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Inhibition of β-Catenin Signaling in Articular Chondrocytes Results in Articular Cartilage Destruction

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Abstract

Objective—Osteoarthritis is a degenerative joint disease whose molecular mechanism is currently unknown. Wnt/ β -catenin signaling has been demonstrated to play a critical role in the development and function of articular chondrocytes. To determine the role of β -catenin signaling in articular chondrocyte function, we generated *Col2a1-ICAT*–transgenic mice to inhibit β -catenin signaling in chondrocytes.

Methods—The expression of the *ICAT* transgene was determined by immunostaining and Western blot analysis. Histologic analyses were performed to determine changes in articular cartilage structure and morphology. Cell apoptosis was determined by TUNEL staining and the immunostaining of cleaved caspase 3 and poly(ADP-ribose) polymerase (PARP) proteins. Expression of Bcl-2, Bcl-x_L, and Bax proteins and caspase 9 and caspase 3/7 activities were examined in primary sternal

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AUTHOR CONTRIBUTIONS

Analysis and interpretation of data. Zhu, M. Chen, Rosier, O'Keefe, D. Chen.

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Dr. D. Chen had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

Study design. D. Chen.

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Statistical analysis. Zhu.

chondrocytes isolated from 3-day-old neonatal *Col2a1-ICAT*–transgenic mice and their wild-type littermates and in primary chicken and porcine articular chondrocytes.

Results—Expression of the *ICAT* transgene was detected in articular chondrocytes of the transgenic mice. Associated with this, age-dependent articular cartilage destruction was observed in *Col2a1-ICAT*– transgenic mice. A significant increase in cell apoptosis in articular chondrocytes was identified by TUNEL staining and the immunostaining of cleaved caspase 3 and PARP proteins in these transgenic mice. Consistent with this, Bcl-2 and Bcl-x_L expression were decreased and caspase 9 and caspase 3/7 activity were increased, suggesting that increased cell apoptosis may contribute significantly to the articular cartilage destruction observed in *Col2a1-ICAT*–transgenic mice.

Conclusion—Inhibition of β -catenin signaling in articular chondrocytes causes increased cell apoptosis and articular cartilage destruction in *Col2a1-ICAT*–transgenic mice.

Osteoarthritis (OA) is a degenerative joint disease. Numerous genetic and environmental factors have been proposed as contributors to the development of OA. The progression of the disease is slow and eventually results in degeneration and loss of the articular cartilage in various joints, including fingers, knees, hips, and spine. The disease process leads to limitation of joint movement, joint deformity, joint stiffness, inflammation, and severe pain. The mechanism of OA pathogenesis remains undefined.

OA is mediated by a dynamic interplay between the articular chondrocytes, their matrix, and synovial cells. Articular chondrocytes, the cell type present in articular cartilage, are responsible for producing and maintaining the extracellular matrix (ECM), which supports the appropriate biomechanical function of the tissue. Phenotypic changes seen in OA are often accompanied by ECM degradation, which eventually results in cartilage destruction (1). Synovial fibroblasts are mesenchyme-derived cells that form a thin lining of synovial tissue surrounding the fibrous capsule of the joint. The physiologic roles of synovial tissue are to produce synovial fluid that lubricates the joints and to supply nutrients to the articular chondrocytes.

Articular cartilage plays a crucial role in joint function during adulthood, and destruction of articular cartilage has severe consequences. While most cartilages are replaced by bone during development, only a few cartilages are permanent, including articular cartilage. Articular chondrocytes, which function to maintain the matrix, are unique among growth plate chondrocytes in that they exhibit very limited mitotic activity, have a slow rate of matrix synthesis and degradation, and do not progress fully to the terminally differentiated phenotype displayed by cells in the growth plate (2). Very little is known regarding the mechanisms involved in establishing and maintaining this articular chondrocyte phenotype.

