# Wnt-β-catenin signaling pathway inhibition by sclerostin may protect against degradation in healthy but not osteoarthritic cartilage

JIANG WU, LONG MA, LONG WU and QUNHUA JIN

Orthopedics Ward 3, The General Hospital of Ningxia Medical University, Yinchuan, Ningxia 750004, P.R. China

Received November 7, 2015; Accepted October 4, 2016

#### DOI: 10.3892/mmr.2017.6278

Abstract. The aim of the present study was to determine the regulation of sclerostin (SOST) in osteoarthritis (OA) and its effect on articular cartilage degradation. Human cartilage samples from healthy and OA subjects were assessed by Safranin O staining and immunohistochemistry. Primary chondrocytes were pre-incubated with 250 ng/ml SOST, 10 ng/ml interleukin-1- $\alpha$  (IL-1 $\alpha$ ) or a combination of the two. The effects of treatment on the Wnt-β-catenin signaling pathway and cartilage degradation were examined by reverse transcription-quantitative polymerase chain reaction and western blotting. SOST was detected in the cartilage focal area, demonstrating secretion by osteocytes and chondrocytes. SOST has been identified to inhibit the Wnt-β-catenin signaling pathway by binding to low-density lipoprotein-related receptors 5 and 6, and catabolic factors were decreased in healthy chondrocytes. However, SOST did not influence human OA chondrocytes. IL-1α activated the Wnt-β-catenin signaling pathway and promoted cartilage degradation, which was inhibited by SOST in healthy and OA cartilage. The results of the present study suggested that SOST is important in maintaining the integrity of healthy, but not end-stage OA, cartilage.

# Introduction

Osteoarthritis (OA) is a disease affecting the joints characterized by the thinning and disintegration of cartilage, synovial inflammation and subchondral bone remodeling (1). The primary feature of OA is the gradual loss of articular cartilage (2). OA progression is complex and involves numerous processes, including aggrecan and type II collagen degradation due to increased cleavage by activated proteolytic enzymes including matrix metalloproteinases (MMPs) and a disintegrin and metalloproteinase with thrombospondin motifs (ADAMTSs) (3). Focal loss of articular cartilage in OA may be associated with alterations in the subjacent bone via modified load transmission and/or direct signaling between neighboring tissues (4).

Wnt signaling is critical for the regulation of adult bone turnoveenhancing this signaling pathway has been investigated as a potential therapeutic strategy for osteoporotic and inflammatory bone loss, to induce bone production and inhibit soluble antagonists (5). By contrast, increased Wnt-β-catenin signaling has been demonstrated to stimulate tissue degradation rather than formation in adult cartilage (6). Increased levels of  $\beta$ -catenin have been observed in chondrocytes at areas of cartilage degeneration (7), and it is upregulated in cartilage, particularly in the superficial cartilage zone (8). This may result in chondrocyte hypertrophy leading to cartilage damage (9). Furthermore, stimulation of chondrocytes with Wnt-\beta-catenin has been demonstrated to increase the expression levels of various factors involved in OA, including runt-related transcription factor 2 (RUNX-2), MMP-13, ADAMTS-4, and ADAMTS-5, facilitating cartilage matrix degradation (9).

Sclerostin (SOST), encoded by the Sost gene, is a secreted cysteine-knot protein of the differential screening selected gene abberative in neuroblastoma family, which acts as an antagonistic ligand for the Wnt coreceptors, low-density lipoprotein-related receptor (LRP)5 and LRP6, and an inhibitor of the canonical Wnt/ $\beta$ -catenin signaling pathway (10). Although the alterations in SOST in human osteoarthritic cartilage have been described (11), the complex role of SOST during OA progression remains unclear. A previous study has demonstrated that SOST is additionally expressed by chondrocytes in mineralized cartilage (12). However, the therapeutic effects of SOST in OA cartilage remain controversial. Chan et al (13) demonstrated in vitro that increased chondrocyte SOST may protect against cartilage degradation in OA, and Bouaziz et al (14) used SOST-knockout mice to reveal that the loss of SOST promotes OA in mice via β-catenin-dependent Wnt signaling pathways. However, Roudier et al (15) used SOST-knockout mice and an OA mouse model to demonstrate that SOST is expressed in articular cartilage, but its loss does not affect cartilage remodeling during aging or following mechanical injury. Whether SOST protects cartilage from degradation via inhibiting Wnt-\beta-catenin remains unknown.

*Correspondence to:* Professor Qunhua Jin, Orthopedics Ward 3, The General Hospital of Ningxia Medical University, 804 Shengli South Street, Yinchuan, Ningxia 750004, P.R. China E-mail: jinqunhua0951@yeah.net

*Key words:* cartilage, osteoarthritis, sclerostin, Wnt-β-catenin, therapy

The present study therefore used healthy and OA chondrocytes to investigate the complex role of SOST in healthy and OA cartilage.

