

Alterations of autophagy in knee cartilage by treatment with treadmill exercise in a rat osteoarthritis model

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Abstract. The aim of the present study was to investigate potential alterations in the articular cartilage in a rat model of monosodium iodoacetate (MIA)-induced osteoarthritis (OA) with or without treatment with moderate treadmill exercise. A total of 30 male Sprague-Dawley rats were randomly divided into three groups (n=10), including the control, OA and OA with treadmill exercise (OAE) groups. Rats were evaluated upon completing the treadmill exercise program (speed, 18 m/min; 30 min/day; 5 days/week for 4 weeks). Interleukin (IL)-1 β and IL-4 levels in the serum and intra-articular lavage fluid (IALF) were measured by ELISA. Alterations in articular cartilage and synovium were also evaluated by histology, immunohistochemistry, western blotting and reverse transcription-quantitative polymerase chain reaction. The results revealed that IL-1 β in the serum and IALF decreased in the OAE group, whereas IL-4 increased, and histological evaluation indicated that the OAE group had a clear treatment response. However, the expression of type II collagen in the articular cartilage increased in the OAE group as compared with the OA group, whereas ADAMTS5 expression decreased. In contrast to light chain 3B (LC3B), the protein expression levels of BECLIN1 and sequestosome 1 (SQSTM1) were increased in the OA group. In addition, a significant increase was observed between OA and OAE groups in LC3B and SQSTM1 protein levels, whereas no change was observed in BECLIN1 levels between the OA and OAE groups in the superficial and deep zones. The results of western blotting demonstrated that LC3II was notably decreased in the OA group and partially increased in the OAE group. The mRNA expression levels of LC3B and SQSTM1 increased in the OA and OAE groups, with a significant difference observed

between the two groups, while a concomitant decrease was detected in BECLIN1 levels. In conclusion, 30 min of treadmill exercise had an evident protective effect in the articular cartilage of rats with MIA-induced OA and may promote autophagy in the articular cartilage.

Introduction

Osteoarthritis (OA) is a chronic joint disease characterized by the progressive degradation of articular cartilage, accompanied by remodeling of the subchondral bone and consequent articular dysfunction. OA affects 20-25% of individuals aged >45 years and 37.4% of individuals aged \geq 60 years old (1,2). Knee OA may cause severe pain, physical disability and reduced quality of life. Furthermore, OA is one of the major reasons for knee arthroplasty and leads to high healthcare costs (3,4).

A variety of risk factors contribute to OA, including age, gender, obesity, history of knee injury, genetic factors and nutritional factors (5). Research in the field of molecular biology has identified an imbalance in the homeostasis of the cartilage matrix during OA, as well as inflammation caused by various biomarkers, such as prostaglandins, adipokines and cytokines among others (6).

Moderate physical exercise is recognized to be an effective therapeutic treatment for knee OA (7). Research revealed that a moderate level of exercise is beneficial in rat knee OA models, and affects subchondral or trabecular bone metabolism (8-10). Various cytokines, growth factors and proteins secreted by osteoclasts and osteoblasts of OA subchondral bone cysts aggravate the disease and are alleviated by moderate physical exercise (11-14).

Autophagy, as an essential cellular homeostatic mechanism in eukaryotic cells, involves continuous biosynthesis and renewal of cellular organelles and macromolecules. It is recognized that autophagy serves an important role in the knee OA process, in which it has been demonstrated that multiple signal transduction pathways may be involved (15). Previous studies reported that autophagy in chondrocytes mainly serves as a protective response of cells to negative stimulation, such as hypoxia (16), glucose starvation or nutrient insufficiency (17), aging (18) and certain inflammatory mediators (19-21). Nevertheless, complex results have been derived from *in vivo* studies (18,19,22). Autophagy may not only be a protective

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or homeostatic mechanism in normal or OA cartilage, but may also be an essential process for the differentiation and maturation of chondrocytes, which occurs constantly during the pathological process of OA (23).

A previous study reported that physical exercise affects autophagy in numerous tissues and organs, including muscle, adipose, heart and brain tissues (24). The present study was designed to detect the alterations in autophagy in the cartilage following treadmill exercise using a rat OA model that was induced by monosodium iodoacetate (MIA) injection. The results may help to develop a novel theory regarding the treatment effect of exercise on OA.

Materials and methods

Experimental animals. The experimental protocol was approved by the Ethics Committee of Shengjing Hospital, China Medical University (Shenyang, China). A total of 30 male Sprague-Dawley rats (230 ± 10 g; 8 weeks old; specific-pathogen-free) were purchased from HFK Bioscience Co., Ltd. (Beijing, China) and kept in plastic cages with sawdust bedding in a controlled environment ($22 \pm 2^\circ\text{C}$; 70% humidity) under a 12:12-h light/dark cycle with the lights turned on at 06:00 a.m. Animals were able to move freely in the cages, were fed a standard diet and had access to tap water *ad libitum*. The body weight of rats was recorded weekly prior to the treadmill exercise each Monday. Adaptive exercise was applied to all animals at a speed of 10 m/min for 10 min/day prior to regular treadmill exercise to eliminate stress.

OA model and treadmill running protocols. The animals were randomly divided into three groups ($n=10$ per group), as follows: Control (sterile saline injection only), OA (MIA injection) and OA with treadmill exercise (OAE). An intra-articular injection of MIA was used in both knees following the protocol described in our previous study (25), in order to generate the OA model according to the study of Guzman *et al.* (26). All rats were kept sedentary in the cages; however, the OAE group began treadmill exercise at 24 h post-injection with a speed of 18 m/min for 30 min/day (moderate exercise) for 5 days/week, which is considered to be beneficial for cartilage damage caused by MIA injection. The treatment schedule is shown in Fig. 1A.