The function of articular chondrocytes is regulated by a variety of growth factors, including Wnt/ β -catenin signaling molecules, which play a critical role in chondrocyte proliferation, differentiation, and apoptosis. Beta-catenin is a key molecule in the canonical Wnt signaling pathway and plays a critical role in multiple steps during chondrocyte formation and maturation (3,4). Inhibitor of β -catenin and T cell factor (ICAT) is an 82-amino-acid small molecule (5) whose crystal structure reveals binding capacity to the armadillo repeats of β -catenin. This binding disrupts the complex formation of β -catenin with T cell factor (TCF)/lymphoid enhancer factor (LEF) (6,7) and thus leads to inhibition of signaling in this pathway. To investigate the role of β -catenin signaling in the maintenance of articular cartilage function under physiologic and pathophysiologic conditions, we generated *Col2a1-ICAT*-transgenic mice in which the *ICAT* transgene was targeted specifically to chondrocytes using the *Col2a1* promoter. We found that the *ICAT* transgene was highly expressed in articular chondrocytes in 6-month-old transgenic mice, leading to inhibition of β -catenin signaling in transgenic mouse chondrocytes. Associated with this, severe articular cartilage destruction was

observed in *Col2a1-ICAT*–transgenic mice. Articular chondrocyte apoptosis was significantly increased, providing novel evidence regarding the contribution of β -catenin signaling to the maintenance of normal articular cartilage function.

MATERIALS AND METHODS

Generation of transgenic mice

To generate the *Col2a1-ICAT* transgene, DNA fragments encoding ICAT were cloned into the *Not* I site of a Cola1(II) (*Col2a1*)–based expression vector, PKN185 (8). The resulting vector contains the FLAG-tagged *ICAT* (*FLAG-ICAT*) complementary DNA including the 5' *Nde* I site *Col2a1* promoter (nucleotide 1940–2971, GenBank accession no. M65161), β -globin intron cassette, SV40 poly(A), and *Col2a1* enhancer (nucleotide 4930–5571, GenBank accession no. M65161). The *FLAG-ICAT* transgene was excised by *Nde* I and *Hind* III digestion. The *ICAT* transgene was then purified and injected into pronuclei of fertilized eggs from C57BL/6J mice. Positive transgenic founder mice were identified by Southern blot analysis, and mice were genotyped by polymerase chain reaction (PCR). The F₃ generation of 2 separate lines of *Col2a1-ICAT*–transgenic mice and their wild-type (WT) littermates were used for phenotype analysis and cellular function studies.

Histology and histomorphometry

Initial radiographic and histologic analyses were performed in 2 lines of the transgenic mice (lines 5 and 7), and similar articular cartilage destruction was found. Analysis of time-dependent articular cartilage destruction was then performed in 1 line of these transgenic mice (line 5). Knee joints from 6-, 9-, 12-, and 15-month-old WT and *Col2a1-ICAT*–transgenic mice were dissected, fixed in 10% formalin, decalcified, and embedded in paraffin. Serial midsagittal sections ($3-\mu$ m thick) of knee joints from 6-month-old mice were cut every 10 μ m from both the medial and lateral compartments. The sections were stained with Alcian blue/hematoxylin and eosin (H&E). Articular cartilage was outlined on the tibial surface, and an area algorithm in the software package ImagePro 4.5 (Leeds Precision Instruments, Minneapolis, MN) was used to determine the pixel area of outlined articular cartilage from each section. Using this approach, average articular cartilage area was determined. Cells in each section were counted, and the ImagePro counting algorithm was used to determine the average number of chondrocytes per unit cartilage area. Six mice per group were used for histomorphometric measurements.

Immunostaining

Tissue sections were deparaffinized by immersing in xylene, then fixed with 4% paraformaldehyde for 15 minutes and treated with 0.5% Triton for 15 minutes followed by fixing with 4% paraformaldehyde for another 5 minutes. Sections were then incubated with a mouse anti-FLAG M2 monoclonal antibody (1:200 dilution; Sigma, St. Louis, MO), rabbit anti–cleaved caspase 3 monoclonal antibody (Asp175, 1:200 dilution; Cell Signaling Technology, Danvers, MA), and rabbit anti–poly(ADP-ribose) polymerase (anti-PARP) polyclonal antibody (1:50 dilution; Abcam, Cambridge, MA) overnight. Secondary incubations were performed with a fluorescence-conjugated secondary antibody for 60 minutes, and sections were mounted with Vectashield (LabVision, Fremont, CA). Slides were visualized under a fluorescence microscope.