## Materials and methods

Human samples. All human samples were obtained with the informed consent of patients, and with approval from the Ningxia Medical University Ethics Committee (Yinchuan, China). Human OA samples (OA group: n=57; female, 42; male, 15; age, 61.6±6.8 years) were obtained from patients undergoing total knee arthroplasty (TKA) for OA. Healthy human specimens (healthy group: n=6; female, 2; male, 4; age, 24.7±5.9 years) were obtained from patients undergoing lower extremity amputation due to destructive injury. X-ray films of knees were used to determine whether patients had OA. Cartilage from the medial condylar, encompassing the maximal cartilage erosion focal area, was used for primary chondrocyte culture and subsequent mRNA and protein extraction. Medial tibial plateaus, encompassing the maximal cartilage erosion focal area, were isolated from patients prior to fixing in 4% paraformaldehyde for sectioning.

Histology. Knees were fixed in 4% paraformaldehyde for 24 h at 4°C, decalcified in 0.5 M ethylenediaminetetraacetic acid at room temperature for 21 days and embedded in optimum cutting temperature compound followed by paraffin. Serial 4- $\mu$ m-thick sagittal sections of the medial tibial plateau were obtained at three depths, at 50- $\mu$ m intervals. Sections were stained with Safranin O and cartilage degradation was determined using the modified Mankin scoring system (16). Samples were divided into three groups: Normal, mid-stage OA and end-stage OA. Paraffin sections for SOST immunostaining were first dewaxed and rehydrated, then antigen retrieval was performed with 0.1% trypsase (Beijing Solarbio Science & Technology Co., Ltd.) and goat serum (Boster Bio-Engineering Ltd. Co., Wuhan, China) was used as a blocking antigen. Next, the slides were incubated with a primary rabbit anti-SOST antibody (cat. no. ab63097; Abcam, Cambridge, MA, USA) for 2 h. Subsequently, biotinylated goat anti-rabbit IgG secondary antibody [cat. no. sv0002; Boster Bio-Engineering Ltd. Co.; 1:100, diluted in PBS containing 1% bovine serum albumin (GibcThermo Fisher Scientific, Inc., Waltham, MA, USA)] was applied for 2 h at room temperature, followed by incubation for 1 h with horseradish peroxidase-conjugated antibody (cat. no. sv0002; Boster Bio-engineering Limited Company, Wuhan, China). Then the slides were colored with 3,3-diaminobenzidin (DAB) and stained with hematoxylin. Positive cells were stained yellow and counted under the light microscope (Olympus CX31, Olympus Corporation, Tokyo, Japan) on the tibial joint cartilage surface (magnification, x25) and expressed as a percentage of the total cells.

*Ex vivo cartilage explant culture*. Full-depth articular cartilage explants were isolated from the medial femoral condyle of patients undergoing TKA or amputation. Cartilage samples were cut into small pieces (~1 mm<sup>3</sup>) and digested individually with 0.25% trypsin (Beijing Solarbio Science & Technology Co., Ltd., Beijing, China) and collagenase II (Sigma-Aldrich; Merck Millipore, Darmstadt, Germany) to isolate chondrocytes.

As it is difficult to isolate chondrocytes from end-stage OA patients, due to limited cartilage on the bone surface, cartilage was not obtained from all OA patients. In total, 42 samples of OA chondrocytes (n=42) and six samples of healthy chondrocytes were cultured. All chondrocytes were cultured in 20% fetal bovine serum (GibcThermo Fisher Scientific, Inc., Waltham, MA, USA) in Dulbecco's modified Eagle's medium (GibcThermo Fisher Scientific, Inc.) for 2 days to ensure cell adhesion. Chondrocytes from healthy subjects were subcultured once at a split ratio of 1:4 using trypsin, generating 24 culture dishes of healthy chondrocytes (n=24). The medium was replaced every 3 days and activation of the Wnt-\beta-catenin signaling pathway was examined by incubating chondrocytes in the absence or presence of 10 ng/ml interleukin-1- $\alpha$  (IL-1 $\alpha$ ; PeproTech, Inc., Rocky Hill, NJ, USA) for 48 h. The effect of SOST was examined by addition of 250 ng/ml recombinant human SOST (PeproTech, Inc.) for 48 h in the absence or presence of 10 ng/ml IL-1a. All chondrocytes were subsequently prepared for mRNA and protein extractions.

RNA extraction and reverse transcription-quantitative polymerase chain reaction (RT-qPCR). RNA was extracted from human chondrocytes using the E.Z.N.A® Total RNA kit (Omega Bio-Tek, Inc., Norcross, GA, USA), and quantified with a NanoDrop 2000 spectrophotometer (Thermo Fisher Scientific, Inc.). Next, RT was performed using RevertAid First Strand cDNA Synthesis kit (Thermo Fisher Scientific, Inc.). The temperature protocol for RT was as follows: 25°C for 5 min; 42°C for 60 min, 70°C for 5 min for termination as described in the kit manual. mRNA expression levels were quantified by qPCR using a SYBR® Green Master mix (Thermo Fisher Scientific, Inc.) and a LightCycler<sup>®</sup> 480 (Roche Diagnostics, Basel, Switzerland). The temperature protocol for the reaction was as follows: 95°C for 10 min, 40 cycles at 95°C for 15 sec, drawing melting curve at 65°C for 15 sec, 60°C for 1 min, 95°C for 15 sec and 60°C for 15 sec. The primers used are presented in Table I. Averaged quantification cycle (Cq) values were normalized to the averaged Cq value of  $\beta$ -actin. Adjusted average Cq values were used to calculate relative expression vs. the control, using the  $2^{-\Delta\Delta Cq}$  method (17).