Sampling and tissue preparation. After 4 weeks of exercise, all animals were anesthetized 2 h subsequent to the last exercise session. Blood samples (3 ml) were obtained immediately and centrifuged at $3,000 \times g$ for 10 min in 4°C to obtain the serum. All rats were euthanized, and the joints of the left knee were separated and fixed in 4% paraformaldehyde solution at room temperature. Intra-articular lavage fluid (IALF) was obtained from the synovial cavity of the right knee by injection and recovery with 0.2 ml PBS three times using a 1-ml syringe. Cartilage was scraped off from the weight-bearing area of the condyles of the right femur and tibia using a scalpel, and was carefully collected. The left knee joints were kept at room temperature, while other samples were kept at -80°C .

Histological analysis. Following 2 weeks of soaking in 4% paraformaldehyde solution, the left knee joints were washed

with PBS five times (10 min/wash) and transferred to a 10% EDTA solution (EDTA-2Na, dissolved in 0.1 M PBS) to decalcify the samples. Subsequently, samples were kept at a constant temperature of 37°C for 1 month and the solution was renewed every 3 days. Dehydration in an ethanol series and embedding in paraffin was performed successively following the decalcification, and serial $4.5\text{-}\mu\text{m}$ sagittal sections were then cut for histological examination. Hematoxylin and eosin (H&E) staining and toluidine blue staining were used to obtain scores according to the Osteoarthritis Research Society International (OARSI) and the modified Mankin scoring systems, respectively, in order to grade and stage OA development in cartilage (27). Synovitis scoring, which reflects the inflammation state in the synovium (28), was also performed by H&E staining.

ELISA of IALF and serum levels. IL-1 β and IL-4 levels in the knee IALF and serum were detected using ELISA kits (cat. nos. ml037361 and ml102825; Shanghai Enzyme-linked Biotechnology Co., Ltd., Shanghai, China), following the manufacturer's protocol. Additionally, the concentration of protein in the IALF was measured using a bicinchoninic acid (BCA) assay kit (cat. no. P0010S; Beyotime Institute of Biotechnology, Shanghai, China) to correct the error in harvesting.

Immunohistochemical assay. Left knee joint sections were deparaffinized in xylene and rehydrated in graded ethanol and water. Enzymatic antigen retrieval (cat. no. AR0026; Boster Biological Technology, Pleasanton, CA, USA) was performed for 30 min at 37°C for antigen retrieval, and left knee joint sections were then stained using a two-step method, according to the protocol of the SPlink Detection kit (cat. no. SP-9001; Zhongshan Golden Bridge Biotechnology Co., Ltd., Beijing, China). The sections were incubated overnight at 4°C with rabbit polyclonal antibodies against type II collagen (1:500 dilution; cat. no. ab34712), light chain 3β (LC3B; 1:100 dilution; cat. no. ab48394), BECLIN1 (1:800 dilution; cat. no. ab62557) (all from Abcam, Cambridge, MA, USA) and sequestosome 1 (SQSTM1; 1:20 dilution; cat. no. 8420-1-AP; ProteinTech Group, Wuhan, China). Sections were then incubated with biotinylated secondary antibody and streptavidin/horseradish peroxidase were incubated for 30 min at room temperature successively according to the instructions of the SPlink Detection kit. Finally, the sections were visualized with 3,3-diaminobenzidine tetrahydrochloride for 1 min and counterstained with hematoxylin for 5 min. The negative control group was assayed according to the abovementioned procedure, by substituting the primary antibody with PBS. The type II collagen content was evaluated based on the optical density of positive area measured using the image analysis software Image-Pro Plus, version 6.0 (Media Cybernetics, Inc., Rockville, MD, USA). To evaluate the positive cells of LC3B, BECLIN1 and SQSTM1 in the different zones of the cartilage (29), three images of each protein were captured under $\times 200$ magnification, representing the cartilages in the loaded parts. The total number and positive number of chondrocytes, as well as the number of different zones, were counted in each image, respectively. Finally, the percentage of positive cells was calculated.

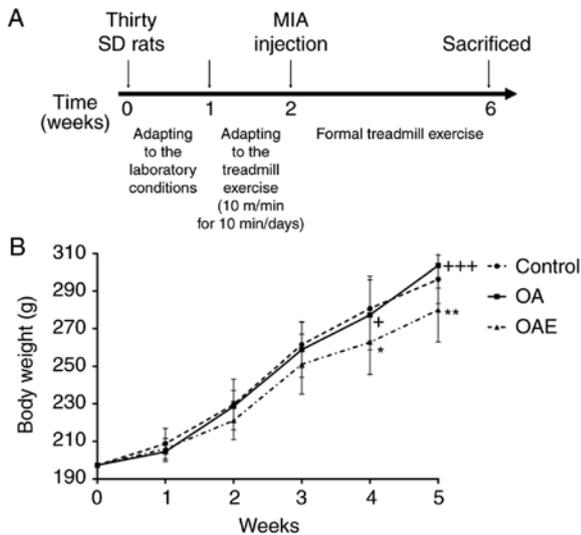


Figure 1. (A) Treatment schedule for the rats. (B) Body weight during the treadmill exercise. All rats received a standard diet, and body weights were measured weekly. No significant difference was detected between the control and OA group during exercise. Differences between control and OAE, OA and OAE groups in weeks 4 and 5 were significant. * $P<0.05$ and ** $P<0.01$ vs. control group; + $P<0.05$ and +++ $P<0.001$ vs. OAE group. Results are expressed as the mean and 95% confidence interval (one-way analysis of variance; $n=10$ rats/group). MIA, monosodium iodoacetate; OA, osteoarthritis; OAE, OA with treadmill exercise.

