°C for 2 h. Membranes were washed 3 times with TBST and incubated with Col2a1 anti-rabbit monoclonal antibodies (1:500) or MMP-3 rat anti-rabbit monoclonal antibodies (1:500) overnight at 4°C. After further washes with TBST 3 times, the membranes were incubated with goat anti-rabbit horseradish peroxidase-labeled secondary polyclonal antibody (1:1,000; cat. no. 7077; Cell Signaling Technology, Danvers, MA, USA) for 1 h at room temperature. Membranes were washed 3 more times with TBST and ECL reagent (Beyotime, Shanghai, China) was added for signal development. The relative expression level of each protein was normalized to the endogenous control (β-actin; 1:1,000; cat. no. 60008-1-Ig, Proteintech, Wuhan, China) using ImageJ 1.48 (NIH, Bethesda, MD, USA).

Statistical analysis. SPSS 13.0 software (SPSS, Inc., Chicago, IL, USA) was used for all statistical analyses and data are
expressed as the mean ± standard error of the mean. One-way analysis of variance was used for multiple group comparisons and a post-hoc Dunnett’s test was performed. P<0.05 was considered to indicate a statistically significant difference.

**Results**

miR-4784 expression. The expression of miR-4784 in OA articular chondrocytes gradually decreased with disease progression. miR-4784 expression in groups OA at week 4 and 8 was 61 (P<0.05) and 54% (P<0.01) of that in the group normal control, respectively (Fig. 1). However, no significant difference in miR-4784 expression was observed between groups OA at week 4 and 8 (Fig. 1).

Expression of Col2a1 mRNA in chondrocytes prior to and following ds-miRNA-4784 transfection. Col2a1 mRNA expression in groups OA at week 4 and 8 prior to transfection were 49 and 38% of that in normal control group, respectively (P<0.01; Fig. 2). However, no significant difference was observed between groups OA at week 4 and 8. Following ds-miRNA-4784 transfection (ds-miRNA-4784+), no significant change in Col2a1 mRNA expression was observed in normal control group. However, Col2a1 mRNA levels in groups OA at week 4 and 8 were increased by 63 and 126%, respectively, following transfection (P<0.01; Fig. 2). Furthermore, no significant difference in Col2a1 expression was identified between groups OA at week 4 and 8 following transfection.

Col2a1 protein expression in chondrocytes prior to and following ds-miRNA-4784 transfection. No significant

![Graph](image1.png)

**Figure 1.** miRNA-4784 expression in normal chondrocytes and OA model chondrocytes at week 4 and 8 after modeling. Data are presented as the mean ± standard error of the mean. n=10. *P<0.01 and **P<0.05. miRNA, microRNA; OA, osteoarthritis.

![Graph](image2.png)

**Figure 2.** Col2a1 mRNA expression in chondrocytes. Shown are normal chondrocytes and chondrocytes at week 4 and 8 after the establishment of the OA model, respectively. + indicates chondrocytes following transfection. Data are presented as the mean ± standard error of the mean. n=10. *P<0.01. OA, osteoarthritis.

![Graph](image3.png)

**Figure 3.** MMP-3 and Col2a1 protein expressions were assessed in chondrocytes using western blotting. + indicates chondrocytes following transfection. (A) MMP-3 and Col2a1 protein expressions using western blotting. (B) The expression of Col2a1 protein increased at week 4 and 8 after transfection (*P<0.01). (C) The expression of MMP-3 protein decreased at week 4 and 8 after transfection (**P<0.01). Data are presented as the mean ± standard error of the mean. n=10. *P<0.01. MMP, metalloproteinase; OA, osteoarthritis.
difference in Col2a1 protein expression was observed in normal control group prior to and following ds-miRNA-4784 transfection. However, Col2a1 protein expression increased significantly in groups OA at week 4 and 8 following transfection compared with pre-transfection levels (P<0.01; Fig. 3A and C).

**MMP-3 protein expression in chondrocytes prior to and following ds-miRNA-4784 transfection.** No significant difference in MMP-3 protein expression was observed in normal control group prior to and following ds-miRNA-4784 transfection. However, the expression of MMP-3 protein significantly decreased following transfection compared with pre-transfection levels in groups OA at week 4 and 8 (P<0.01; Fig. 3A and B).