Western blot analysis. Chondrocyte proteins were extracted using a Whole Cell Lysis assay kit (Nanjing KeyGen Biotech Co., Ltd., Nanjing, China) and quantified using the bicinchoninic acid assay. Membranes were probed with the following primary antibodies: anti-SOST (1:1,000; cat. no. ab63097), anti-β-catenin (1:1,000; cat. no. ab6302), anti-MMP-13 (1:2,000; cat. no. ab39012) and anti-ADAMTS-4 (1:1,000; cat. no. ab84792; Abcam). Protein concentration was determined using the bicinchoninc acid protein assay kit (Boster Bio-engineering Limited Company, Wuhan, China). Equal quantities of protein (20  $\mu$ g) were separated by 10% SDS-PAGE and electroblotted onto a PVDF membrane (EMD Millipore, Billerica, MA, USA). The membrane was blocked in PBS containing 0.1% Tween-20 (Beyotime Institute of Biotechnology, Inc., Haimen, China) and 5% non-fat dry milk. The membrane was then probed with primary antibodies for overnight incubation at 4°C. Horseradish peroxidase-conjugated rabbit anti-human secondary antibody (cat. no. BA1070; 1:2,000; Boster Bio-engineering Limited Company) was then



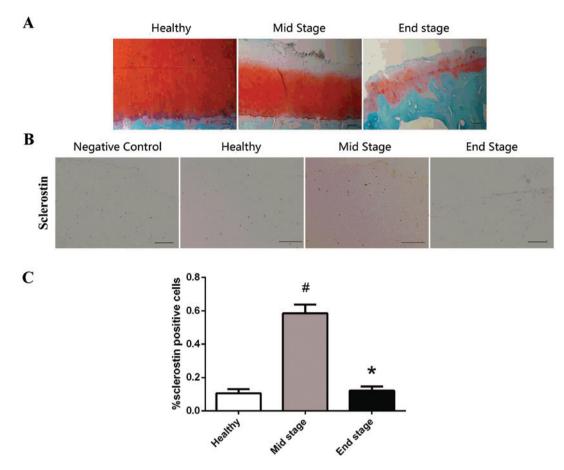


Figure 1. Expression of SOST in cartilage of OA patients and healthy controls. (A) Safranin O staining reveals the progression of cartilage lesions in the medial tibial plateau of healthy and OA cartilage. Scale bar=500  $\mu$ m. (B) Immunohistochemical analysis of SOST in calcified cartilage. Scale bar=200  $\mu$ m. (C) Percentage of SOST-positive cells in cartilage of tibial plateau. The percentage of SOST-positive chondrocytes in the mid-stage OA group was significantly increased compared with the healthy and end-stage OA groups. \*P<0.05 vs. healthy; \*P<0.05 vs. mid-stage. SOST, sclerostin; OA, osteoarthritis.

added for 1 h at 26°C . Finally, the protein bands were detected using ECL solution (GE Healthcare Life Sciences, Shanghai, China) and images were captured using a FluorChem imaging system (Alpha Innotech, San Leandro, CA, USA).  $\beta$ -actin was used as a loading control. Unfortunately, western blot analysis was not performed on healthy group chondrocytes. Only six healthy cartilage samples were used, which were not subcultured to the third generation or cultured for a prolonged time in case of cytometaplasia and apoptosis. Therefore, not enough cells were obtained for western blotting and SOST,  $\beta$ -catenin, MMP-13 and ADAMTS-4 protein expression levels were detected only in OA human chondrocytes.

Statistical analysis. Data are presented as the mean  $\pm$  standard deviation, and were compared using SPSS software (version 22.0; IBM SPSS, Armonk, NY, USA). Two-way analysis of variance was used to compare the percentage of SOST-positive cells between healthy and OA cartilage groups. Alterations in the number of positively stained cells, and the fold change in gene expression are presented graphically as means with 95% confidence intervals. The protein expression levels in different groups were compared using a Student's *t*-test. As certain data, for example gene expression, were not normally distributed, treatment effects were assessed using the non-parametric Mann-Whitney U test. P<0.05 was considered to indicate a statistically significant difference.

#### Results

Expression of SOST is not permanently increased in cartilage during OA progression. The Mankin score of the healthy group (n=6) was 1.3±1.2. Cartilage samples obtained from OA patients (n=57) were divided into two groups, with scores for the mid-stage OA group (n=37) being 7.7±0.9 and the end-stage OA group (n=20) being 13.0±1.2 (Fig. 1A). Immunohistochemical staining was performed to detect the expression of SOST at different stages of OA (Fig. 1B). Few SOST-positive stained chondrocytes were observed in the healthy and end-stage OA groups, with focal localization in the calcified cartilage and deep cartilage near the tidemark. However, the percentage of SOST-positive chondrocytes in the mid-stage OA group was significantly increased compared with the healthy (P=0.002) and end-stage OA groups (P=0.001) (Fig. 1C). No differences were observed between the healthy and end-stage OA groups (P=0.48); therefore, SOST was not permanently increased during OA progression, instead increasing and subsequently decreasing. A high number of positively stained chondrocytes were observed in the calcified cartilage of the healthy group; however, only a few were present in the deep cartilage and none on the surface. Conversely, positively stained chondrocytes were widely distributed in the surface, deep and calcified cartilage in mid-stage OA cartilage, with low SOST expression in the calcified cartilage in end-stage OA samples.