Western blotting. Following removal from storage at -80°C , the cartilage was washed in PBS twice, and ultrasonication was performed in radioimmunoprecipitation assay lysis buffer (cat. no. P0013C) with 1 mM phenylmethylsulfonyl fluoride (cat. no. ST506) (both from Beyotime Institute of Biotechnology). The lysates were centrifuged at $14,000 \times g$ for 5 min at 4°C , and the supernatants were obtained to measure the protein concentration with the BCA assay kit. Equal amounts of protein ($50 \mu\text{g}$) were added to an SDS-PAGE gel [8% for type II collagen and a disintegrin and metalloproteinase with thrombospondin motifs 5 (ADAMTS5); 12% for LC3B and β -actin] following heating with loading buffer (cat. no. NP0007; Thermo Fisher Scientific, Inc., Waltham, MA, USA) at 100°C for 5 min, and then samples were transferred to polyvinylidene difluoride membranes. Following blocking at room temperature for 2 h with 1% bovine serum albumin (cat. no. A8020; Solarbio Science & Technology Co., Ltd., Beijing, China) in Tris-buffered saline with 0.1% Tween-20 (TBST), the membranes were incubated with primary antibodies overnight at 4°C , including the following: Rabbit polyclonal anti-type II collagen (molecular weight, 142 kDa; 1:5,000 dilution; cat. no. ab34712; Abcam), rabbit polyclonal anti-ADAMTS5 (molecular weight, 100 kDa; 1:200 dilution; cat. no. BA3020; Boster Biological Technology), rabbit polyclonal anti-LC3B (molecular weight, 15 kDa for LC3-II and 17 kDa for LC3-I; 1:1,000 dilution; cat. no. ab48394; Abcam) and mouse monoclonal anti- β -actin (molecular weight, 42 kDa; cat. no. AA128; 1:1,000; Beyotime Institute of Biotechnology) antibodies. Following washing five times with TBST for 5 min each time, the peroxidase-conjugated AffiniPure goat anti-rabbit and anti-mouse IgG (H+L) secondary antibodies (cat. nos. ZB5301 and ZB5305; Zhongshan Golden Bridge Biotechnology Co., Ltd.) were applied at a dilution of 1:10,000 in TBST and incubated at room temperature for

1.5 h. The membrane was washed with TBST again, detected with enhanced chemiluminescence (cat. no. P0018; Beyotime Institute of Biotechnology) and quantified using Image-Pro Plus software, version 6.0 (Media Cybernetics, Inc.). β -actin served as the internal reference.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Following removal of the cartilage from the -80°C freezer, a mortar and pestle were used to grind up the samples in a low-temperature environment maintained by liquid nitrogen. Approximately 50 mg cartilage samples was collected for total RNA isolation with RNAiso Plus (cat. no. 9108; Takara Biotechnology Co., Ltd., Dalian, China) and High-Salt Solution for Precipitation (Plant) (cat. no. 9193; Takara Biotechnology Co., Ltd.). A total of $1 \mu\text{g}$ total RNA was then reverse-transcribed into first-strand complementary DNA (cDNA) with a PrimeScript RT reagent kit with gDNA Eraser (cat. no. RR047A; Takara Biotechnology Co., Ltd.). The cDNA samples were used to conduct qPCR with an ABI Prism 7500 Fast Real-Time PCR system (Applied Biosystems; Thermo Fisher Scientific, Inc.) in a $20\text{-}\mu\text{l}$ system using SYBR Premix Ex Taq II (cat. no. RR820A; Takara Biotechnology Co., Ltd.) with a two-step method (95°C for 30 sec, 95°C 5 sec, 40 cycles of 60°C for 34 sec). All kits and reagents were used according to the manufacturer's protocol. The following primers were used: BECLIN1 forward, 5'-CTGGACCGAGTGACCATT CA-3' and reverse, 5'-AGACACCATCCTGGCGAGTT-3'; LC3B forward, 5'-TTCTTCCTCCTGGTGAAT GG-3' and reverse, 5'-CTGGGAGGCATAGACCATGT-3'; SQSTM1 forward, 5'-CTGTGGTGGGAAGCTCGCTAT-3' and reverse, 5'-GAAAGATGAGCTTGCTGTGCT-3'; β -actin forward, 5'-CACCCGCGAGTACAACCTTC-3' and reverse, 5'-CCC ATACCCACCATCACACC-3'. The expression levels of the PCR products were calculated using the $2^{-\Delta\Delta\text{C}_q}$ method, with β -actin serving as the housekeeping gene (30).

Statistical analysis. All data are expressed as the mean and 95% confidence interval (CI), and were analyzed using IBM SPSS statistical software version 20 (IBM Corp., Armonk, NY, USA). To evaluate the normality of the results, the Kolmogorov-Smirnov and Shapiro-Wilk tests (when sample size was ≤ 5) were performed. Comparisons among groups were determined by one-way analysis of variance with Tukey's tests for pairwise comparisons, following the Levene's test to examine homogeneity of variance. If homogeneity of variance was missing, Games-Howell method was used as a replacement. $P<0.05$ was considered to indicate a statistically significant difference.

Results

Weight change. There were no significant changes between the control and the OA group throughout the procedure. At the same time, there was a significant difference between the OAE and the control, the OAE and the OA group at the fourth and fifth week (Fig. 1B).

