

# Naringin and bone marrow mesenchymal stem cells repair articular cartilage defects in rabbit knees through the transforming growth factor- $\beta$ superfamily signaling pathway

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Received July 25, 2019; Accepted April 21, 2020

DOI: 10.3892/etm.2020.9187

**Abstract.** The present study aimed to assess the effect of a combination of naringin and rabbit bone marrow mesenchymal stem cells (BMSCs) on the repair of cartilage defects in rabbit knee joints and to assess possible involvement of the transforming growth factor- $\beta$  (TGF- $\beta$ ) signaling pathway in this process. After establishing an articular cartilage defect model in rabbit knees, 20 New Zealand rabbits were divided into a sham operation group (Sham), a model group (Mod), a naringin treatment group (Nar), a BMSC group (BMSCs) and a naringin + BMSC group (Nar/BMSCs). At 12 weeks after treatment, the cartilage was evaluated using the International Cartilage Repair Society (ICRS)'s macroscopic evaluation of cartilage repair scale, the ICRS's visual histological assessment scale, the Modified O'Driscoll grading system, histological staining (hematoxylin and eosin staining, toluidine blue staining and safranin O staining) and immunohistochemical staining (type-II collagen, TGF- $\beta$ 3 and SOX-9 immunostaining). Using the above grading systems to quantify the extent of repair, histological quantification and macro quantification of joint tissue repair showed that the Nar/BMSCs group displayed

repair after treatment in comparison to the untreated Mod group. Among the injury model groups (Mod, Nar, BMSCs and Nar/BMSCs), the Nar/BMSCs group displayed the highest degree of morphological repair. The results of histological and immunohistochemical staining of the repaired region of the joint defect indicated that the BMSCs had a satisfactory effect on the repair of the joint structure but had a poor effect on the repair of cartilage quality. The Nar/BMSCs group displayed satisfactory therapeutic effects on both repair of the joint structure and cartilage quality. The expression level of type-II collagen was high in the Nar/BMSCs group. Additionally, staining of TGF- $\beta$ 3 and SOX-9 in the Nar/BMSCs group was the strongest compared with that of any other group in the present study. Naringin and BMSCs together demonstrated a more efficient repair effect on articular cartilage defects in rabbit knees than the use of either treatment alone in terms of joint structure and cartilage quality. One potential mechanism of naringin action may be through activation and continuous regulation of the TGF- $\beta$  superfamily signaling pathway, which can promote BMSCs to differentiate into chondrocytes.

## Introduction

Knee osteoarthritis is a degenerative joint disease induced by articular cartilage degeneration. Its main clinical features are arthralgia and joint deformity and dysfunction (1). It is characterized by a high incidence and recurrent seizures of the joint and worsens with time. As the condition advances, prosthetic replacement surgery is usually required. Over a million people worldwide receive new diagnoses of articular cartilage disease annually (2) and the condition lowers the quality of life of these patients (3).

Articular cartilage, an avascular tissue, can be classified into hyaline cartilage, fibrocartilage and elastic cartilage. Hyaline cartilage is located on the surface of the knee joint

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**Key words:** naringin, rabbit, knee joint, cartilage defect, bone mesenchymal stem cells, transforming growth factor- $\beta$  superfamily signaling pathway

and has critical functions that differ from the other two types of cartilage, such as the secretion of lubricating substances and friction reduction. Hyaline cartilage plays a vital role in maintaining the normal functions of the knee joint (4).

It is well accepted that knee joint cartilage has a poor blood supply. Moreover, it lacks stromal stem cells and has an extremely poor ability for self-repair (5). When cartilage is repaired, it is usually spontaneously repaired into fibrocartilage, not hyaline cartilage, therefore losing the extra function of this cartilage form (6). There are a number of differences between fibrocartilage and hyaline cartilage in terms of tissue structure and mechanical properties (7). Articular cartilage defects, the predominant factor in the pathological process of osteoarthritis, lead to degeneration of the joint (8).

Much attention has been paid to natural small molecule compounds based on plant derivatives for the potential treatment of osteoarticular disease, attributed to the feasibility of their collection and their bioactivity and therapeutic effects (9). A previous study indicated that these compounds may be beneficial in the treatment of osteoarticular diseases through the regulation of signaling transduction pathways (10).

In traditional Chinese medicine (TCM), the pathogenesis of osteoarthritis is thought to involve liver and kidney deficiencies, as well as malnutrition of the muscles and bones (11-13). *Drynaria fortunei* is a commonly used treatment in TCM which is thought to benefit the liver and kidneys, strengthen tendons and bones and target the key treatment points of osteoarthritis. *Drynaria fortunei* is thought to be therapeutically beneficial in the treatment of knee osteoarthritis (14). The present study was designed to investigate the specific mechanism of action of *Drynaria fortunei*, as current knowledge is lacking. Naringin is a plant derivative and also an active ingredient of *Drynaria fortunei* and is thought to have the same effect as the corresponding dosage of the crude *Drynaria fortunei* (15).

Mesenchymal stem cells (MSCs) are cells with marked proliferative abilities and the ability to differentiate into cartilage cells (16). Myeloid tissue is one of the main sources of MSCs. MSCs that exist in the bone marrow are termed bone marrow mesenchymal stem cells (BMSCs). They are one of the most commonly used cells in research into articular cartilage repair (17). A number of compounds are thought to exhibit potential for promotion of the differentiation of BMSCs, such as insulin-like growth factor I and vascular endothelial growth factor (18).

The transforming growth factor (TGF)- $\beta$  superfamily includes TGF- $\beta$ s and bone morphogenetic proteins. The TGF- $\beta$  signaling pathway plays a vital role in cartilage cell proliferation and differentiation. SOX-9, a downstream factor in this pathway, plays an indispensable function in articular cartilage differentiation and is a transcriptional regulator of chondrogenesis (19). SOX-9 is necessary to support a normal cartilage cell phenotype (20). Overexpression of SOX-9 results in increased synthesis of aggrecan, which can provide a suitable living environment for articular cartilage (21).

Naringin, the active ingredient of *Drynaria fortunei*, can be taken orally or injected directly into a cartilage defect site in combination with BMSCs and has previously been suggested to stimulate the repair of articular cartilage defects in rabbit knees (22). The present study aimed to conduct an in-depth investigation into the repair of joint structure and cartilage and

assess the role of the TGF- $\beta$  superfamily signaling pathway in this process. A scoring system, histological staining and immunohistochemical staining were used to quantify the efficacy of naringin induced repair and the potential mechanisms of naringin action were explored (Fig. 1).

## Materials and methods

**Extraction and separation of rabbit BMSCs.** Following a previously published protocol (23), a 2-week-old New Zealand rabbit was used for the collection of BMSCs. Rabbits were anesthetized by intravenous injection of 30 mg/kg pentobarbital sodium into the ear margin. A bone marrow puncture needle was then inserted into the posterior superior iliac spine of the rabbit. A total of 20 ml of low molecular weight heparin calcium (Tianjin Chase Sun Pharmaceutical Co., Ltd.) was injected to extract the bone marrow. The inner wall of the syringe was moistened with low molecular weight heparin 5 min before extraction. The extracted bone marrow was then injected into a sterile centrifuge tube and was diluted 1:5 with L-DMEM (Beijing Solarbio Science & Technology Co., Ltd.). The bone marrow-DMEM mixture was centrifuged at  $1,200 \times g$  for 5 min at 25°C. The supernatant was discarded and subsequently cells were resuspended in 20 ml DMEM containing 10% fetal calf serum (Gibco; Thermo Fisher Scientific, Inc.). The extracted cells were added to a culture flask and placed in an incubator at 37°C and 5% CO<sub>2</sub>. The medium was replaced with fresh medium 48 h later. The full-marrow adherent method (24) was employed for separation and purification of rabbit BMSCs. The culture medium was replaced every 3 days until the cells reached 80% confluency, when they were passaged for further experiments.