Cell isolation and cell culture

Three-day-old neonatal mice were euthanized and genotyped using tail tissues obtained at the time of death. The anterior rib cage and sternum were harvested en bloc, washed with phosphate buffered saline (PBS), and then digested with Pronase (Roche Applied Science, Indianapolis,

IN) dissolved in PBS (2 mg/ml) in a 37°C water bath with continuous shaking for 60 minutes. This was followed by incubation in a solution of collagenase D (3 mg/ml dissolved in serum-free Dulbecco's modified Eagle's medium [DMEM]; Roche Applied Science) for 90 minutes at 37°C. The soft tissue debris was thoroughly removed. The remaining sterna and costosternal junctions were further digested in a fresh collagenase D solution in petri dishes in a 37°C incubator for 5 hours with intermittent shaking. This step allows remnant fibroblasts to attach to the petri dish while the chondrocytes remain afloat in the medium. The digestion solution was filtered through Swinnex filter (Millipore, Bedford, MA) to remove all residual bone fragments. The solution was centrifuged, and the cells were resuspended in complete medium (DMEM with 10% fetal bovine serum [FBS], 1% penicillin/streptomycin, 100 mM L-glutamine, and 50 μ g/ml ascorbic acid, pH 7.1). The cells were counted and plated at the appropriate density. To remove any remaining fibroblasts, 24-hour cultures were treated with 0.05% trypsin for 1 minute to lift the fibroblasts from the culture dish while allowing the chondrocytes to remain attached.

For chicken chondrocytes, 6-week-old chicks were euthanized and the femora and tibiae were carefully separated by removing the surrounding tissue at the knee joint. Articular cartilage was cut off and digested with trypsin (1 mg/ml; Sigma) on a rotator for 30 minutes. This was followed by incubation in a solution of hyaluronidase (1 mg/ml; Sigma) for 60 minutes. The remaining tissues were further digested in collagenase A (1 mg/ml; Roche Applied Science) on the rotator overnight. The digestion solution was filtered through Swinnex filter to remove all residual bone fragments. The solution was centrifuged, and the cells were resuspended in complete medium (DMEM with 5% FBS, 1% penicillin/streptomycin, and 50 μ g/ml ascorbic acid, pH 7.1). Cells were counted and plated at the appropriate density.

Porcine articular chondrocytes were isolated from the articular cartilage of elbow and metatarsal joints of freshly slaughtered pigs. Briefly, articular cartilage was shaved from the articular surfaces and minced into 2×2 -mm chunks. Articular chondrocytes were then released from the native ECM with a 1.5-hour treatment in 0.05% (weight/volume) Pronase (Calbiochem, San Diego, CA) followed by 16-hour digestion in 0.2% (w/v) collagenase (Worthington, Lakewood, NJ). Cells were then pelleted and resuspended in 20 ml of cell culture media containing DMEM, 1% penicillin/streptomycin, 10% FBS, and 50 µg/ml ascorbic acid before counting and plating.

Luciferase assay

Primary sternal chondrocytes were transfected with the Top-flash reporter construct (0.5 μ g/ well, 12-well plate) and control SV40 Ranilla luciferase construct (0.01 μ g/well) using Lipofectamine 2000 reagent (Invitrogen, Carlsbad, CA) in the absence or presence of BIO (1 μ *M*), the glycogen synthase kinase 3 β (GSK-3 β) inhibitor. Cell extracts were harvested, and the luciferase activity was measured 2 days after transfection by the Promega Dual Luciferase system (Promega, Madison, WI).

Western blot analysis

Primary chondrocytes were lysed on ice for 30 minutes in a buffer containing 50 mM Tris HCl, pH 7.4, 150 mM NaCl, 1% Nonidet P40, and 0.1% sodium dodecyl sulfate (SDS) supplemented with protease inhibitors (10 μ g/ml leupeptin, 10 μ g/ml pepstatin A, and 10 μ g/ml aprotinin) and phosphatase inhibitors (1 mM NaF and 1 mM Na₃VO₄). Proteins were fractionated by SDS– polyacrylamide gel electrophoresis, transferred to a nitrocellulose membrane, and detected using anti–Bcl-2 and anti–Bcl-x_L monoclonal antibodies (1:1,000 dilution; Santa Cruz Biotechnology, Santa Cruz, CA) and an anti-Bax monoclonal antibody (1:1,000 dilution; Santa Cruz Biotechnology).