Gene	Accession no.	Tm (°C)	Product size (bp)	Forward	Reverse
SOST	NM_002427.3	60	126	TTCTCCTTCGGGACCTCAAT	TCTCTCACCTCTGCCCATTC
β-catenin	NM_005099.4	58	122	GCTTTGTGTCGTCTTGAACG	TCAGCAATCCCTTTCTCACC
LRP5	AB257751.1	58	140	TTCATCTACTGGACCGACTGG	TTGGTTCCGACGACCTTG
LRP6	AB257752.1	59	114	ATCAAGCACCAAAGGCACT	GTGGAAGGGCTGTTAGAAGAA
RUNX-2	NM_001024630.3	60	100	GACGAGGCAAGAGTTTCACC	GGTTCCCGAGGTCCATCTAC
MMP-13	NM_001012329.1	60	129	TATGACTATGCGTGGCTGGA	CCATTTGTGGTGTGGGAAGT
ADAMTS-4	NM_005099.4	59	126	GGCTGTGATCGCATCATTGG	CACATTGTTGTATCCGTACCTGA
ADAMTS-5	NM_007038.3	59	172	TGTGAAGAGACCTTTGGTTCC	TTCTGTGATGGTGGCTGAAG
COL2A1	X02420	60	195	CATTCATCCCACCCTCTCAC	TTCCTGTCTCTGCCTTGACC
β-actin	NM_001101	59	204	TGACGTGGACATCCGCAAAG	CTGGAAGGTGGACAGCGAGG-3

Table I. Primer sec	nuences used for	polymerase chain	reaction analysis.

Tm, annealing temperature; SOST, sclerostin; LRP, low-density lipoprotein-related receptor; RUNX-2, runt-related transcription factor 2; MMP, matrix metalloproteinase; ADAMTS, a disintegrin and metalloproteinase with thrombospondin motifs; COL2A1, collagen type II alpha 1 chain.

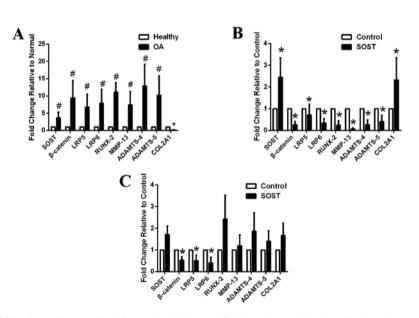


Figure 2. Expression levels of Wnt- $\beta$ -catenin associated factors in healthy and OA chondrocytes incubated in the absence or presence of SOST. (A) The Wnt- $\beta$ -catenin signaling pathway was overactivated in OA chondrocytes (n=6) compared with healthy chondrocytes (n=6). (B) SOST treatment increased the mRNA expression levels of the anabolic marker, COL2A1, but inhibited the mRNA expression levels of the Wnt signaling-associated factors,  $\beta$ -catenin, LRP5 and LRP6 and catabolic markers, RUNX-2, MMP-13 and ADAMTS-4,5 in healthy chondrocytes (n=6). (C) SOST inhibited the mRNA expression levels of the Wnt signaling-associated factors,  $\beta$ -catenin, LRP5 and LRP6 in OA chondrocytes, but did not influence downstream factors (n=6). (D) Western blot analysis of OA chondrocytes incubated with SOST and/or IL-1 $\alpha$ . <sup>#</sup>P<0.05 vs. healthy chondrocyte <sup>\*</sup>P<0.05 vs. control. OA, osteoarthriti SOST, sclerostin; LRP, low-density lipoprotein-related recepto; RUNX-2, runt-related transcription factor 2; MMP, matrix metalloproteinas; ADAMTS, a disintegrin and metalloproteinase with thrombospondin motif; COL2A1, collagen type II alpha 1 chain.

Wht- $\beta$ -catenin signaling is overactivated in OA chondrocytes.  $\beta$ -catenin (P=0.011), LRP5 (P=0.023) and LRP6 (P=0.012) mRNA expression levels were upregulated in OA chondrocytes, suggesting that the Wht- $\beta$ -catenin signaling pathway was overactivated and may be important in cartilage degradation (Fig. 2A). In addition, mRNA expression levels of the cartilage catabolic factors, RUNX-2 (P=0.015), MMP-13 (P=0.015) and ADAMTS-4 (P=0.008), ADAMTS-5 (P=0.002) were upregulated, whereas the anabolic factor, COL2A1 (P=0.003) was downregulated, in the OA compared with the healthy chondrocyte group, thus accelerating cartilage matrix breakdown (Fig. 2A). Furthermore, SOST (P=0.021) mRNA expression levels were increased in the OA group, consistent with the results of the immunohistochemical analysis. Increased SOST expression may be a reaction to the activation of the Wnt- $\beta$ -catenin signaling pathway and may inhibit the overactivated signaling pathway to maintain the integrity and normal structure of cartilage. If this is the case, SOST may be a potential therapeutic to delay OA progression.