**MMP-3 mRNA expression in chondrocytes prior to and following ds-miRNA-4784 transfection.** The expression of MMP-3 mRNA in groups OA at week 4 and 8 prior to transfection were 3.12 and 3.95-fold higher than that in normal control group (P<0.01, Fig. 4). However, no significant difference was observed between groups OA at week 4 and 8. Following transfection with ds-miRNA-4784, no significant difference in MMP-3 mRNA expression was observed in normal control group compared with pre-transfection level. Furthermore, the expression of MMP-3 mRNA in groups OA at week 4 and 8 was significantly decreased following transfection compared with pre-transfection levels (P<0.01; Fig. 4). No significant difference in MMP-3 mRNA expression was observed between groups OA at week 4 and 8 following transfection.

**Discussion**

OA is one of the most common types of chronic arthritis and seriously affects the quality of life in the elderly population (1-3). Major pathological features of OA include a series of biological and/or morphological changes, such as the degeneration of articular cartilage, hyperosteoegeny and/or sclerosis (16). Currently, OA is typically treated conservatively, with the main aim being the pain relief; however, effective radical treatment remains insufficient (17). Furthermore, certain patients may experience disease recurrence due to ineffective treatment (18).

The occurrence of OA is associated with many parameters, including genetic and immune factors; however, the exact pathogenesis remains unclear (3). The roles of various miRNAs in OA chondrocytes have been widely studied and have provided a novel research direction, which has helped to elucidate the pathogenesis of OA (13). miRNAs are a group of ubiquitous endogenous small single-stranded non-coding RNAs that are associated with the regulation of 30% of genes (19). The abnormal expression of various miRNAs, including miRNA-140, -126 and -146, has been observed in patients with OA (13,14). These miRNAs advance the development of OA by promoting the expression of MMPs and reducing the mechanism of collagen (20).

In the present study, a rabbit OA model was established to assess the changes in miRNA-4784 expression in chondrocytes at 4 and 8 weeks following the model construction. The results revealed that the expression of miRNA-4784 in chondrocytes gradually decreased with prolonged duration of the disease, indicating that miRNA-4784 is downregulated in OA chondrocytes. However, the mechanism by which this occurs remains unclear. Previous studies have revealed that the expression of a series of miRNAs, including miRNA-140, decrease in chondrocytes during early OA; this change may be associated with miRNA inhibition by cytokines such as metalloproteinases or collagen (13,20).

Changes in the expression of Col2a1 and MMP-3 in OA chondrocytes were also assessed in the present study. Col2a1 expression serves an important role in maintaining the normal function of chondrocytes, the downregulation of which indicates the degeneration of articular cartilage (21-23). MMP-3 is an important chondroitin-degrading enzyme whose upregulation promotes the degeneration of articular cartilage (24). In the present study, compared with normal control group, the expression of Col2a1 in OA chondrocytes was significantly reduced at the miRNA and protein levels, which is consistent with the results of previous studies (25-27). Furthermore, an increase in MMP-3 expression was observed in OA chondrocytes, indicating that there was a degenerative change in the cartilage during early stage OA. Additionally, miRNA-4784 levels increased following transfection with exogenous ds-miRNA-4784, while Col2a1 expression also increased in OA chondrocytes. The expression of MMP-3 mRNA and protein were decreased, suggesting that ds-miRNA-4784 transfection improves the function of OA chondrocytes.

The results of the present study also demonstrated that miRNA-4784 expression gradually decreases with prolonged disease duration. Transfection-induced miRNA-4784 upregulation maintained the stability of chondrocytes, promoted the expression of Col2a1 and inhibited the expression of MMP-3. These results confirm that miRNA-4784 serves a role in the development and progression of OA. In this investigation, we only studied the mechanism of miRNA-4784 in the pathogenesis of OA from experimental animals, but we did not detect the cytokines and RNA in clinical patients. Therefore, more research is needed to further confirm the role of miRNA-4784 in the pathogenesis of OA. The present study may provide novel insights into the pathogenesis of OA and a theoretical basis for the application of gene therapy in future treatments for OA.
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Availability of data and materials
The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions
JL and QY were major contributors in writing the manuscript and participated in the analysis and discussion of the data. JL and YYa were responsible for the cell culture. XC and YYe performed RT-qPCR and western blotting. All authors read and approved the final version of the manuscript.

Ethics approval and consent to participate
The present study was approved by the Ethics Committee of the 174th Hospital of Chinese PLA (Chenggong Hospital Affiliated to Medical College of Xiamen University, Xiamen, China).

Patient consent for publication
Not applicable.

Competing interests
The authors declare that they have no competing interests.

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