TUNEL staining

A TUNEL staining kit (DeadEnd Fluorometric TUNEL System; Promega) was used to assess programmed cell death by catalytically incorporating fluorescein-12-dUTP at 3'-OH DNA ends using the terminal deoxynucleotidyl transferase and recombinant enzyme. After deparaffinization, the tissue sections were placed in equilibrium buffer and then in a solution containing the equilibrium buffer, nucleotide mix, and terminal deoxynucleotidyl transferase and recombinant enzyme and incubated at 37°C for 1 hour. The reaction was stopped with $2\times$ saline–sodium citrate. Hoechst 33342 was used to stain nuclei. Results were visualized under a fluorescence microscope. Articular cartilage area from the tidemark to the surface and from the anterior to posterior cartilage margins was outlined in each slide. Chondrocyte numbers within the outlined regions were counted on 4 nonconsecutive sections from each joint sample. The apoptotic cell rates were determined by counting the numbers of TUNEL staining–positive cells in the cartilage area divided by the total cell number. Six *Col2a1-ICAT*–transgenic mice and 6 WT littermates were analyzed.

Caspase activity assay

Caspase 3/7 activity in primary mouse sternal chondrocytes and chicken and porcine articular chondrocytes was measured by adding Apo-ONE Caspase-3/7 Reagent (Apo-ONE homogeneous caspase 3/7 assay; Promega) to each well of a black 96-well tissue culture plate that was either blank or that had cells in culture. After incubation for 2 hours, the fluorescence of each well was measured. Caspase 9 activity was measured by adding Caspase-Glo 9 Reagent (Promega) to each well of a white 96-well tissue culture plate that was either blank or that had cells in culture. After incubation for 2 hours, the fluorescence of each well was measured. The caspase 8 activity was determined by adding the Caspase-Glo 8 Reagent (Promega) to the cell culture.

RESULTS

Generation of Col2a1-ICAT-transgenic mice and verification of signaling phenotype

Col2a1-ICAT-transgenic mice were engineered to overexpress ICAT in chondrocytes under control of the mouse *Col2a1* promoter. As mentioned, ICAT is a small peptide which selectively binds to the armadillo repeats of β -catenin and disrupts the interaction between β -catenin and TCF. Overexpression of ICAT specifically inhibits β -catenin signaling without disturbing cell adhesion (9). This unique feature of ICAT makes it an ideal molecule for the selective blockade of β -catenin signaling in target cells. In the present studies, we first examined the expression of the *ICAT* transgene in articular cartilage by immunostaining using an anti-FLAG antibody and found that the *FLAG-ICAT* transgene was specifically expressed in articular chondrocytes in 2-week-old and 6-month-old *Col2a1-ICAT*-transgenic mice (Figures 1a and b). To determine whether β -catenin reporter assays using primary sternal chondrocytes that were transfected with the Top-flash reporter construct and treated with the GSK-3 β inhibitor, BIO. We found that BIO-induced reporter activity was completely inhibited in chondrocytes derived from *Col2a1-ICAT*-transgenic mice (Figure 1c).

To further demonstrate the expression of the *ICAT* transgene in vitro, we examined FLAG-ICAT expression in isolated sternal chondrocytes via Western blotting using the anti-FLAG antibody. The results showed that the expression of ICAT protein could be detected in sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice but not in cells from the WT littermates (Figure 1d). The results from in vitro immunostaining assays further showed the expression of the *FLAG-ICAT* transgene in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (Figure 1e). To determine changes in β -catenin downstream genes, we examined the expression of TCFs and LEF and found that the messenger RNA expression of

Tcf3, *Tcf4*, and *Lef* was significantly reduced in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (Figure 1f). The results demonstrate that we have established a transgenic mouse model with overexpression of ICAT in chondrocytes and the associated inhibition of β -catenin signaling specifically in chondrocytic cells.

Articular cartilage destruction in Col2a1-ICAT-transgenic mice

At age 9 months, more severe destruction of the articular cartilage was observed (Figure 3a), and by age 12 months, the progressive destruction of the articular cartilage resulted in severe loss of the entire articular cartilage surface in *Col2a1-ICAT*–transgenic mice. Detached fragments of bone covered with cartilage were also observed (Figure 3b). Histologic analysis showed that 7 of 10 (70%), 6 of 9 (67%), and 6 of 11 (55%) *Col2a1-ICAT*–transgenic mice at ages 6, 9, and 12 months, respectively, had articular cartilage destruction. In contrast, no articular cartilage damage was found in 6-month-old WT mice, and only minor articular cartilage damage was found in <10% of age-matched 9- and 12-month-old WT mice (10 mice were analyzed).