**Preparation of tested animals.** In total, 20 New Zealand rabbits (3-month-old, male) were selected, with a weight of ~2.5 kg prior to surgery. They were placed in separate cages and allowed to freely move within the cage. The animals had access to food and water *ad libitum*. A 12-h light-dark cycle was maintained. The temperature was maintained at 21-25°C and the relative humidity was maintained at 40-70%.

The animals were acclimatized for  $\geq 7$  days before subsequent experiments. The present study followed the National Institutes of Health Guidelines Regarding the Care and Use of Laboratory Animals. Animal treatment in the experimental process conformed to the relevant regulations on ethical standards of animals specified in the Guiding Proposal for Being Kind to Experimental Animals issued by the Ministry of Science and Technology and in Ethical Issues in Animal Experimentation (11,25). The animal experiments were approved by the Science and Technology Department of Beijing University of Chinese Medicine (Beijing, China) and the animal ethics review committee of the Institute of Basic Theories of Chinese medicine, Chinese Academy of Chinese Medical Sciences, approval no. 201706058 (Beijing, China).

**Animal models.** Animal models for articular cartilage defects were prepared according to a previously described method (22,23). New Zealand rabbits were purchased from Beijing Changyang Xishan Experimental Animal Farm. Rabbits were anesthetized using intravenous administration

Table I. New Zealand white rabbit baseline information.

Parameter	Sham operation group	Model group	Naringin group	BMSCs group	Naringin+BMSCs group
Weight, kg	2.80±0.41	2.78±0.17	2.83±0.22	3.10±0.24	3.20±0.34
Sex, male/female	4/0	4/0	4/0	4/0	4/0
Age, month	6	6	6	6	6

of 3% sodium pentobarbital (30 mg/kg) in the ear vein. Knee tissues were cut layer by layer under aseptic conditions after anesthesia (skin, subcutaneous tissue, muscle, joint capsule). A corneal trephine (Beijing North Sanyou Medical Devices Co., Ltd.) was used to drill a hole on the articular surface of the trochlea from the femoral condyle to the subchondral bone, with a diameter of 5 mm and a depth of 3 mm. The defect site was filled according to the experimental scheme. The patella was reset after completion. The knee joint capsule and skin were sutured. Penicillin ( $10 \times 10^4$  U) was injected intramuscularly postoperatively each day for 3 days. All rabbits were allowed to move their knee joints freely in their cages without restriction and clinical signs were observed daily. No animal was excluded owing to abnormal clinical findings. The rabbits were anesthetized using intravenous administration of 3% sodium pentobarbital (30 mg/kg) in the ear vein. Measurement of anesthetic depth, respiratory rate and pattern and assessment of mucous membrane color and heart rate monitoring was performed throughout anesthesia. The animals were sacrificed by an intravenous air bolus injection into the ear vein containing 30 ml air, which was injected into the ear margin. Animal death was confirmed by respiratory arrest, cardiac arrest and changes in mucosal color. Rabbits in different groups were sacrificed at the 12th week after surgery. There were no statistically significant differences in body weight among the different groups (Table I).

**Preparation of naringin solution.** A naringin solution was prepared with 10 g naringin and 100 ml deionized water. The gavage concentration (calculated according to the weight of naringin) was 0.0084 g/kg daily (26,27).

**Laboratory animal grouping and intervention method.** All 20 New Zealand rabbits were randomly distributed into 5 groups (4 rabbits and 8 knee joints in each group). In the sham operation group (sham), the joint capsule was cut and seamed, while the knee joint was not treated. In the model group (Mod), deionized water was administered via gavage every day after successful establishment of the articular cartilage defect model. In the naringin group (Nar), naringin solution was administered via gavage every day after successful establishment of the articular cartilage defect model. In the BMSCs group (BMSCs), rabbit BMSCs were immediately implanted at the joint defect site after successful establishment of the articular cartilage defect model. Moreover, deionized water was administered via gavage every day. In the naringin + BMSCs group (Nar/BMSCs), after successful establishment of the articular cartilage defect model, BMSCs were implanted at the defect site immediately and the naringin solution was administered to the rabbits every day. The daily volume of

Table II. International Cartilage Repair Society macroscopic evaluation of cartilage repair.

Category	Score
Degree of defect repair	
In level with surrounding cartilage	4
75% repair of defect depth	3
50% repair of defect depth	2
25% repair of defect depth	1
No repair of defect depth	0
Integration to border zone	
Complete integration with surrounding cartilage	4
Demarcating border-1 mm	3
Three-quarters of graft integrated, one-quarter with a notable border-1 mm in width	2
One-half of graft integrated with surrounding cartilage, one-half with a notable border-1 mm	1
From no contact, to one-quarter of graft integrated with surrounding cartilage	0
Macroscopic appearance	
Intact smooth surface	4
Fibrillated surface	3
Small, scattered fissures or cracks	2
Several small or few large fissures	1
Total degeneration of grafted area	0
Overall repair assessment	
Grade I, normal	12
Grade II, nearly normal	8-11
Grade III, abnormal	4-7
Grade IV, severely abnormal	1-3

deionized water in the Mod and BMSC groups was 0.25 ml. The volume of gavage in Nar and Nar/BMSCs groups was 0.25 ml.

**Macroevaluation.** In order to reexamine the intra-articular structure and to cut out the complete double knee joint of rabbits, arthrotomy was carried out using the same approach as that used during modeling. The International Cartilage Repair Society (ICRS) macroscopic evaluation of cartilage repair (Table II) was adopted to systematically evaluate the degree of articular cartilage repair in the different groups (28). The main parameters monitored were the degree of defect repair, integration to the border zone and macroscopic appearance. All specimens were independently evaluated by two professionals

with intermediate professional titles. The investigators who performed the evaluations were blind to the treatment.