SOST inhibits the Wnt- $\beta$ -catenin signaling pathway in healthy and OA chondrocytes, with beneficial effects observed in



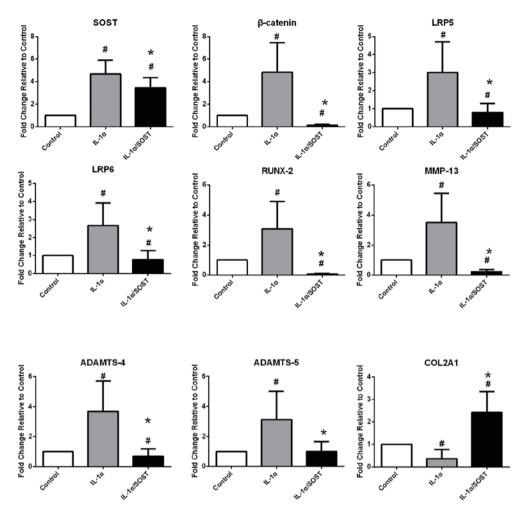


Figure 3. mRNA expression levels of Wnt- $\beta$ -catenin-associated genes in healthy chondrocytes incubated with IL-1 $\alpha$  in the absence or presence of SOST. mRNA expression levels of the Wnt- $\beta$ -catenin-associated factors,  $\beta$ -catenin and LRP5/6 and the catabolic markers, RUNX-2, MMP-13 and ADAMTS-4,5 were increased, whereas the anabolic marker, COL2A1 was decreased following IL-1 $\alpha$  treatment in healthy chondrocytes. These effects were inhibited by SOST (n=6). The increasing of ADAMTS-5 by IL-1 $\alpha$  was inhibited by SOST, but has no difference with the control group after the decreasing. \*P<0.05 vs. control; \*P<0.05 vs. IL-1 $\alpha$ . IL-1 $\alpha$ , interleukin-1- $\alpha$ ; SOST, sclerostin; LRP, low-density lipoprotein-related recepto; RUNX-2, runt-related transcription factor 2; MMP, matrix metalloproteinas; ADAMTS, a disintegrin and metalloproteinase with thrombospondin motif; COL2A1, collagen type II alpha 1 chain.

healthy chondrocytes only. SOST, as an inhibitor of the Wnt- $\beta$ -catenin signaling pathway, may decrease the expression of downstream factors, including MMPs and ADAMTSs, to protect cartilage from degradation. Healthy (Fig. 2B) and OA (Fig. 2C) chondrocytes were incubated with 250 ng/ml SOST for 48 h; this decreased β-catenin and LRP5/6 mRNA expression levels in the two groups.  $\beta$ -catenin (healthy group, P=0.012: OA group, P=0.022), LRP5 (healthy group, P=0.026; OA group, 0.0016) and LRP6 (healthy group, P=0.015; OA group, P=0.013). Therefore, SOST may inhibit the Wnt-β-catenin signaling pathway by binding to LRP5/6 in chondrocytes. In addition, SOST decreased RUNX-2 (P=0.015), MMP-13 (P=0.006) and ADAMTS-4 (P=0.013), ADAMTS-5 (P=0.015), and increased COL2A1 (P=0.002) mRNA expression levels in healthy chondrocytes (Fig. 2B). This indicated that SOST may assist in the maintenance of cartilage integrity and reduce cartilage damage. Although SOST treatment decreased β-catenin and LRP5/6 mRNA expression levels, it did not influence RUNX-2 (P=0.065), MMP13 (P=0.083), ADAMTS4 (P=0.074), ADAMTS-5 (P=0.063) and COL2A1 (P=0.068) mRNA expression levels in OA chondrocytes compared with the control group (Fig. 2C). This indicated that SOST may not have beneficial effects on OA chondrocytes despite decreasing  $\beta$ -catenin expression levels via binding to LRP5/6.

IL-1a may regulate cartilage degradation via activation of the Wnt- $\beta$ -catenin signaling pathway; this may be inhibited by SOST. Chondrocytes were incubated with 10 ng/ml IL-1α for 48 h to simulate the inflammatory environment, determine the activation effect on the Wnt- $\beta$ -catenin signaling pathway and quantify the effects of IL-1 $\alpha$  on chondrocytes. Following treatment with IL-1 $\alpha$  in the absence or presence of SOST, RT-qPCR was performed on healthy (Fig. 3) and OA (Fig. 4) chondrocytes and western blotting was performed on OA chondrocytes to detect protein expression of SOST,  $\beta$ -catenin, ADAMTS-4 and MMP-13 (Fig. 5). The mRNA expression levels of β-catenin (P=0.002), LRP5 (P=0.004) and LRP6 (P=0.014) in healthy (Fig. 3) were increased following treatment with IL-1a. Furthermore, mRNA expression levels of the cartilage catabolic factors, RUNX-2, MMP13 and ADAMTS4/5 were significantly increased, and the anabolic factor, COL2A1 was decreased, following treatment with IL-1 $\alpha$  in healthy (P-values were 0.015, 0.009, 0.022, 0.012;