Histological analysis of knee cartilage. The joint cartilage of the control group exhibited a slight degeneration in the superficial zone (SZ), including cartilage matrix edema

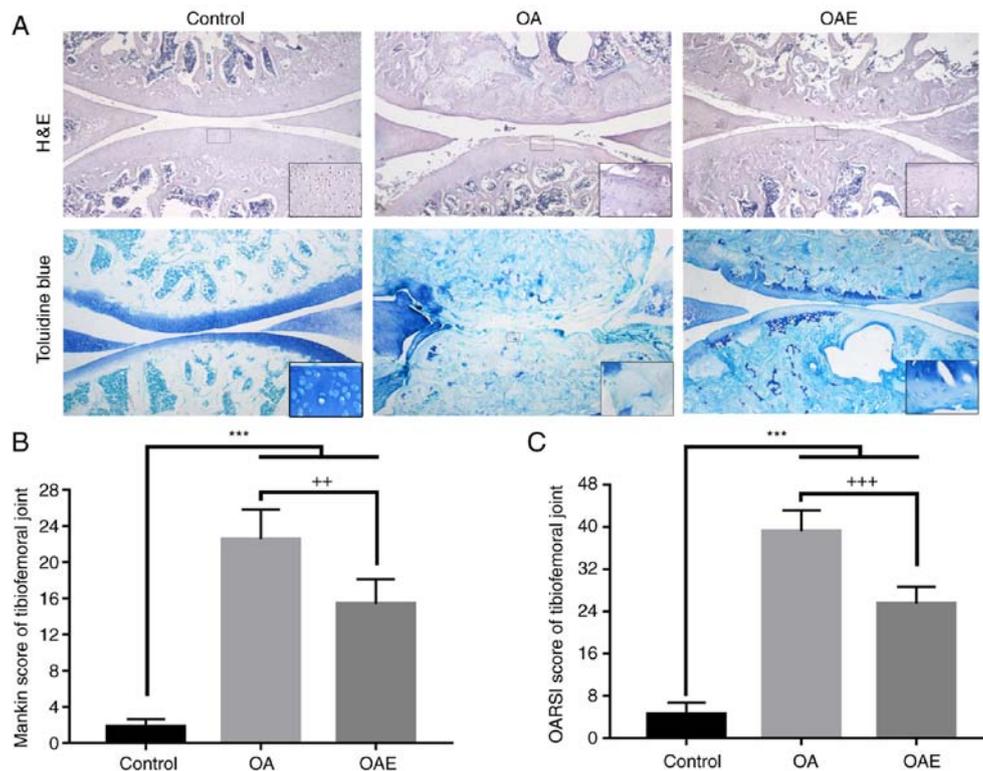


Figure 2. Histological staining and evaluation of the tibiofemoral joint. (A) H&E and toluidine blue staining of sections in the sagittal plane. Images were captured at x40 magnification, and local magnification of a selected square was used to observe the details for each group (x200 for H&E and x400 for toluidine blue). (B) Mankin score and (C) OARSI score for the tibiofemoral joint. Differences between the control vs. OA, control vs. OAE, and OA vs. OAE groups were statistically significant. *** $P < 0.001$, ** $P < 0.01$ and *** $P < 0.001$ vs. OA group. Results are expressed as the mean and 95% confidence interval (one-way analysis of variance; $n = 10$ rats/group). OA, osteoarthritis; OAE, OA with treadmill exercise; H&E, hematoxylin and eosin.

and chondrocyte proliferation (Fig. 2A). By contrast, in the OA group, severe damage was observed in sections stained with H&E and toluidine blue, such as denudation in the deep zone (DZ) and microfractures in the fibrocartilage. In the OAE group, the cartilage exhibited different alterations, from superficial fibrillation to matrix loss in the mid zone. Furthermore, the Mankin score in the control group was 1.80 (95% CI, 0.95-2.65), while it was 22.55 (95% CI, 19.27-25.83) in the OA group and 15.40 (95% CI, 12.69-18.11) in the OAE group (Fig. 2B). The OARSI score was 4.55 (95% CI, 2.36-6.74) in the control group, 39.20 (95% CI, 35.23-43.17) in the OA group, and 25.50 (95% CI, 22.36-28.64) in the OAE group (Fig. 2C). The OA and OAE group scores were significantly increased compared with the control, while there was a marked reduction in the OAE group in comparison with the scores in the OA group.

Type II collagen and ADAMTS5 alterations. As demonstrated via immunohistochemical analysis, type II collagen expression in the OA group was much weaker in comparison with that in the control group, and this reduction was notably attenuated by exercise in the OAE group (relative percentages, 43 and 69% in the OA and OAE groups, respectively; Fig. 3A and B). The expression of type II collagen detected via western blotting exhibited a similar trend to the immunohistochemistry results [control group, 1.87 (95% CI, 1.28-2.46); OA group, 0.30 (95% CI, 0.05-0.55); and OAE group, 0.97 (95% CI, 0.35-1.58); Fig. 3C and D]. On the contrary, the expression of ADAMTS5 was markedly increased in the OA group and

moderately increased in the OAE group [control group, 0.16 (95% CI, -0.07-0.38); OA group, 1.12 (95% CI, 0.83-1.42), and OAE group, 0.47 (95% CI, 0.08-0.87; Fig. 3C and D)].

Inflammation in synovium, and IL-1 β and IL-4 levels detected by ELISA. As compared with the control group, H&E staining of the synovium of the OA group displayed an enlargement of the synovial lining cell layer and inflammatory cell infiltration, accompanied by an increase of the cellularity (Fig. 4A). However, the synovial stroma was mostly normal in the OAE group. The synovitis score in the control group was 0.90 (95% CI, 0.22-1.58), while the scores in the OA and OAE groups were 3.20 (95% CI, 2.49-3.91) and 2.00 (95% CI, 1.12-2.88), respectively (Fig. 4A and B).