**Histological and immunohistochemical evaluation.** The specimens were fixed in 10% neutral formalin at room temperature for 24 h. For decalcification the specimens were placed in EDTA-buffered solution (Beijing Zhongshan Jinqiao Biotechnology Co. Ltd.) for 4 weeks. The decalcifying solution was changed twice per week. The articular cartilage of the repaired area was cut out after obtaining satisfactory decalcification (Achieved when the syringe needle punctured the tissue without resistance and the tissue was soft and elastic.), so as to be trimmed into a sheet with a thickness of 4 mm. Freshly prepared 4% methanal was used for fixation for 48 h at room temperature, followed by paraffin embedding and slicing. The specimens were cut into 4- $\mu$ m thick slices for histological and immunohistochemical staining, including hematoxylin and eosin (H&E) staining (H&E staining kit; Beijing Solarbio Science & Technology Co., Ltd.) to observe morphological changes in cartilage cells; toluidine blue staining (Toluidine Blue O cartilage stain solution; Beijing Solarbio Science & Technology Co., Ltd.) to observe the secretion of acid mucopolysaccharide and indirectly evaluate the cartilage matrix; safranin O staining (Safranin O-Fast Green FCF cartilage stain kit; Beijing Solarbio Science & Technology Co., Ltd.) to display the conditions of osteogenesis and chondrogenesis; type-II collagen immunostaining (rabbit anti-collagen II antibody; Beijing Bioss Biotechnology Co., Ltd.) to detect specific indicators of cartilage cells and exhibit the content and quality of hyaline cartilage. Slices were subsequently blocked in 3% H<sub>2</sub>O<sub>2</sub> (Beijing Zhongshan Jinqiao Biotechnology Co. Ltd.) for 10 min at room temperature before staining. Slices were incubated with rabbit anti-collagen II antibody (Beijing Bioss Biotechnology Co., Ltd.; cat. no. bs-10589R; 1:200) at 37°C for 2 h, followed by incubation with horseradish peroxidase conjugated goat anti-rabbit/mouse secondary antibody (1:1; cat. no. PV6000; Beijing Zhongshan Jinqiao Biotechnology Co., Ltd) at room temperature for 30 min. TGF- $\beta$ 3 immunostaining was performed to evaluate the expression and secretion of TGF- $\beta$ 3. Slices were incubated with rabbit anti-TGF- $\beta$ 3 antibody (Rabbit anti-TGF- $\beta$ 3 antibody; Beijing Bioss Biotechnology Co., Ltd.; cat. no. bs-1910R; 1:300) at 37°C for 2 h, followed by incubation with horseradish peroxidase conjugated goat anti-rabbit secondary antibody (1:1; cat. no. PV6000; Beijing Zhongshan Jinqiao Biotechnology Co., Ltd) at room temperature for 30 min. SOX-9 immunostaining was used to evaluate the expression and secretion of SOX-9. Slices were incubated rabbit anti-SOX9 antibody (rabbit anti-SOX9 antibody; Beijing Bioss Biotechnology Co., Ltd.; cat. no. bs-10725R; 1:200) at 37°C for 2 h, followed by incubation with horseradish peroxidase conjugated goat anti-rabbit secondary antibody (1:1; PV6000; Beijing Zhongshan Jinqiao Biotechnology Co., Ltd.) at room temperature for 20 min. After staining, the slices were photographed and examined using an optical microscope (Nikon Eclipse 600; Nikon Corporation) equipped with a digital camera (Nikon DXM1200F; Nikon Corporation) to record images of the stained slices.

A total of 40 slices were prepared from each rabbit and from each slice three images were captured. All staining procedures were conducted according to the manufacturers' instructions.

Table III. International Cartilage Repair Society visual histological assessment scale.

Feature	Score
Surface	
Smooth/continuous	3
Discontinuities/irregularities	0
Matrix	
Hyaline	3
Mixture, hyaline/fibrocartilage	2
Fibrocartilage	1
Fibrous tissue	0
Cell distribution	
Columnar	3
Mixed/columnar-clusters	2
Clusters	1
Individual cells/disorganized	0
Cell population viability	
Predominantly viable	2
Partially viable	1
<10% viable	0
Subchondral bone	
Normal	3
Increased remodeling	2
Bone necrosis/granulation tissue	1
Detached/fracture/callus at base	0
Cartilage mineralization (salcified cartilage)	
Normal	2
Abnormal/inappropriate location	0
Toluidine blue stain	
Normal	4
Slight reduction	3
Moderate reduction	2
Severe reduction	1
No staining	0
Percent toluidine blue in defect	
75-100%	4
50-75%	3
25-50%	2
0-25%	1
No toluidine blue staining	0

The ICRS Visual Histological Assessment Scale (Table III) and the modified O'Driscoll grading system (Table IV) were utilized to evaluate articular cartilage repair level in the different groups (29-31). The main parameters included surface, matrix and cell distribution, cell population viability, nature of the predominant tissue, structural characteristics and absence of cellular changes due to degeneration. All specimens were independently evaluated by two professionals.

**Statistical analysis.** SPSS 19.0 software (IBM Corp.) and Prism 7 (GraphPad Software, Inc.) were used for statistical analysis. Data are presented as the mean value  $\pm$  SD. All

Table IV. Modified O'Driscoll grading system.

A, Nature of the predominant tissue	
Parameter	Score
Cellular morphology	
Hyaline articular cartilage	4
Incompletely differentiated mesenchyme	2
Fibrous tissue or bone	0
Safranin-O staining of the matrix	
Normal or nearly normal	3
Moderate	2
Slight	1
None	0
B, Structural characteristics	
Parameter	Score
Surface regularity	
Smooth and intact	3
Superficial horizontal lamination	2
Fissures, 25-100% of the thickness	1
Severe disruption. including fibrillation	0
Structural integrity	
Normal	2
Slight disruption. including cysts	1
Severe disintegration	0
Thickness	
100 per cent of normal adjacent cartilage	2
50-100 per cent of normal cartilage	1
0-50 per cent of normal cartilage	0
Bonding to the adjacent cartilage	
Bonded at both ends of graft	2
Bonded at one end, or partially at both ends	1
Not bonded	0
C, Freedom from cellular changes of degeneration	
Parameter	Score
Hypocellularity	
Normal cellularity	3
Slight hypocellularity	2
Moderate hypocellularity	1
Severe hypocellularity	0
Chondrocyte clustering	
No clusters	2
<25 per cent of the cells	1
25-100 per cent of the cells	0
Freedom from degenerative changes in adjacent cartilage	
Normal cellularity. no clusters, normal staining	3
Normal cellularity, mild clusters, moderate staining	2
Mild or moderate hypocellularity, slight staining	1
Severe hypocellularity. poor or no staining	0

statistical inferences were subjected to two-sided tests.  $P < 0.05$  was considered to indicate a statistically significant difference. The confidence interval of parameter estimation was 95%. Statistical differences were analyzed using Kruskal-Wallis tests followed by post-hoc testing using Dunn's multiple comparison tests.

## Results

*Morphological observation of tissue repair tissue on the joint defect site.* The effect of each treatment on the repair of the surface structure of the articular cartilage defects was studied on the basis of morphological observations. 'Degree of defect repair', 'In level with the surrounding cartilage', 'Integration to border zone', 'Macroscopic appearance' markers.

The main findings were as follows: i) Sham group: There were no surface defects on the knee joint; however, there was a slight abrasion at the edge (Fig. 2A). ii) Mod group: The defect site displayed some tissue granulation with an obvious sunken defect area. There were obvious differences between the areas of normal cartilage tissue and the areas of regenerated tissue. Moreover, an obvious boundary existed between the areas of normal tissue and the areas of defective tissue (Fig. 2B). iii) Naringin group: At the defect site, tissue was semi-transparent and granulated. The defect was not completely filled with regenerated tissue. Compared with the Mod group, the defect area was smaller, with a reduced depth. The boundary between the defect and the surrounding cartilage tissue was fuzzy but distinguishable (Fig. 2C). iv) BMSCs group: Tissue at the defect appeared transparent, in a similar manner to the cartilage tissue. When compared with the Mod group, the defect area was smaller, the depth was shallow, having clearly been reduced and the boundary appeared fuzzy. Although the thickness of the regenerated cartilage was similar to that of the surrounding normal cartilage, the area at the edge of the defect was not as smooth as that observed in normal cartilage (Fig. 2D). v) Nar/BMSCs group: The repaired defect site was integrated with the surrounding tissue. Repairing tissue at the defect site had become milk white, semi-transparent and smooth. It was hard to distinguish repaired tissue at the defect site from the surrounding normal cartilage tissue. The defect site had also developed a smooth surface (Fig. 2E). The characteristic manifestations of the above groups can also be found in the corresponding images before the index detection (Fig. 2F-J). In summary morphological observations in the Naringin group and in the BMSCs group suggested better tissue healing than in the Mod group. However, the Nar/BMSCs group showed the most signs of healing from the defect model animals (Fig. 2).