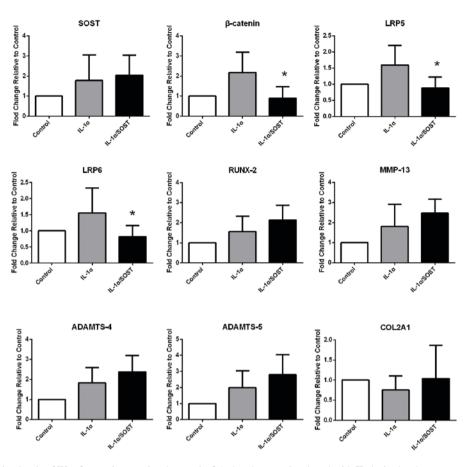


Figure 4. mRNA expression levels of Wnt- $\beta$ -catenin-associated genes in OA chondrocytes incubated with IL-1 $\alpha$  in the absence or presence of SOST. mRNA expression levels of the Wnt- $\beta$ -catenin-associated factors,  $\beta$ -catenin and LRP5/6 were not influenced by IL-1 $\alpha$  in OA chondrocytehowever, the expression of these factors was inhibited by SOST. Catabolic and anabolic markers were not influenced by IL-1 $\alpha$  alone or with SOST (n=6). \*P<0.05 vs. IL-1 $\alpha$ . OA, osteo-arthriti; IL-1 $\alpha$ , interleukin-1- $\alpha$ ; SOST, sclerostin; LRP, low-density lipoprotein-related recepto; RUNX-2, runt-related transcription factor 2; MMP, matrix metalloproteinas; ADAMTS, a disintegrin and metalloproteinase with thrombospondin motif; COL2A1, collagen type II alpha 1 chain.

Fig. 3) but not OA (P-values were 0.62, 0.53, 0.71, 0.53; Fig. 4) chondrocytes. These results indicated that IL-1 $\alpha$ may accelerate the degradation of cartilage by activating the Wnt- $\beta$ -catenin signaling pathway in healthy but not OA cartilage. Furthermore, chondrocytes exposed to IL-1 $\alpha$ demonstrated increased mRNA expression levels of SOST in the healthy (P=0.002) but not the OA group (P=0.56). Chondrocytes were treated with a combination of 250 ng/ml SOST and 10 ng/ml IL-1a to investigate whether SOST inhibited Wnt-β-catenin following overactivation. Compared with IL-1 $\alpha$  only, IL-1 $\alpha$  plus SOST treatment decreased  $\beta$ -catenin and LRP5/6 mRNA expression levels in healthy (P=0.002, 0.012, 0.012) and OA chondrocytes (P=0.022, 0.002, 0.015) and decreased RUNX-2, MMP-13 and ADAMTS4/5, and increased COL2A1 in healthy (P=0.023, 0.009, 0.013, 0.016; Fig. 3) but not OA (P=0.68, 0.57, 0.55; Fig. 4) chondrocytes. SOST therefore inhibited overactivation of the Wnt-β-catenin signaling pathway in healthy and OA chondrocytehowever, it did not decrease the expression of downstream catabolic factors to induce 'anti-catabolic' effects on OA chondrocytes.

### Discussion

The etiology and progression of OA remain to be fully elucidated owing to the complexity of the diseastherefore, the role of SOST in cartilage and subchondral bone during degradation requires further investigation. Whether SOST protects cartilage from degradation remains controversial. SOST is typically expressed in osteocytes, particularly mature osteocytes, which are surrounded by mineralized matrix (18,19). However, later studies have suggested that SOST is expressed by hypertrophic chondrocytes in calcified cartilage around the growth plate (20-23). Full-depth articular cartilage explants from the trochlear groove of ovine knee joints were used to examine the activation of IL-1 $\alpha$  in the Wnt- $\beta$ -catenin signaling pathway; chondrocytes were revealed to express SOST, which is regulated by IL-1 (24). Increased SOST in human chondrocytes may inhibit the Wnt-β-catenin signaling pathway and downstream MMPs and ADAMTS in vivo, thus protecting cartilage from degradation (13). Bouaziz et al (14) demonstrated using destabilization of the medial meniscus (DMM) mice that SOST is only secreted by calcified matrix-embedded cells, and is increased during the development of OA but decreased in end-stage OA. In addition, this study demonstrated using SOST-knockout mice that loss of SOST accelerates degradation of cartilage in OA. However, Roudier et al (15) revealed using human articular cartilage that SOST is increased in OA cartilage, and that loss of SOST in the joint enhanced the bone mass of subchondral bonhowever, there was no difference between the cartilage