ELISA revealed that, compared with the control group, the IL-1 β level was significantly increased in the serum and IALF of the OA group, whereas the expression of IL-4 was notably diminished. In the OAE group, the variation in IL-1 β and IL-4 levels was alleviated compared with that in the OA group, although the differences between the two groups were not significant (Fig. 4C). More specifically, the serum IL-1 β levels were 23.18 (95% CI, 19.77-26.58 pg/ml), 33.75 (95% CI, 28.83-38.66 pg/ml) and 27.56 (95% CI, 23.10-32.01 pg/ml) in the control, OA and OAE groups, respectively. The IL-1 β levels in IALF were 62.31 (95% CI, 56.71-67.91 pg/mg protein), 97.48 (95% CI, 87.26-107.69 pg/mg protein) and 87.18 (95% CI, 78.05-96.31 pg/mg protein) in the three groups, respectively. Furthermore, the IL-4 levels in the control, OA and OAE groups were 83.98 (95% CI,

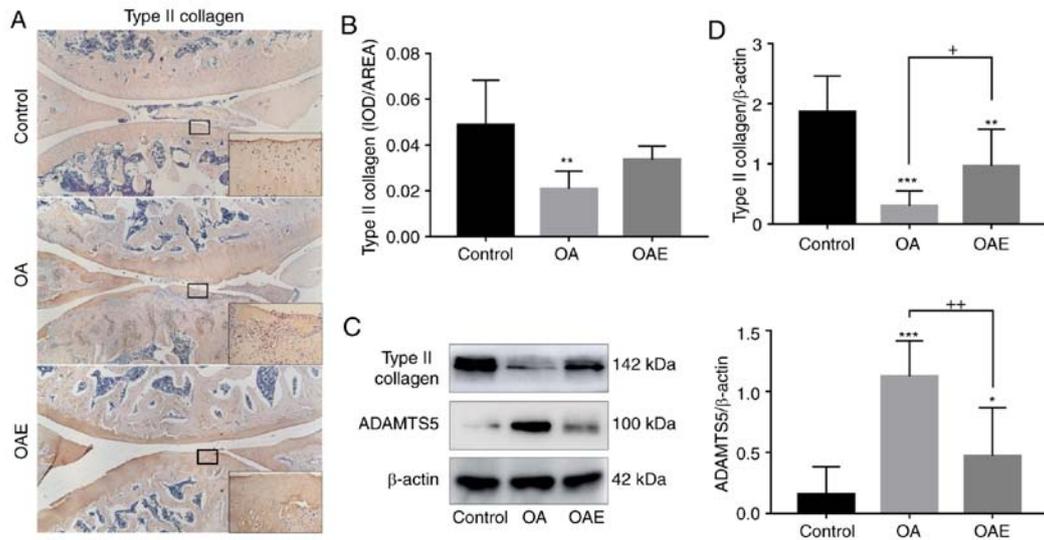


Figure 3. Type II collagen and ADAMTS5 alterations among the groups. (A) Immunohistochemical staining images and (B) quantified levels of type II collagen. Images were captured at x40 magnification, and local magnification of a selected square at x200 was applied. The difference between the control and OA groups was significant (n=5). (C) Western blot analyses and (D) quantified results of the protein levels of type II collagen and ADAMTS5, with β-actin serving as the internal reference. For type II collagen and ADAMTS5, differences between the control vs. OA, control vs. OAE, and OA vs. OAE groups were significant (n=3). Results are expressed as the mean and 95% confidence interval (one-way analysis of variance). *P<0.05, **P<0.01 and ***P<0.001 vs. control group; *P<0.05 and **P<0.01. OA, osteoarthritis; OAE, OA with treadmill exercise; ADAMTS5, a disintegrin and metalloproteinase with thrombospondin motifs 5.