*Pathological staining and observation of the location of repaired tissues at the joint defect site.* The repair of joint tissues was observed. The structural repairs to the articular cartilage defects in the treatment groups administered BMSCs were semi-transparent and there were no clear signs of inflammation, however, in the groups that were not administered BMSCs, articular cartilage defects were not smooth and the boundaries between new tissue and surrounding undamaged tissues were readily observed. The treatment groups administered BMSCs appeared to display improved structural repair

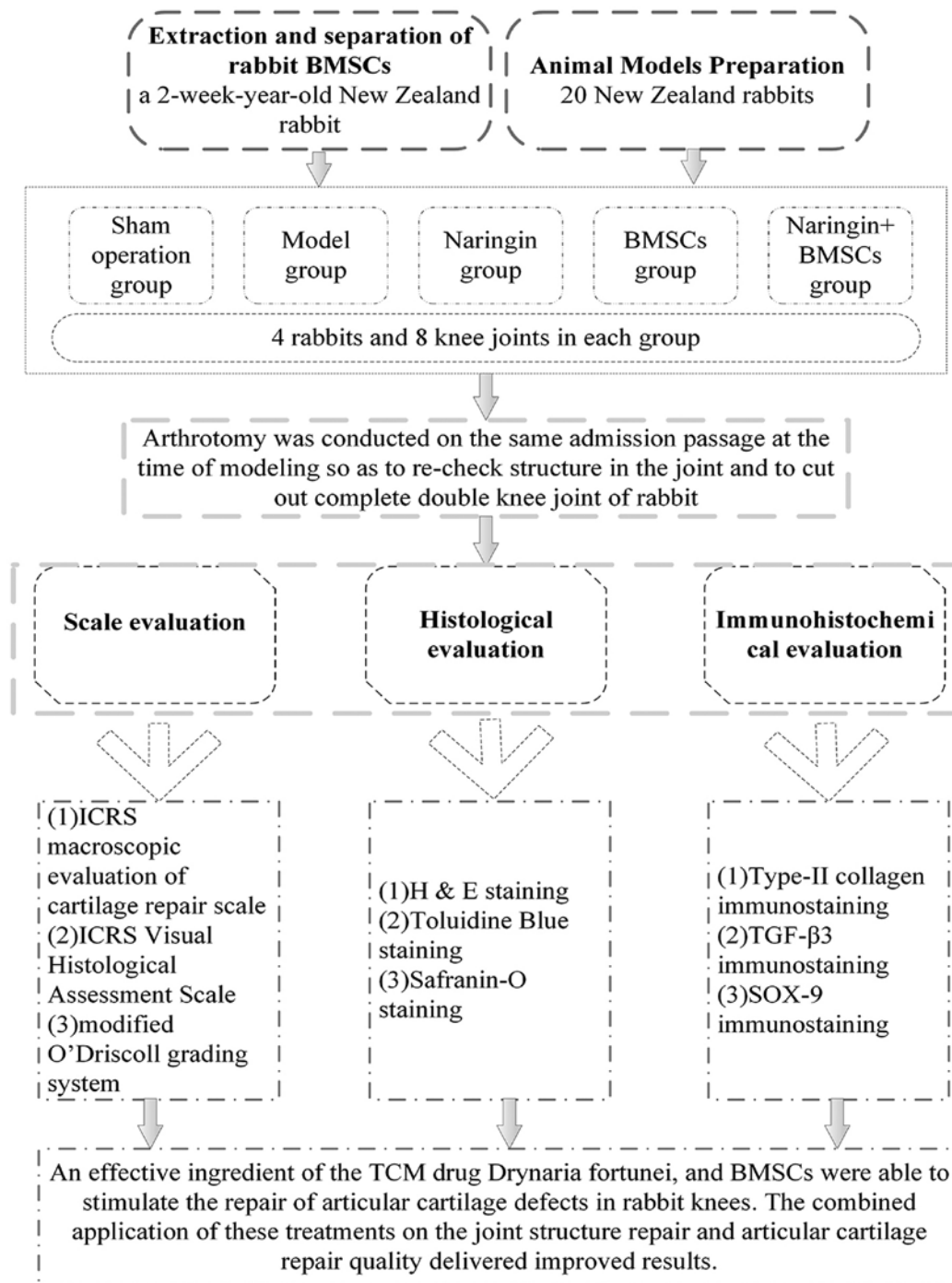


Figure 1. Flow chart describing the structure of experiments performed to evaluate naringin and bone marrow mesenchymal stem cell induced repair of articular cartilage defects in rabbit knees. BMSCs, bone mesenchymal stem cells; ICRS, International Cartilage Repair Society; TGF-3, transforming growth factor-β3; SOX-9, sry-type high-mobility-group box-9.

to those that were not administered BMSCs. In terms of the quality of the repair of articular cartilage defects, in the groups that were not administered naringin, cartilage damage was obvious, the hyaline cartilage was severely worn, the thickness of the cartilage layer was reduced and the tide mark was blurred. However, in the groups administered naringin, hyaline cartilage appeared at the site of the cartilage damage, and the tide line was clear, which was close to the nature of surrounding tissue. The Nar/BMSCs group was observed to have the most efficient repair regarding structural and quality repairs of the articular cartilage defects, as shown in Fig. 3.

H&E staining was used to observe the cartilage morphology. Cells in the repaired tissues of the Mod group were arranged irregularly. Moreover, an obvious gap could be observed (Fig. 3B). The Nar group was observed to have poor structural repair at the defect site. Although the number of repairing cells was limited, the structures and arrangements of the repairing cells generated at the defect site were similar to those of the surrounding normal cartilage cells (Fig. 3C). The BMSCs group was observed to have an efficient overall contour restoration; however, the edge of the repairing tissue was irregular, with obvious gaps (Fig. 3D). The regenerative