At age 15 months, radiography showed ectopic bone formation in *Col2a1-ICAT*–transgenic mice (Figure 4a). Histologic analyses revealed surface fibrillation and vertical cleft development at the knee articular surface in *Col2a1-ICAT*–transgenic mice (Figure 4b). Complete loss of subchondral bone, which was replaced by the formation of new woven bone secondary to the articular destruction, was found in transgenic mice (Figure 4b). Overall, these phenotypic changes resembled the clinical features commonly observed in OA patients. The modified Chamber's scoring system reported by Glasson et al (10) was used to determine that the average histologic grade in *Col2a1-ICAT*–transgenic mice was 4 of 6 (i.e., moderate loss of ~40–60% of the articular cartilage).

Increase in chondrocyte apoptosis in Col2a1-ICAT-transgenic mice

We investigated the underlying mechanism of the pathogenesis of the articular cartilage destruction in *Col2a1-ICAT*--transgenic mice. No significant changes in cell proliferation were observed in articular chondrocytes of transgenic mice, as assessed by Ki-67 staining (data not shown). We then analyzed the expression of matrix genes via real-time reverse transcriptase-PCR using primary sternal chondrocytes isolated from neonatal WT and *Col2a1-ICAT*-- transgenic mice and found that the expression of type X collagen was reduced in primary chondrocytes from transgenic mice (data not shown), suggesting that the chondrocyte maturation process was delayed. Increased expression of matrix metalloproteinase 13 (MMP-13) is often associated with OA (11,12). However, in *Col2a1-ICAT*--transgenic mice, MMP-13 expression was also reduced (data not shown). These results suggest that articular

cartilage destruction observed in *Col2a1-ICAT*-transgenic mice may not be caused by the premature acceleration of articular chondrocyte differentiation.

We then analyzed cell apoptosis by TUNEL staining and found that the number of apoptotic cells in the articular cartilage was significantly increased in *Col2a1-ICAT*-transgenic mice (Figures 5a and b). Quantitative analysis showed a 75% increase in the number of apoptotic cells detected in the knee articular cartilage of the transgenic mice compared with that of the WT mice (Figure 5b). To further verify changes in cell apoptosis, we performed immunostaining using antibodies against the cleaved caspase 3 and PARP in the articular cartilage using knee joint sections derived from 6-month-old *Col2a1-ICAT*-transgenic mice and littermate controls. We found that staining of both cleaved caspase 3 and PARP was increased in *Col2a1-ICAT*-transgenic mice (Figures 5c and d). These results are consistent with the TUNEL staining results and indicate that apoptosis of articular chondrocytes is indeed increased in transgenic mice, suggesting that articular cartilage destruction in these mice may be caused by an increase in articular chondrocyte apoptosis.

It has been reported that the expression of Bcl-2 and Bcl- x_L in human articular chondrocytes protects cells from nitric oxide–induced apoptosis (13). In the present study, we found that the expression of Bcl-2 and Bcl- x_L proteins was decreased in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (Figure 6a). In contrast, the expression of Bcl-2/ cytochrome c in the cytokine-induced activation of apoptosis. Thus, we examined changes in caspase 9 activity and found that it was increased in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (Figure 6b). In contrast, no changes were found in caspase 8 activity (data not shown). Caspases 3 and 7 are downstream of caspase 9 and have been shown to be involved in chondrocyte apoptosis and development of OA (14,15). In the present studies, we found that caspase 3/7 activity was also significantly increased in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (Figure 6b).

To further determine the regulatory role of β -catenin signaling in caspase activities in articular chondrocytes, we also measured caspase 9 and caspase 3/7 activities in primary chicken and porcine articular chondrocytes following activation or inhibition of canonical Wnt signaling. Consistent with the results from *Col2a1-ICAT*–transgenic mice, we found that Wnt-3A (100 ng/ml) significantly inhibited caspase 9 activity in chicken articular chondrocytes, while Dkk-1 (1 µg/ml) stimulated caspase 3/7 activity (Figure 6c). In contrast, Wnt-3A (100 ng/ml) inhibited both caspase 9 and caspase 3/7 activities in porcine articular chondrocytes (Figure 6d). In addition, we found that Bax protein expression was reduced by treatment with Wnt-3A (100 ng/ml) in primary porcine articular chondrocytes (Figure 6e).