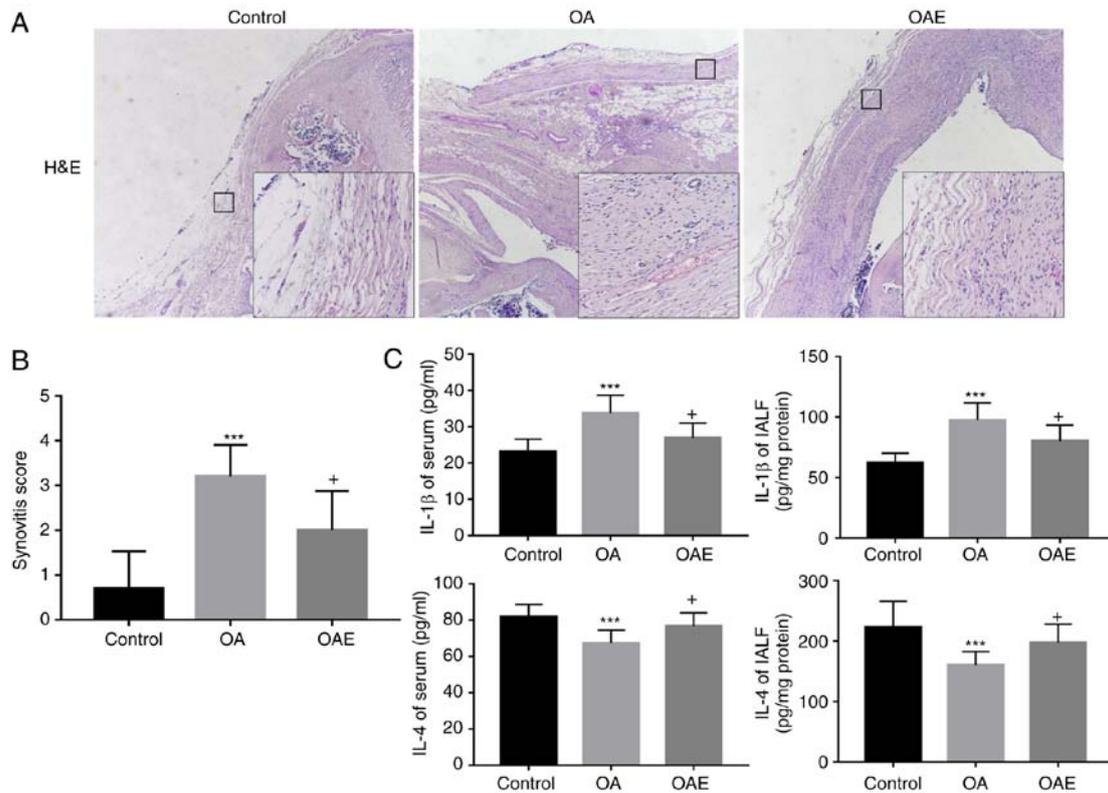


Figure 4. (A) H&E staining for histological evaluation in synovium, and (B) synovitis score. Images were captured at x40 magnification, and local magnification of a selected square at x200 was applied. Differences in synovitis score in the control vs. OA, control vs. OAE and OA vs. OAE groups were statistically significant (n=5). (C) ELISA of IL-1β and IL-4 levels in the serum and IALF. Treadmill exercise reduced the increase in IL-1β and reversed the decrease in IL-4 caused by monosodium iodoacetate injection, in the serum and IALF (n=10). Results are expressed as the mean and 95% confidence interval (one-way analysis of variance). *P<0.05 vs. OA group; ***P<0.001 vs. control group. OA, osteoarthritis; OAE, OA with treadmill exercise; H&E, hematoxylin and eosin; IALF, intra-articular lavage fluid; IL, interleukin.

76.33-91.63 pg/ml), 67.35 (95% CI, 62.21-72.49 pg/ml) and 74.75 (95% CI, 69.02-80.48 pg/ml) in the serum, respectively; these levels in the IALF were 206.40 (95% CI, 174.82-237.98 pg/mg

protein), 160.34 (95% CI, 144.32-176.36 pg/mg protein) and 172.63 (95% CI, 144.76-200.50 pg/mg protein), respectively (Fig. 4C).

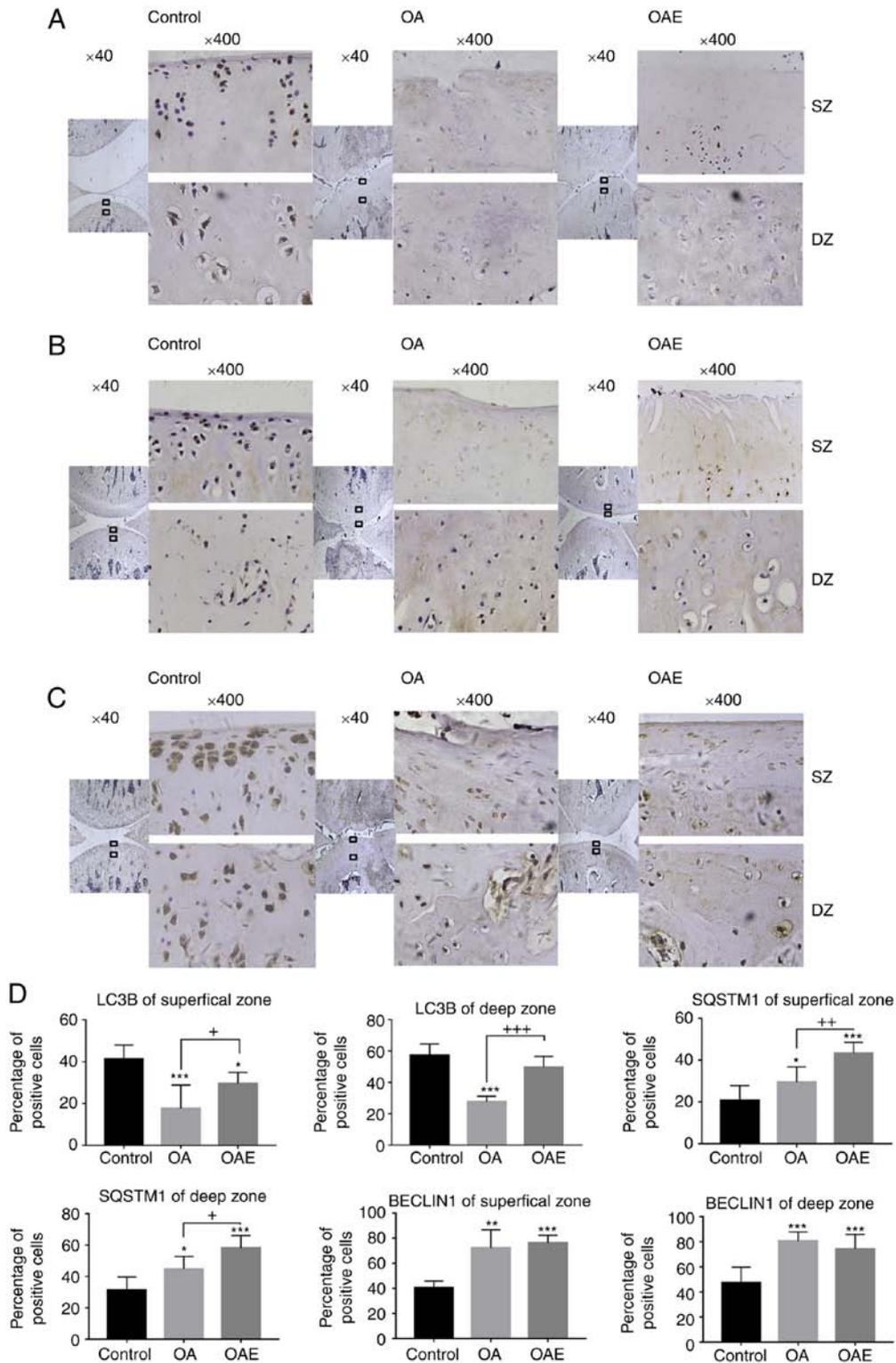


Figure 5. Immunohistochemical staining for LC3B, SQSTM1 and BECLIN1. (A) LC3B exhibited a decrease in the SZ and DZ of the cartilage in both the OA and OAE groups, while treadmill exercise increased LC3B expression in OA rats. (B) SQSTM1 expression was increased in the OA and OAE groups, while treadmill exercise evidently increased SQSTM1 level in the SZ and DZ of the cartilage. (C) BECLIN1 expression was increased in the OA and OAE groups in both the SZ and DZ of the cartilage, with no significant difference between the OA and OAE groups. (D) Results are expressed as the mean and 95% confidence interval (one-way analysis of variance; n=5). *P<0.05, **P<0.01 and ***P<0.001 vs. control group; *P<0.05, **P<0.01 and ***P<0.001. OA, osteoarthritis; OAE, OA with treadmill exercise; LC3B, light chain 3B; SQSTM1, sequestosome 1; SZ, superficial zone; DZ, deep zone.