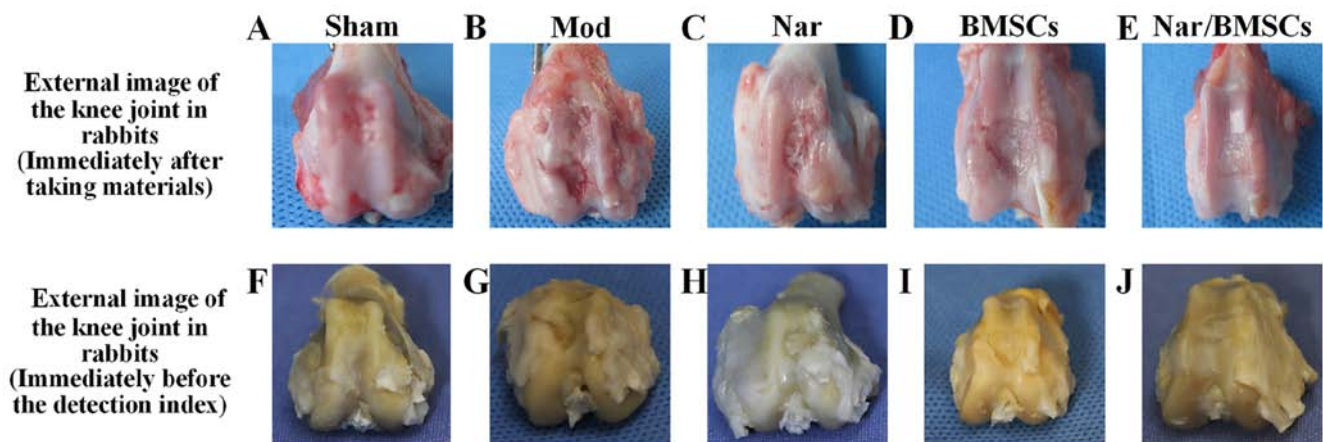


Figure 2. Gross appearance and assessment results of articular cartilage defects in the rabbit models at 12 weeks. General appearance of rabbit knee joints at 12 weeks post-operation, including the (A) Sham group, (B) Mod group, (C) Nar group, (D) BMSCs group and (E) Nar/BMSCs group. General appearance of rabbit knee joints before the histological and immunohistochemical assessment, including the (F) Sham group, (G) Mod group, (H) Nar group, (I) BMSCs group and (J) Nar/BMSCs group. BMSC, bone marrow mesenchymal stem cells; Mod, model; Nar, naringin.

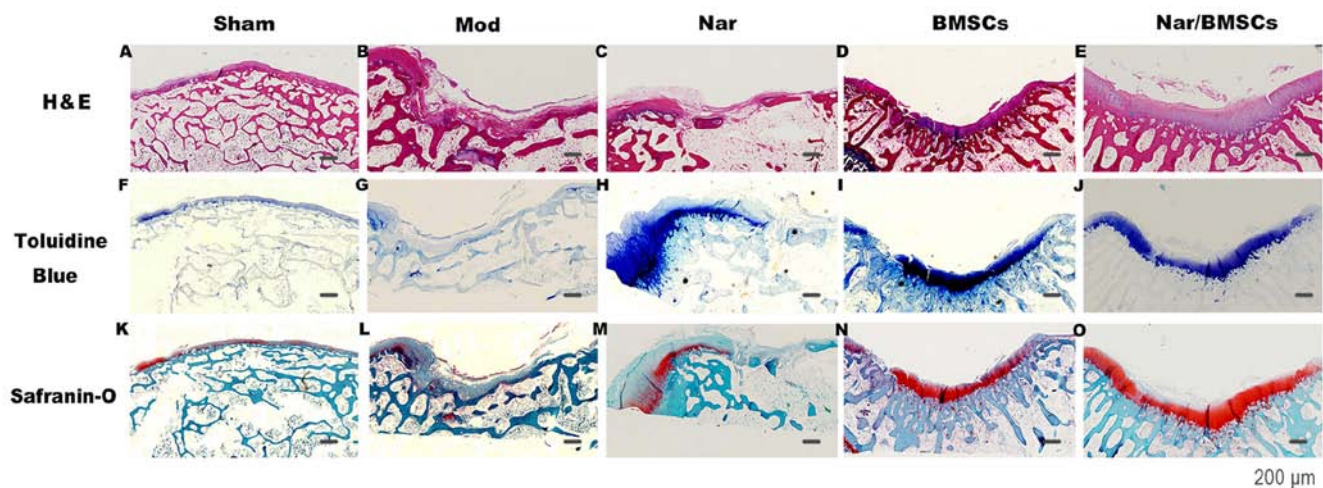


Figure 3. Histological findings of the repair tissue at articular cartilage defect sites (n=8 knees/group). Scale bars represent 200  $\mu$ m. Representative H&E staining at 12 weeks in the (A) Sham group, (B) Mod group, (C) Nar group, (D) BMSCs group and (E) Nar/BMSCs group. Toluidine blue staining at 12 weeks in the (F) Sham group, (G) Mod group, (H) Nar group, (I) BMSCs group and (J) Nar/BMSCs group. Safranin-O staining at 12 weeks in the (K) Sham group, (L) Mod group, (M) Nar group, (N) BMSCs group and (O) Nar/BMSCs group. H&E: Continuous pink on the joint surface illustrates that the cytoplasm of the cartilage is stained pink; Toluidine Blue, continuous dark blue on the joint surface indicates the presence of chondrocytes and cartilage matrix; Safranin O, continuous red on the joint surface indicates the cartilage matrix. BMSC, bone marrow mesenchymal stem cells; H&E, hematoxylin and eosin; Mod, model; Nar, naringin.

tissue of the Nar/BMSCs group appeared smooth. Differences between regenerative tissues and adjacent normal cartilage were not clear (Fig. 3E). The cartilage and cartilage matrix were evaluated using toluidine blue staining. Cells in the repair tissues of the Mod group were irregularly arranged and full of fibroblast-like cells, without signs of repaired deep tissue or the generation of glycosaminoglycan (GAG; Fig. 3G). The staining of regenerative tissue in the Nar group was clear, but repaired tissues were not completely blended with the adjacent normal cartilage. Although the repair quantity was limited, the repair quality appeared satisfactory (nuclear staining was clear, the cartilage matrix was uniformly distributed; Fig. 3H). The thickness of the stained tissue containing regenerative cells in the BMSCs group was similar to that of the surrounding cartilage. However, the junction of repaired tissue and normal cartilage was still distinguishable in this group (Fig. 3I). The regenerative tissue in the Nar/BMSCs

group stained positive and appeared homogeneous, similar to that of the adjacent normal cartilage. The boundary between the normal cartilage and regenerative cartilage was not clear and the cellular morphology and distribution in the new cartilage were almost identical to that of the normal cartilage (Fig. 3J).

Safranin O staining was used to evaluate articular cartilage repair. Joint cartilage was stained in red. The structural repair of the defect site of the BMSCs group was apparent in that Safranin O staining showed a clear structure and the articular surface was almost smooth. However, the cartilage matrix was not complete as the nuclear structure was not clear (Fig. 3N). The articular cartilage regeneration conditions of the groups administered naringin were apparent, the collagen content of the cartilage matrix was clear and there was red staining. The Nar/BMSCs treatment appeared to demonstrate the best