To determine the potential contribution of the ERK and phosphatidylinositol 3-kinase/Akt pathways to the apoptosis observed in *Col2a1-ICAT*–transgenic mice, we examined levels of phosphorylated ERK-1/2 and Akt in WT and transgenic mouse chondrocytes. No changes in either phosphorylated ERK-1/2 or Akt levels were observed in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice (data not shown), suggesting that the ERK and Akt pathways are not involved in *Col2a1-ICAT*–induced cell apoptosis. Overall, these results suggest that increased cell apoptosis is mediated by the Bcl-2/caspase 9/caspase 3/7 pathway.

DISCUSSION

In the present studies, we selectively blocked β -catenin signaling in chondrocytes by generating *Col2a1-ICAT*-transgenic mice. Specific expression of the *ICAT* transgene in articular chondrocytes of 6-month-old transgenic mice was observed, and severe articular cartilage

destruction was identified. We found that increased cell apoptosis in articular chondrocytes may contribute to the articular cartilage destruction in these transgenic mice.

Seven transgenic founder mice were initially generated. We could not establish transgenic lines for 2 of the founders since newborn mice were extremely small and did not survive past age 1 month. Ultimately, 2 transgenic lines (lines 5 and 7) with moderate phenotype were established and analyzed. Although severe articular cartilage destruction was observed, articular cartilage defects were not detected in 100% of the *Col2a1-ICAT*–transgenic mice, especially in older animals. Since ICAT may not bind to β -catenin in an irreversible manner, it is possible that β -catenin signaling may not have been completely blocked in some of the transgenic mice.

The phenotypic stability and survival of the chondrocytes are essential for the maintenance of a proper cartilage matrix. It has been reported that activation of Wnt/ β -catenin signaling inhibits cell apoptosis (16–18). In contrast, inhibition of Wnt proteins promotes programmed cell death in different types of cancer cells (19–23). The Wnt antagonist secreted Frizzled-related protein 1 (sFRP-1) promotes osteoblast and osteocyte apoptosis (24), and Wnt-7A inhibits apoptosis in articular chondrocytes (25). Serum withdrawal–induced apoptosis can be prevented by the canonical and the noncanonical Wnt proteins in osteoblasts (26). Furthermore, in human OA cartilage, FrzB-2 (encoding sFRP-4 protein) is highly expressed and is associated with chondrocyte apoptosis (27). These observations are consistent with our present finding that in *Col2a1-ICAT*–transgenic mice, cell apoptosis is significantly increased in articular chondrocytes. These findings indicate that Wnt/ β -catenin signaling proteins are involved in the antiapoptosis signaling pathway.

FRZB encodes sFRP-3, a glycoprotein that antagonizes the signaling of Wnt ligands through Frizzled membrane-bound receptors (28). In vitro transfection assays demonstrated that sFRP-3 can inhibit β -catenin nuclear translocation and TCF/LEF-dependent transcriptional activation (29). The mutation of sFRP-3 involving the Arg324Gly substitution causes sFRP-3 to lose its ability to antagonize Wnt signaling and may predispose humans to OA by altering the development or stability of cartilage in weight-bearing joints (29,30). These findings suggest that activation of Wnt/ β -catenin signaling may also be associated with the development of OA and further demonstrate that Wnt/ β -catenin signaling is critical for joint development, maintenance, and OA pathogenesis.

Beta-catenin plays an important role in articular chondrocyte function as well as in embryonic cartilage development (3). Although *Col2a1-ICAT*–transgenic mice show severe cartilage destruction, the possibility that phenotypic changes are due to defects in epiphyseal subchondral bone that occurred during early cartilage development cannot be ruled out in the present studies. Due to the limitation of the transgenic approach in this study, we will not be able to separate the effect of ICAT in articular chondrocytes from its effect on growth plate chondrocytes. To determine the specific role of β -catenin in articular chondrocytes in a cell-autonomous manner in adult mice, cell-specific deletion of the β -catenin gene in an inducible manner in articular chondrocytes in adult mice is required. Since tissue-specific knockout of the β -catenin gene (targeted by *Col2a1-Cre*) results in perinatal lethality (3), we have recently generated *Col2a1-CreER*–transgenic mice (31,32), and we are planning to generate mice with deletion of the β -catenin gene specifically in adult articular cartilage using this tamoxifen-inducible *Cre/loxP* system. These mouse models will provide more information about the interrelationship or exclusivity of the growth plate and articular cartilage effects when there are alterations in β -catenin signaling.