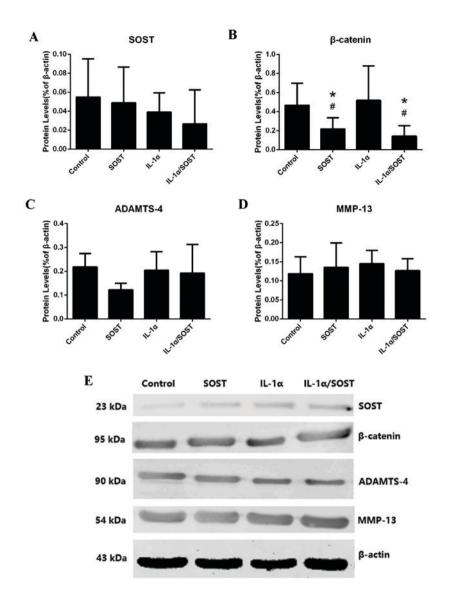


Figure 5. Protein expression levels of Wnt- $\beta$ -catenin-associated genes in OA chondrocytes incubated with IL-1 $\alpha$  in the absence or presence of SOST. (A) Protein levels of SOST in OA chondrocytes was not influenced by IL-1 $\alpha$  (P=0.76), SOST (P=0.43) and IL-1 $\alpha$ /SOST (P=0.17; n=6) compared with control group (n=6). (B) Wnt- $\beta$ -catenin-associated factors,  $\beta$ -catenin were influenced by SOST (P=0.035) and IL-1 $\alpha$ /SOST (P=0.024) in OA chondrocytes, but not influenced by IL-1 $\alpha$  (P=0.15). (C) Protein levels of ADAMTS-4 in OA chondrocytes was not influenced by IL-1 $\alpha$  (P=0.56), SOST (P=0.23) and IL-1 $\alpha$ /SOST (P=0.31; n=6) compared with control group (n=6). (D) Protein levels of MMP-13 in OA chondrocytes were not influenced by IL-1 $\alpha$  (P=0.13), SOST (P=0.21) and IL-1 $\alpha$ /SOST (P=0.37; n=6) compared with control group (n=6). (E) Western blot analysis of OA chondrocytes incubated with SOST and/or IL-1 $\alpha$ . <sup>#</sup>P<0.05 vs. IL-1 $\alpha$ /SOST. (A) Protein levels of ADAMTS, a disintegrin and metalloproteinase with thrombospondin motif; COL2A1, collagen type II alpha 1 chain.

of SOST-knockout mice and DMM mice. Furthermore, increased SOST in cartilage did not affect cartilage remodeling during aging or following mechanical injury.

The present study demonstrated that SOST was expressed only in the focal area of calcified cartilage and deep cartilage adjacent to the tidemark in the medial tibial plateau of healthy and end-stage OA human articular cartilage. The expression of SOST was significantly increased in mid-stage OA and the positively stained chondrocytes were closer to the surface of cartilage compared with the healthy and end-stage OA groups. This is consistent with the study by Bouaziz *et al* (14) using DMM mice. SOST was not continuously increased in cartilage during the development of OA, instead first increasing and subsequently decreasing, which is in contrast to previous studies (13,25,26). In early-stage OA, SOST in the cartilage may be secreted by osteocytes in the subchondral bone as a result of stimuli, including mechanical loading, but is not secreted by chondrocytes. Furthermore, in the present study almost every hypertrophic chondrocyte was positively stained. SOST therefore may not be secreted by healthy chondrocytes but by hypertrophic ones. In early-stage OA, chondrocytes in the cartilage did not express SOST; therefore, it must be secreted by osteocytes in the subchondral bone, from which it penetrates into cartilage via microchannels or vessels. Therefore, SOST-positive chondrocytes were observed only in the calcified cartilage. In mid-stage OA, the chondrocytes became hypertrophic and began to secrete SOST, resulting in a significant increase in SOST expression compared with early-stage OA (27). In end-stage OA, SOST expression was significantly reduced compared with mid-stage OA. This is not consistent with some previous studies and further investigation is required to confirm these results.

In the present study, the Wnt- $\beta$ -catenin signaling pathway was overactivated in OA chondrocytes, and matrix breakdown factors including MMPs and ADAMTSs were upregulated. Inhibiting the Wnt-β-catenin signaling pathway may decrease these mediators and therefore be a potential therapeutic approach for the treatment of OA. SOST has been identified to inhibit the Wnt-β-catenin signaling pathway by binding to LRP5/6 in healthy and OA chondrocytehowever, it decreased expression levels of cartilage catabolic factors only in healthy chondrocytes, with no beneficial effects observed on OA chondrocytes. This finding is inconsistent with previous studies in animal OA models (13,26). It may be that SOST expression is affected by mechanical loading in osteoblasts in subchondral bone, and alterations of load bearing on cartilage may stimulate signal transduction between subchondral bone and cartilage. Due to the weight-bearing diversity across different areas of cartilage within the joint, SOST expression and its effect may vary between areas (28). As there is a marked difference in walking and limb alignment between humans and rodents, mechanical loading in the same area of cartilage within the joint may cause different consequences of gene expression, regulation and transduction of factors in the Wnt-β-catenin signaling pathway (29). In previous studies using rodent OA models, different experimental results may be due to a number of reasons. For example, using different joint cartilage regions for quantification due to the small scale and blurred boundaries of different parts of rodent joints, ignoring species specificity, in particular mechanical loading within the joint, and interactions of different signaling pathways. To avoid these errors, the present study used cartilage from the medial femoral condyle for chondrocyte isolation and primary culture to determine the effects of SOST on cartilage, and the medial tibial plateau for paraffin sectioning and immunohistochemistry to measure SOST expression in cartilage at different stages of OA. However, only six samples of healthy human cartilage were obtained. To generate sufficient chondrocyte numbers, cells were subcultured at a ratio of 1:4 to obtain 24 bottles of chondrocytes in the healthy group. Due to differing conditions in vivo and in vitro, chondrocytes were passaged only to the first generation in case of cytometaplasia and apoptosis. Consequently, not enough chondrocytes were obtained for western blot analysis, which may lead to inaccurate results within the present study. SOST may therefore only be a precautionary measure to prevent cartilage degradation, and not a potential therapeutic strategy to restore integrity or healthy cartilage structure during OA progression.