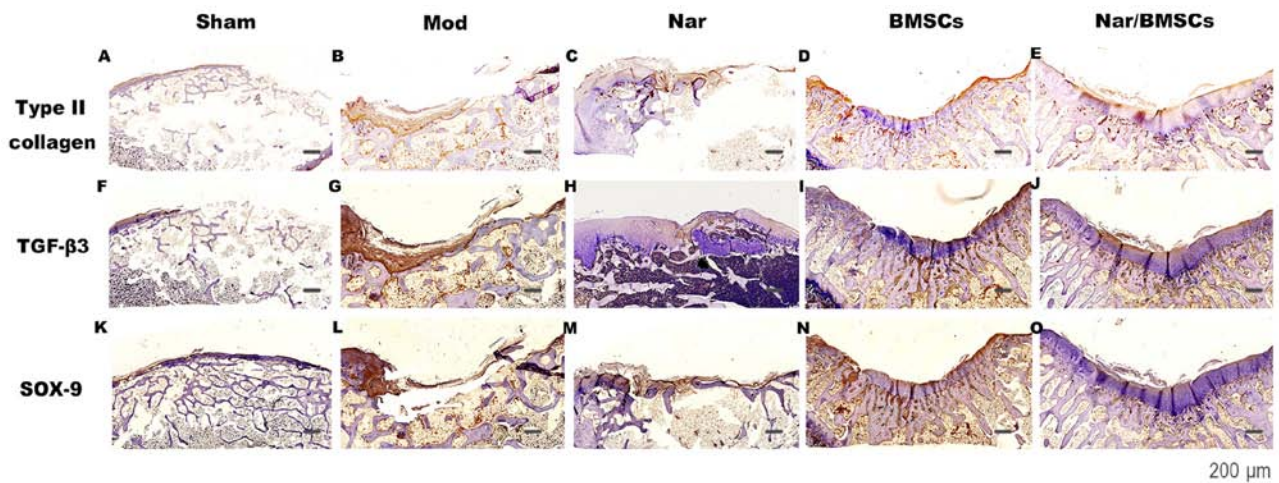


Figure 4. Immunohistological findings of the repair tissue at articular cartilage defect sites (n=8 knees/group). Scale bars represent 600  $\mu$ m. Type II collagen immunostaining at 12 weeks in the (A) Sham group, (B) Mod group, (C) Nar group, (D) BMSCs group and (E) Nar/BMSCs group. TGF- $\beta$ 3 immunostaining at 12 weeks in the (F) Sham group, (G) Mod group, (H) Nar group, (I) BMSCs group and (J) Nar/BMSCs group. SOX-9 immunostaining at 12 weeks in the (K) Sham group, (L) Mod group, (M) Nar group, (N) BMSCs group and (O) Nar/BMSCs group. Type II collagen, TGF- $\beta$ 3 and SOX-9 staining: Brown on the joint surface was positive. BMSC, bone marrow mesenchymal stem cells; Mod, model; Nar, naringin; TGF, transforming growth factor  $\beta$ .

therapeutic effect. The cartilage matrix was a uniform red and the color of the tissue structure was clear and uniform in the Nar/BMSCs group. The defect site was occupied by cartilage cells, which were highly consistent with the surrounding normal cartilage. Compared with the corresponding position of the defect in Mod group, the appearance of those areas stained in red was similar to that of adjacent normal cartilage (Fig. 3O).

*Immunohistochemical staining of the repair tissue on the joint defect site.* Type-II collagen immunostaining was used to evaluate the quality of the cartilage repair in the articular cartilage defects. The extracellular matrix of hyaline cartilage cells is mainly composed of type-II collagen and proteoglycan. TGF- $\beta$ 3 and SOX-9 are key factors in the TGF- $\beta$  superfamily signaling pathway. TGF- $\beta$ 3 is crucial in cartilage cell proliferation, differentiation and other processes. The expression profile of these markers was evaluated in Fig. 4.

In the Sham group, the expression of type-II collagen was uniform and smooth (Fig. 4A). In the Mod group, there was no significant expression of type-II collagen, with the defect having a sunken surface and uneven tissue (Fig. 4B). In comparison with the Mod group, the BMSCs group presented higher levels of type-II collagen. The Nar and Nar/BMSCs groups, which both received naringin treatment, showed more abundant type-II collagen expression than the group without naringin. The structural repair of joint defect in the Nar group appeared unsatisfactory, some of the joint defects were not repaired, but the content of type-II collagen in the repaired tissue was high, with strong positive staining. Despite a limited number of cells in the area of damage, the cells that were present were mainly regenerative cells of hyaline cartilage (Fig. 4C). Although a clear structural repair of the joint defect was exhibited in the BMSCs group, the type-II collagen content of the repaired tissue was high, with weak positive staining and a limited quantity of regenerative hyaline cartilage (Fig. 4D). The Nar/BMSCs group displayed the best therapeutic effect, the expression of type-II collagen was brownish yellow, and it was

superior to other groups in terms of the uniform distribution of lines on the articular surface. Meanwhile, the expression of type-II collagen in Nar/BMSCs group was indicated by strong positive staining and the staining intensity appeared greater than in other groups. The density of chondrocytes and cartilage matrix in the Nar/BMSCs group was similar to that of the normal cartilage. The original defects were filled with novel hyaline cartilage tissue. A possible reason for this is that the implanted BMSCs differentiated into chondrocytes (Fig. 4E).

TGF- $\beta$ 3 and SOX-9 are the main indicators of activation of the TGF- $\beta$  superfamily of signaling pathways (32). SOX-9 is regulated and controlled by the TGF- $\beta$  superfamily signaling pathway and serves as the main indicator of downstream signal transmission (33). Areas of positive staining for TGF- $\beta$ 3 in the groups without oral administration of naringin were relatively weak (Fig. 4G and I), while they were stronger in the groups with naringin treatment (Fig. 4J). SOX-9 was uniformly expressed on the surface of the joint in the Sham group (Fig. 4K). Areas of positive staining for SOX-9 in the groups without oral administration of naringin were relatively weak (Fig. 4L). The Nar group showed more SOX-9 expression than the Mod group (Fig. 4M). Although BMSCs group was implanted with bone marrow mesenchymal stem cells, the expression of SOX-9 was lower than in Nar group (Fig. 4N). While they were stronger in the groups with naringin treatment (Fig. 4O). Notably, the TGF- $\beta$ 3 and SOX9 immunohistochemical staining reaction in the Nar/BMSCs group was the most obvious in comparison with that of other groups. The staining was positive in the Nar/BMSCs group, and distributed evenly in the articular surface. The repaired articular defect structure was clear. The regenerative cell matrix of the regenerated hyaline cartilage in the defect repair site was rich in GAG. The arrangement of chondrocytes tended to be regular, and the surface of the cartilage was largely smooth. A satisfactory repair effect can be found from the staining of the above Nar/BMSCs group. (Fig. 4O).

Two professionals conducted quantitative assessment of each sample according to: (i) The International Cartilage

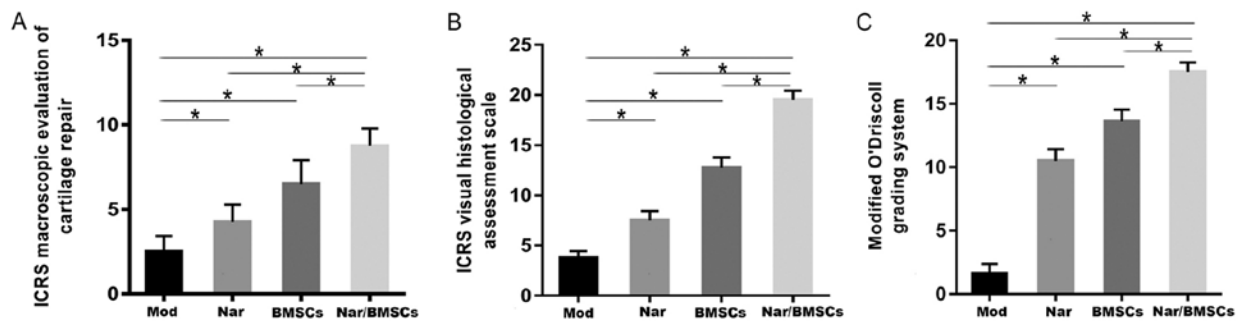


Figure 5. Histological evaluation of the tissue sections. (A) Sections were histologically evaluated based on the ICRS macroscopic evaluation of cartilage repair. (B) Sections were histologically evaluated based on the ICRS Visual Histological Assessment Scale. (C) Sections were histologically evaluated based on modified O'Driscoll grading system. Error bars represent 95% confidence interval. \* $P < 0.05$ , BMSC, bone marrow mesenchymal stem cells; ICRS, International Cartilage Repair Society; Mod, model; Nar, naringin.