In summary, in this study we provide novel evidence that selective inhibition of β -catenin signaling in articular chondrocytes causes articular cartilage destruction in *Col2a1-ICAT*-transgenic mice. The findings not only suggest that increased cell apoptosis may contribute to

the articular cartilage damage in these mice, they also establish the idea that therapeutic approaches to OA involving inhibition of β -catenin signaling might not be helpful.

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Figure 1. Generation of *Col2a1-ICAT*-transgenic mice and verification of signaling phenotype

Two lines of *Col2a1-ICAT*–transgenic mice were established. The expression of the *FLAG-ICAT* transgene and the effect of inhibitor of β -catenin and T cell factor (ICAT) on β -catenin signaling in chondrocytes were examined. **a** and **b**, Immunofluorescence staining was performed using tissue sections from long bones of 2-week-old mice (**a**) and 6-month-old mice (**b**). Safranin O–fast green–stained histologic sections from 2-week-old mice and Alcian blue/ hematoxylin and eosin (H&E)–stained histologic sections from 6-month-old mice were used as controls. Using the anti-FLAG M2 antibody, the expression of the *FLAG-ICAT* transgene was detected in articular chondrocytes of *Col2a1-ICAT*–transgenic mice but not in those of their wild-type (WT) littermates. **c**, To determine changes in β -catenin signaling in *Col2a1-ICAT*–transgenic mice, primary sternal chondrocytes isolated from 3-day-old *Col2a1-ICAT*–transgenic mice and their WT littermates were transfected with Top-flash reporter construct and treated for 48 hours with glycogen synthase kinase 3 β inhibitor, BIO (1 μ M). Values are the mean and SEM. BIO stimulated Top-flash reporter activity in WT chondrocytes (* = *P* < 0.05 versus unstimulated WT chondrocytes, by unpaired *t*-test). In transgenic mouse chondrocytes, the basal reporter activity was reduced and the stimulatory effect of BIO was

completely inhibited. **d** and **e**, To further demonstrate *FLAG-ICAT* transgene expression in vitro, we performed Western blot (**d**) and immunostaining (**e**) assays using primary sternal chondrocytes derived from 3-day-old *Col2a1-ICAT*–transgenic mice and their WT littermates. Western blot and immunostaining data showed that FLAG-ICAT was detected only in chondrocytes derived from *Col2a1-ICAT*–transgenic mice and not in those from their WT littermates. **f**, The expression of β -catenin signaling downstream genes, *Tcf3*, *Tcf4*, and *Lef*, was examined by reverse transcriptase–polymerase chain reaction assays. The mRNA expression of *Tcf3*, *Tcf4*, and *Lef* was reduced in chondrocytes derived from *Col2a1-ICAT*–transgenic mice. (Original magnification × 10 in **a**; × 20 in **b** and **e**.) DAPI = 4', 6-diamidino-2-phenylindole.



Knee joints of 6-month-old mice



Figure 2. Articular cartilage destruction in 6-month-old *Col2a1-ICAT*-transgenic mice a, Changes in knee joint structure of *Col2a1-ICAT*-transgenic mice were analyzed by histology using Alcian blue/H&E and orange G staining. Mild cartilage degeneration was observed at the articular surface of knee joints in 7 of 10 6-month-old *Col2a1-ICAT*-transgenic mice. In the weight-bearing area, the articular layer was thinner or even completely missing in transgenic mice. Chondrocyte clusters (**arrow**) were found in *Col2a1-ICAT*-transgenic mice. No articular cartilage damage was found in 6-month-old WT mice. Boxed areas in upper panels are shown at higher magnification in lower panels. **b**-**d**, Histologic and histomorphometric analyses showed that articular area (**b** and **c**) and articular chondrocyte numbers (**d**) were significantly reduced in *Col2a1-ICAT*-transgenic mice. Values are the mean and SEM. * = P < 0.05 versus WT mice, by unpaired *t*-test (n = 6 mice per group). (Original magnification × 10.) ROI = region of interest (see Figure 1 for other definitions).