In addition, the route of injecting SOST may have a marked impact on whether SOST may be a potential therapeutic in OA. Intraperitoneal injection of SOST as a systemic drug delivery method would be convenient and effective. However, there is a very limited blood supply within cartilage tissue, although certain microchannels and micrangium penetrate calcified cartilage and the tidemark to deliver mediators between cartilage and subchondral bone. Whether SOST is small enough to access these channels remains unknown; therefore, the concentration of SOST in cartilage may not be enough to elicit an effect. Furthermore, SOST has been reported to inhibit osteoblast differentiation, proliferation and activity, resulting in reduced osteoblastic bone formation (30), which may lead to decreased subchondral bone stiffness and increased mechanical stress in cartilage. Intra-articular injection may have an advantage in maintaining the concentration of SOST; however, whether SOST permeates cartilage into the calcified region and subchondral bone remains unknown. It is currently difficult to state which drug-delivery method is more suitable.

To determine whether SOST inhibited overactivation of the Wnt- $\beta$ -catenin signaling pathway in healthy and OA chondrocytes, chondrocytes were incubated with IL-1 $\alpha$  to activate Wnt signaling. IL-1 $\alpha$  treatment increased mRNA expression levels of  $\beta$ -catenin and downstream catabolic factors, including MMP-13 and ADAMTS-4/5, in healthy chondrocytes, and SOST inhibited this phenomenon. Notably, IL-1 $\alpha$  and SOST did not influence OA chondrocytes. This may be due to the reaction to the simulation of inflammatory factors of hypertrophic chondrocytes in OA not being as sensitive as healthy chondrocytehealthy chondrocytes may react to inflammatory stimuli and thus contribute to the maintenance of the normal structure and integrity of the cartilage, whereas hypertrophic chondrocytes may have lost this ability.

However, the results of the present study obtained by comparing SOST expression at different stages of OA and the incubation of chondrocytes with SOST ex vivo may not be accurate. All structures within the joint are affected and may interact during the progression of OA, including the cartilage, subchondral bone and synovium. Chondrocytes and cells in the bone may react independently to identical environmental stimuli, leading to altered cell phenotypes in OA. Alternatively, cellular alterations in one cell type may have an affect on another cell type. Previous in vitro and in vivo studies have demonstrated that chondrocytes and osteoblasts influence each other (31,32). Extensive research has demonstrated that cartilage and subchondral bone are not separate in the progression of OA, therefore the progression of the disease should be considered in the context of the interaction of these two compartments (33,34). Recently, molecular crosstalk between osteoblasts/osteocytes and chondrocytes has been demonstrated, revealing a previously unappreciated complexity (35). The dense subchondral vasculature in close proximity to the cartilage and the microchannels that infiltrate the subchondral mineralization region has been revealed to allow communication between bone and cartilage (36). Uncalcified cartilage may be observed dipping through the calcified cartilage into the bone and marrow spaces, which may provide a molecular diffusion pathway with potential nutritional, metabolic and biomechanical effects. In addition, as this interface is involved in OA, these areas may enable trafficking of humoral mediators between tissues (37). In the present study, no crosstalk was observed between subchondral bone and cartilage as it is challenging to simulate the physical and chemical environments within the joint. As alterations in one compartment may influence the other in OA, the normalization of cells in one compartment may have beneficial effects on the other. The two compartments may

subsequently influence each other to create a more favorable cycle, thus delaying the progression of OA.

Although the present study revealed that SOST benefited healthy but not OA chondrocytes, this does not mean that SOST may not influence chondrocytes, owing to the complexity of mediator translation between cartilage and subchondral bone. One possibility is that in the presence of SOST or as a result of pharmacologic inhibition, a compensatory molecule, for example another Wnt signaling inhibitor, is upregulated in the cartilagalternatively, the crosstalk between cartilage and subchondral bone may mask the effect of SOST inhibition. Further studies are required to clarify the complex role of the Wnt signaling pathway in OA and the interaction of SOST in the two compartments, which may contribute to an improved understanding of the pathogenesis and future therapies of human OA.

In conclusion, the results of the present study demonstrated that SOST is secreted by chondrocytes at various stages of OA and is not permanently increased in cartilage, first being increased and subsequently decreased. Furthermore, SOST inhibited the Wnt- $\beta$ -catenin signaling pathway in healthy and OA chondrocytehowever, SOST downregulated catabolic factors to benefit only healthy chondrocytes. SOST may prevent cartilage degradation; however, it may be that SOST cannot restore the integrity or normal structure of cartilage during OA progression.

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