Repair Society (ICRS) macroscopic evaluation of cartilage repair (Table II), (ii) The ICRS Visual Histological Assessment Scale (Table III) and (iii) the modified O'Driscoll grading system (Table IV). The main parameters assessed were the degree of defect repair, integration to the border zone and macroscopic appearance, surface, matrix and cell distribution, cell population viability, nature of the predominant tissue, structural characteristics and absence of cellular changes due to degeneration. Statistical analysis of the scores of the above items showed that the extent and quality of repair was increased in the Nar group in comparison to the Mod group; in the Nar/BMSCs group in comparison to the BMSCs group; in the BMSCs group in comparison to the Mod group and in the Nar/BMSCs in comparison to the Nar group. In comparison with the other four groups, the Nar/BMSCs group exhibited the best quality of repair, as indicated in Fig. 5.

## Discussion

Knee osteoarthritis is a common clinical disease (34). Articular cartilage injury is the main pathological factor resulting in knee osteoarthritis. Pathological changes in the morphology of the knee cartilage are thought to be the most prominent features during osteoarthritis. Knee cartilage is avascular, aneural and alymphatic, and lacks the ability to self-repair, thus the repair of articular cartilage damage is a challenge for clinicians (35). Cartilage cells serve as the main cellular components of articular cartilage and the articular cartilage surface is mainly formed by hyaline cartilage, which helps to maintain the cartilage cells in an undifferentiated state and to help the cells secrete extracellular matrix (36). The extracellular matrix mainly consists of two types of macromolecules, type-II collagen and proteoglycan.

The structure of normal cartilage when visualized with H&E staining presents a smooth cartilage surface, regular arrangement and almost round chondrocytes. The cartilage matrix under toluidine blue staining appears bluish-purple. The cartilage matrix under safranin O staining appears dark red. The collagen in cartilage tissue is mainly type II collagen. The brownish yellow part of type II collagen staining can be used as a specific locator of cartilage tissue, which indicates a satisfactory repair of cartilage following injury (37,38).

During the observation of osteoarthritis in the present study, H&E staining showed unevenly distributed cells, a

decrease in the total cell number, apoptosis of the cells and uneven staining. The density of toluidine blue staining was uneven, the staining color was uneven and discoloration was apparent. The phenomenon of light staining or loss of staining appeared in safranin O staining and the dyeing area decreased. Type-II collagen showed a decreased collagen distribution and superficial nuclear staining.

The BMSCs group displayed typical features of stem cells, which are undifferentiated cells in their original state. They possess a self-replicative ability and a multidirectional differentiation potential, existing in the bone marrow, as well as in other tissues and organs throughout life, and are responsible for the repair and updating of the tissue (39). They can be induced to differentiate into a variety of cells. As the ideal seed cells for cartilage tissue engineering, these cells have been studied from numerous perspectives (40,41). The BMSCs in this present study displayed excellent proliferative and differentiation capabilities, as well as a strong viability. This is consistent with the characteristics of BMSCs in previous experiments. BMSCs can be cultured and expanded in order to obtain seed cells, which have been used to treat cartilage defects in numerous animal models (42,43). The use of tissue engineering techniques to study the repair process following cartilage damage is considered a promising approach (2).

Naringin is a component of the TCM treatment, *Drynaria fortunei*, and is thought to be beneficial for the treatment of osteoarthritis, especially in slowing the occurrence of inflammation (44). Oral administration of naringin has been indicated to have the potential for the treatment of cartilage destruction in osteoarthritis models (45,46). As a multi-functional substance (47), naringin has been suggested to have multiple therapeutic effects, such as the capacity to inhibit the expression of enzymes which are key mediators in the development of osteoarthritis, regulate inflammatory protein secretion and promote cartilage cell infiltration (48,49). These studies highlight the potential of naringin in the treatment of osteoarticular degenerative diseases, which may help to restore the microenvironment of the cartilage defects and improve the state of the tissue. However, studies investigating the function of naringin on signaling pathways in osteoarthritis are rare.

Previous studies have clearly indicated that rabbit BMSCs and naringin solution can be applied to repair rabbit knee articular cartilage defects. The combination of both can help to achieve a better repair (20). However, the specific mechanisms

of action remain unknown; therefore, further investigation into the repair mechanisms would be of great clinical significance.

Biological processes such as the formation, metabolism and apoptosis of articular cartilage are regulated by multiple signaling pathways, among which, the TGF- $\beta$  superfamily signaling pathway is prominent in maintenance of the articular cartilage, through regulating mesenchymal cell migration, aggregation and differentiation into cartilage cells; and through regulating the formation of cartilage matrix (50). There are three types of mammalian TGF- $\beta$ , namely  $\beta$ 1,  $\beta$ 2 and  $\beta$ 3 (51). The TGF- $\beta$  superfamily signaling pathway is important in the development of the growth plate and the formation of permanent articular cartilage. Previous studies have indicated that TGF- $\beta$  plays an important regulatory role in the maintenance of the normal cartilage cell phenotype, as well as in the promotion of proliferating cells to differentiate into mature cartilage cells (52-54). Similarly, the TGF- $\beta$  signaling pathway helps to maintain the cartilage cell phenotype and the stability of the cartilage tissue environment (55). Not only in mature cartilage, but also in the process of cartilage formation, blocking the TGF- $\beta$  superfamily signaling pathway has been shown to have harmful effects (56). TGF- $\beta$ 3 has a stronger ability to induce MSCs to differentiate into cartilage than that of TGF- $\beta$ 1 (57,58). Therefore, TGF- $\beta$ 3 was selected as one of the main evaluation indexes of pathway activation in the present study.

SOX-9 can be regulated by the TGF superfamily signaling pathway. TGF- $\beta$  can promote the proliferation and differentiation of cartilage cells by maintaining SOX-9 expression levels (59). SOX-9 is thought to be one of the effective transcription factors that are regulated by type-II collagen and is deemed a modular switch of cartilage differentiation (60). SOX9 is a key transcription factor in development and is active during chondrogenic differentiation, it directly binds to enhancers in the chondrocyte genome by homodimerizing on pairs of 7-bp SOX-domain DNA recognition sites (61). Moreover, it is a key transcription factor of signaling pathways which induce cartilage differentiation of BMSCs (62). During the cartilage differentiation process, SOX-9 can be combined with the enhancer element of the type-II collagen gene to increase the expression of type-II collagen (63) and to promote the differentiation of MSCs into chondrocytes (64). The absence of SOX-9 results in the failure of MSCs to differentiate into cartilage cells, leading to damage to the cartilage primordium (65). Sufficient SOX-9 expression levels can not only facilitate the enrichment of MSCs but can also help to inhibit the conversion of proliferated cartilage cells into a mast cell phenotype. The maintenance of SOX-9 expression levels highly depends on the continuous signal stimulation of the TGF- $\beta$  superfamily signaling pathway. In summary, Sox-9 has a positive regulating function on the differentiation and maturation of the cartilage. Therefore, SOX-9 was selected as one of the downstream evaluation indexes of pathway activation in the present study (32,66).

The BMSCs group in the present study was found to repair the articular cartilage defect of the rabbit knee and the therapeutic effect on articulation structure repair was deemed satisfactory; however, the curative effect on the cartilage quality repair was poor, which was related to the inactivation of the TGF- $\beta$  superfamily signaling pathway. Compared with that

of the Mod group, the BMSC group prominently improved the degree of defect repair and integration to the border zone, the defective joint structure was restored to a certain degree. In the BMSCs group, BMSCs implanted at articular cartilage defects may repair the structures of the joint. However, restoring the structures did not appear to be sufficient and the quality of the repair was important. The quality of repair was significantly different between the regenerated tissue of the BMSCs group and the normal cartilage. The content of hyaline cartilage in the regenerated tissue was lower than that in the repaired tissue, which indicated that the BMSCs were not satisfactory in terms of the repair quality of the articular cartilage defect.