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b

Wild type



Knee joints of 9-month-old mice





Knee joints of 12-month-old mice

Figure 3. Progressive destruction of articular cartilage in 9- and 12-month-old Col2a1-ICAT-transgenic mice

a, Histologic analysis showed moderate destruction of articular cartilage tissue, and defects were extended to the calcified cartilage area in 6 of 9 9-month-old *Col2a1-ICAT*-transgenic mice (**arrow**). **b**, Severe destruction of articular cartilage and subchondral bone was detected in 6 of 11 12-month-old *Col2a1-ICAT*-transgenic mice. In contrast, only minor articular cartilage damage was found in <10% of 9- and 12-month-old wild-type mice (10 mice were analyzed). (Original magnification \times 10.)

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Figure 4. Ectopic and new bone formation in articular cartilage area of *Col2a1-ICAT*-transgenic mice

a, Radiography shows ectopic bone formation (**arrow**) in the knee joint of a 15-month-old Col2a1-ICAT-transgenic mouse. **b**, Histologic analysis reveals the complete loss of articular tissue at the weight-bearing area in a 15-month-old transgenic mouse. The destruction extends to the subchondral bone region. Boxed areas in upper panels are shown at higher magnification in lower panels. The high-magnification picture shows the deep cleft (**black arrows**) and new woven bone formation (**blue arrow**) in the knee joint area of a 15-month-old Col2a1-ICAT-transgenic mouse. (Original magnification \times 10.)



Figure 5. Increased cell apoptosis in articular cartilage of *Col2a1-ICAT*-transgenic mice

a, Cell apoptosis was determined by TUNEL staining. A significant increase in numbers of apoptotic cells was found in articular chondrocytes of 6-month-old *Col2a1-ICAT*-transgenic mice (**arrows**). DAPI staining was used as a positive control. The Alcian blue/H&E-stained histologic pictures from the same tissue blot are also shown. **b**, Quantification of apoptotic cells showed a 75% increase in the number of apoptotic cells in *Col2a1-ICAT*-transgenic mice. Values are the mean and SEM. * = P < 0.05 versus WT mice, by unpaired *t*-test (n = 3 mice per group). **c** and **d**, The immunostaining of cleaved caspase 3 (**c**) and poly(ADP-ribose) polymerase (PARP) (**d**) proteins was performed using knee joint sections derived from 6-month-old WT and *Col2a1-ICAT*-transgenic mice. The immunostaining of both cleaved caspase 3 and PARP was significantly increased in articular cartilage of *Col2a1-ICAT*-transgenic mice (*arrows*). (Original magnification × 20.) ROI = region of interest (see Figure 1 for other definitions).

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Figure 6. Changes in Bcl-2, Bcl- x_L , and Bax expression and caspase 9 and caspase 3/7 activities in primary sternal chondrocytes of *Col2a1-ICAT*-transgenic mice and in primary chicken and porcine articular chondrocytes

a, Primary sternal chondrocytes were isolated from 3-day-old *Col2a1-ICAT*–transgenic mice and their WT littermates. The expression of Bcl-2, Bcl-x_L, and Bax proteins was detected by Western blot analysis. The expression levels of Bcl-2 and Bcl-x_L proteins were decreased in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice, while the expression level of Bax protein was increased. **b**, Caspase 9 and caspase 3/7 activities were measured and found to be significantly increased in primary sternal chondrocytes derived from *Col2a1-ICAT*–transgenic mice. **c** and **d**, Caspase 9 and caspase 3/7 activities were also measured in primary chicken (**c**) and porcine (**d**) articular chondrocytes. Wnt-3A (100 ng/ml) significantly inhibited caspase 9 activity and Dkk-1 (1 µg/ml) stimulated caspase 3/7 activity in chicken articular chondrocytes after 48-hour treatment. In contrast, Wnt-3A (100 ng/ml, 48hour treatment) inhibited both caspase 9 and caspase 3/7 activities in porcine articular chondrocytes. Values in **b**–**d** are the mean and SEM. * = *P* < 0.05 versus WT mice or versus control (Cont), by unpaired *t*-test (n = 4 wells per group). **e**, Bax protein expression was reduced by 48-hour treatment with Wnt-3A (100 ng/ml) in primary porcine articular chondrocytes. See Figure 1 for other definitions.