Autophagy markers in the articular cartilage. As demonstrated by immunohistochemical assay, LC3B was expressed widely in the SZ and DZ of normal cartilage, with a positive detection rate of 41 and 57% of cells, respectively (Fig. 5). However, LC3B

level in the OA group was markedly reduced to 18 and 26% in the SZ and DZ, respectively. In the OAE group, these values were 30 and 50%, respectively. In contrast to LC3B levels, the number of SQSTM1-positive cells gradually increased in the

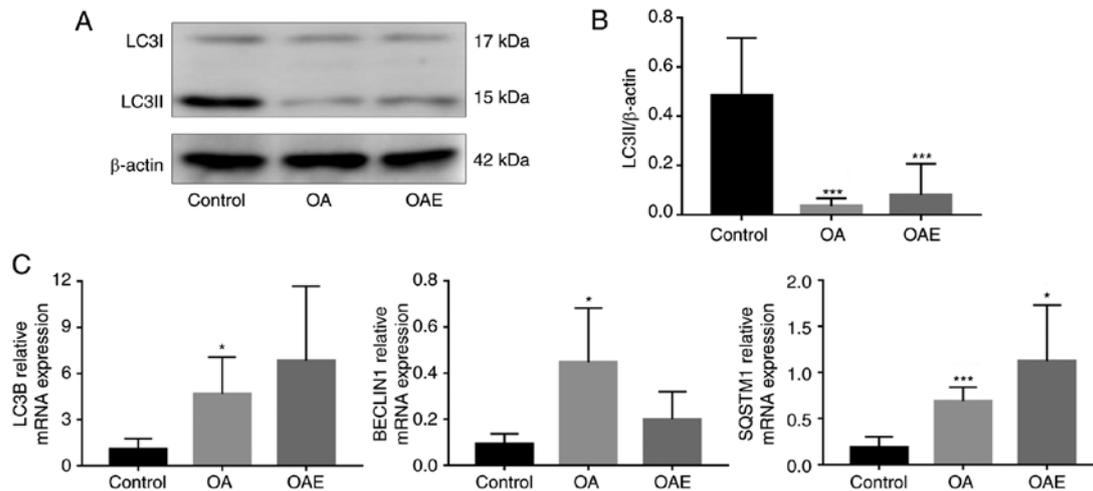


Figure 6. Western blot analysis of LC3B protein level, and reverse transcription-quantitative polymerase chain reaction of LC3B, SQSTM1 and BECLIN1 mRNA expression levels. (A) Western blot analyses and (B) quantified levels of LC3B protein, illustrating a marked decrease in the OA and OAE group compared to the control (n=3). No significant difference was detected between the OA and OAE groups. (C) Relative mRNA expression levels of LC3B, BECLIN1 and SQSTM1 (n=4). There was no significant difference between the OA and OAE groups. Results are expressed as the mean and 95% confidence interval (one-way analysis of variance). *P<0.05 and ***P<0.001 vs. control group. OA, osteoarthritis; OAE, OA with treadmill exercise; LC3B, light chain 3B; SQSTM1, sequestosome 1.

control, OA and OAE groups, with 20, 29 and 43% of cells detected respectively in the SZ, and 31, 45 and 58% detected respectively in the DZ. In terms of BECLIN1, it was notably increased in the different zones in both the OA and OAE groups compared with the control group. The positive detection rates of BECLIN1 were 41, 72 and 76% in the SZ, and 47, 80 and 74% in the DZ in the control, OA and OAE groups, respectively (Fig. 5).

The western blot analyses of LC3B protein are presented in Fig. 6A, while the quantitative analysis of LC3II levels among the groups is presented in Fig. 6B. A marked decrease in LC3II levels was observed in both the OA and OAE groups, with a ratio relative to the control group of 0.073 and 0.167, respectively. RT-qPCR analysis also indicated that, compared with the control group, the mRNA expression levels were significantly increased in the OA group of LC3B and BECLIN1 and in the OA and OAE groups of SQSTM1 (Fig. 6C). Although the difference of LC3B, BECLIN1 and SQSTM1 between the OA and OAE groups was not statistically significant, the mRNA expression levels of LC3B and SQSTM1 exhibited an increasing tendency in OAE. In contrast to LC3B and SQSTM1, the mRNA expression of BECLIN1 was markedly decreased in the OAE group compared with the OA group. The relative mRNA expression of LC3B in the control, OA and OAE groups was 1.08 (95% CI, 0.40-1.76), 4.67 (95% CI, 2.27-7.07) and 6.83 (95% CI, 1.99-11.68), respectively. The relative BECLIN1 levels in these groups were 0.093 (95% CI, 0.049-1.137), 0.447 (95% CI, 0.213-0.682) and 0.2 (95% CI, 0.081-0.319), respectively. Similarly, the relative SQSTM1 levels were 0.188 (95% CI, 0.074-0.302), 0.689 (95% CI, 0.539-0.840) and 1.122 (95% CI, 0.515-1.729), respectively (Fig. 6C).