The expression levels of TGF- $\beta$ 3 and SOX-9 in the Mod and BMSCs groups were relatively low, which indicated that the BMSCs did not activate the TGF- $\beta$  superfamily signaling pathway during the repair of the articular cartilage defects in rabbit knees. Naringin, the active ingredient of the TCM treatment *Drynaria fortune* (67), was effective at improving the quality of the repair of the articular cartilage defects in the rabbit knees, which may be related to the activation of the TGF- $\beta$  superfamily signaling pathway. However, its effect on repairing the defected joint structure was not deemed satisfactory, which may be attributed to the lack of adequate repair cells.

Although the Nar group's contour repair for joint defects was stronger than that of the Mod group, the joint defects did not demonstrate a satisfactory level of repair to the structure. This could be attributed to an insufficient number of repairing seed cells in the articular cartilage defect and could also be related to a poor intrinsic healing potential (68,69). Drug treatment alone appears to be unable to significantly promote the recovery of the joint defect structure. Toluidine blue and safranin O staining revealed that, compared with that of the Mod group, the Nar group had significantly improved cartilage quality in the newly repaired tissue at the defect site. According to the results of type-II collagen staining, the newly repaired tissue was composed of predominantly hyaline cartilage. These findings demonstrated that naringin helped to promote an improved quality of cartilage regeneration and repair in articular cartilage defects of the rabbit knees.

The expression levels of TGF- $\beta$ 3 and SOX-9 were higher in the Nar group compared with those in the Mod group, which indicated that naringin stimulated the secretion of these two specific factors. TGF- $\beta$ 3 can act as the chemotactic molecule that induces chondrogenesis (70). A study into Sox9 post-translational regulation found that the Sox9 protein levels were closely related to the formation of normal cartilage (62). Moreover, SOX-9 can be used as a marker of cartilage formation in degenerated cartilage cells. These data demonstrate that naringin, the effective ingredient of the TCM treatment *Drynaria fortunei*, may be effective at promoting repair of the articular cartilage defects in rabbit knees through activation of the TGF- $\beta$  superfamily signaling pathway.

BMSCs together with naringin, can promote the repair of articular cartilage defects in rabbit knees. The repair of the joint structure defects and the quality of the repair of the articular cartilage were both deemed satisfactory and were found to be enhanced compared to that of the individual application of BMSCs or naringin. This may be associated with the simultaneous activation of the TGF- $\beta$  superfamily signaling pathway in the presence of sufficient seed cells.

Both the BMSCs and the Nar/BMSCs groups exhibited satisfactory effects on the structural repair of the joint defects. Regarding the degree of the defect repair and the association with the surrounding tissues, the Nar/BMSCs group presented greater improvements than the BMSCs group. The Nar/BMSCs group displayed better repair effects in articular cartilage regeneration and cartilage matrix repair as a result of the treatment with naringin. Regarding type-II collagen, the Nar/BMSCs group appeared identical to the surrounding normal articular cartilage tissue and the repaired tissue consisted of a considerable quantity of type-II collagen. The repaired tissue maintained a fairly smooth surface profile and its hyaline cartilage content was higher than that of the BMSCs group. This indicated that naringin may increase the directional differentiation of BMSCs into cartilage tissue in the articular cartilage defect. Furthermore, naringin improved the quality of cartilage regeneration and repair in articular cartilage defects of the rabbit knees.

The expression levels of TGF- $\beta$ 3 and SOX-9 were significantly higher in the Nar/BMSCs group compared to that in the BMSCs group. Particularly, compared to all other groups in the present study, the expression levels of TGF- $\beta$ 3 and SOX-9 were highest in the Nar/BMSCs group. This is consistent with the conclusion that Sox-9 has the potential to restore degenerated cartilage structure and functions (71).

The BMSCs implanted in the articular cartilage defects in the rabbit knees appeared to provide sufficient seed cells for repair to occur. Oral administration of naringin exerted a continuous regulatory function on the TGF- $\beta$  superfamily signaling pathway, which promoted the directional differentiation of BMSCs into cartilage tissue at articular cartilage defects in rabbit knees. Notably, the majority of the differentiated cartilage cells expressed a hyaline cartilage phenotype.

The present study demonstrated that BMSCs had a satisfactory effect on the repair of the joint structure of articular cartilage defects in rabbit knees. Naringin helped to improve the quality of cartilage repair in articular cartilage defects in rabbit knees. BMSCs combined with naringin, appeared to further improve the repair of articular cartilage defects in rabbit knees, thus delivering satisfactory results for joint structure and cartilage quality. It appears that the repair mechanism of action of naringin on articular cartilage defects may be through the continuous regulation of the TGF- $\beta$  superfamily signaling pathway and that combined application with BMSCs may produce better results than the use of naringin alone. These findings may be attributed to the directional differentiation of BMSCs into cartilage cells after regulation of the TGF- $\beta$  superfamily signaling pathway.

The present study is a preliminary study focused on articular cartilage defect repair of rabbit knees using a combination of naringin and BMSCs. Future research will investigate the effects of different concentrations of naringin on articular cartilage defect repair in rabbit knees, the target tissue of naringin (such as chondrocytes or cartilage matrix) and the direct effects of naringin on cartilage cells. Furthermore, the direct functional mechanism of action of naringin on the TGF- $\beta$  superfamily signaling pathway and the mechanism of naringin's influence on BMSCs should be further investigated.

In conclusion, naringin, an effective ingredient of the TCM *Drynaria fortunei*, and BMSCs were able to stimulate

the repair of articular cartilage defects in rabbit knees. The combined application of these treatments on joint structural repair and articular cartilage repair quality delivered improved results, which may be closely associated with sustained positive regulation of the TGF- $\beta$  superfamily signaling pathway and with the directional differentiation of BMSCs into cartilage cells.

## Acknowledgements

The authors would like to thank Professor Mingjing Zhao (Key Laboratory of Chinese Internal Medicine of Ministry of Education and Beijing, Dongzhimen Hospital, Beijing University of Chinese Medicine, Beijing, China) for technical assistance with this study.

## Funding

This study was supported by the Fundamental Research Funds for the Central Universities (grant nos. 2015-JYB-JSMS058, 2018-JYBZZ-JS091).

## 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

PL conceived and designed the study. CY, JC, HL, JY and FW performed the animal surgeries. YL, CY and PL performed the histological and immunohistochemical staining. CY, JC, HL, JY, ZY and YQ acquired the data. CY, JC, ZY and PL analyzed and interpreted data. CY and JC drafted the manuscript. PL and ZY critically revised the manuscript for important intellectual content. All authors read and approved the final manuscript.

## Ethics approval and consent to participate

The animal experiments were approved by the Science and Technology Department of Beijing University of Chinese Medicine (Beijing, China) and the Animal Ethics Review Committee of the Institute of Basic Theories of Chinese Medicine, Chinese Academy of Chinese Medical Sciences (Beijing, China) approval no. 201706058.

## Patient consent for publication

Not applicable.

## Competing interests

The authors declare that they have no competing interests.

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