Discussion

A number of studies have illustrated the therapeutic effect of moderate treadmill exercise in different models of OA (9-13). It has also been reported that systemic factors, such as body weight, and local factors, including muscle strength, are

associated with the occurrence and development of OA (31). In the present study, it was observed that the weight gain in the OAE group was significantly lower compared with the other two groups (Fig. 1B). However, excessive exercise can promote the occurrence of OA (32), suggesting that these prophylactic treatments for OA are limited.

In the present study, an OA model induced by MIA injection was established, with the typical pathological alterations observed in histological analysis, while moderate treadmill exercise was applied in the OAE group and the condition has improved significantly. Alterations in cartilaginous damage were represented by the degradation of type II collagen, the major form of collagen in the matrix. Type II collagen, which is synthesized by chondrocytes and secreted to the extracellular matrix, may be degraded by various proteases, including ADAMTS5 (33). The synthesis of type II collagen is inhibited by IL-1β and the expression of ADAMTS5 is activated simultaneously. In the present study, the histological appearance of the synovium exhibited severe inflammation in OA group which was partially relieved in OAE group. IL-1β, a traditional inflammatory factor, was significantly increased in the OA group, while its expression was attenuated in the OAE group, in both the serum and IALF. It was also observed that the level of IL-4, an anti-inflammatory cytokine that inhibits the proinflammatory effect of IL-1β, was inversely correlated with that of IL-1β. These proteins have been reported to have an important association with the occurrence of OA (34,35). Compared with our previous study (25), the results of the present study indicated a consistent trend in the expression levels of IL-1β and type II collagen.

Autophagy, which is highly conserved in eukaryotes, has been demonstrated to serve an important role in various systems and relevant diseases, including OA. It has been reported that increased expression of LC3B and BECLIN1 mRNA results in increased autophagy in OA chondrocytes and cartilage at the outset, which represents a compensatory response to stress (22). However, in cartilage with mild or severe OA, these two proteins

are strongly expressed in the OA cell clusters, as well as the middle zone and DZ. Their expression is however reduced in the SZ of the cartilage, and this is accompanied by an increase in cell death and cartilage damage (36,37). To the best of our knowledge, studies on the alteration in autophagy in an OA model following treatment with exercise are lacking in the literature.

In the present study, the results appeared to be consistent with the findings of a previous study demonstrating that LC3B was decreased in different zones of the cartilage in OA (18). However, BECLIN1 was strongly expressed in the OA group in the SZ and DZ, and the increase was statistically significant; these results were partly in accordance with those obtained by Sasaki *et al* (22). This may be explained by the complexity of autophagy in different periods and zones of OA, and the distinction between a rat model induced via MIA injection and spontaneous OA in the elderly. SQSTM1 is reported to be a link between LC3 and ubiquitinated substrates (38), and the present study observed an increase in its expression in OA cartilage. A possible explanation for this effect is that the upregulated transcriptional level supplemented the consumed protein during degradation in the lysosome (39). Another important finding of the current study was that exercise markedly promoted the expression of LC3B and SQSTM1 in whole cartilage with immunohistochemical assay compared with the OA group. Notably, no differences were observed in BECLIN1 expression between the OA and OAE groups, which may be partly explained by the excessive expression in OA cartilage. In addition, western blot analysis revealed that LC3II, which serves as a marker of autophagosome formation, was markedly decreased in the OA group and this was partially alleviated in the OAE group. It was also demonstrated that, although the mRNA expression levels of the three genes of interest was elevated following prolonged stimulation with MIA, only BECLIN1 and SQSTM1 protein levels were increased in cartilage, while LC3B protein was decreased. Furthermore, the decrease detected in BECLIN1 mRNA expression in OA following treadmill exercise may be due to the complex role of BECLIN1 in autophagy and apoptosis (40). It can be hypothesized that the excessive consumption of LC3B limited the speed of autophagy and caused the accumulation of BECLIN1 and SQSTM1 proteins in the OA group, which may explain the upregulated mRNA expression of LC3B and simultaneous decrease of its protein expression. Finally, the results demonstrated that the mRNA expression levels of LC3B and SQSTM1 exhibited an increasing trend in the OAE group in comparison with the OA group, although this result was not found to be significant, likely due to the small sample size.

Our previous study demonstrated that multiple sessions of short-term exercise were associated with an improved chondroprotective effect compared with one long session of exercise for the same total time (25). The important role of autophagy has been recently reported in chondrogenesis in the regulation of the growth plate (41). In addition, a previous study indicated that the proliferation of chondrocytes was associated with mechanical stimulation (42). Therefore, it can be hypothesized that the increased expression of autophagy-associated proteins in cartilage caused by moderate treadmill exercise is associated with chondrocyte differentiation and proliferation. The present study may provide a basis for a new therapeutic strategy for OA via moderate exercise. Future studies on this

topic are recommended, as several questions remain unanswered at present.

Although the effects of exercise on autophagy have been examined in numerous organs and systems, there is little evidence for bone and cartilage, particularly in OA. The findings of the present study, while preliminary, suggest that moderate exercise has a marked effect on autophagy in the cartilage in OA, although the limitations of the detection method for autophagy caused in animal models should be considered. Furthermore, the limitations of small sample size and bias during observation cannot be neglected.

In conclusion, the present study provided strong evidence on the hypothesis that moderate treadmill exercise has a protective effect on the articular cartilage of rats with OA induced via MIA injection. In spite of several limitations, the present study suggested that moderate treadmill exercise may promote autophagy in all zones of the articular cartilage in rats with MIA-induced OA.

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Availability of data and material

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

XZ, HZ and LB conceived and designed the study. XZ, YG, YY and XL conducted the experiments. HZ, YG and XL performed the statistical analysis, and designed the figures. XZ wrote the manuscript. YY and LB reviewed and edited the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The present study was approved by the Medical Research and New Technology Ethics Committee of Shengjing Hospital, China Medical University (Shenyang, China).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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