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Distinct roles of Smad pathways and p38 pathways in cartilage-specific gene expression in synovial fibroblasts

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The role of TGF-β/bone morphogenetic protein signaling in the chondrogenic differentiation of human synovial fibroblasts (SFs) was examined with the adenovirus vector–mediated gene transduction system. Expression of constitutively active activin receptor–like kinase 3 (ALK3CA) induced chondrocyte-specific gene expression in SFs cultured in pellets or in SF pellets transplanted into nude mice, in which both the Smad and p38 pathways are essential. To analyze downstream cascades of ALK3 signaling, we utilized adenovirus vectors carrying either Smad1 to stimulate Smad pathways or constitutively active MKK6 (MKK6CA) to activate p38 pathways. Smad1 expression had a synergistic effect on ALK3CA, while activation of p38 MAP kinase pathways alone by transduction of MKK6CA accelerated terminal chondrocytic differentiation, leading to type X collagen expression and enhanced mineralization. Overexpression of Smad1 prevented MKK6CA-induced type X collagen expression and maintained type II collagen expression. In a mouse model of osteoarthritis, activated p38 expression as well as type X collagen staining was detected in osteochondrocytes and marginal synovial cells. These results suggest that SFs can be differentiated into chondrocytes via ALK3 activation and that stimulating Smad pathways and controlling p38 activation at the proper level can be a good therapeutic strategy for maintaining the healthy joint homeostasis and treating degenerative joint disorders.

Nonstandard abbreviations used: anterior cruciate ligament (ACL); bone morphogenetic protein (BMP); constitutively active activin receptor–like kinase 3 (ALK3CA); constitutively active MKK6 (MKK6CA); hemagglutinin (HA); matrix metalloproteinase (MMP); medial meniscus (MM); osteoarthritis (OA); receptor-regulated Smad (R-Smad); synovial fibroblast (SF); TGF-β–activating kinase 1 (TAK1).

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Introduction
Injury to the articular cartilage occurs under various pathological conditions such as trauma, inflammation, and aging (1), and cartilage injury is followed by osteoarthritic changes of the affected joints. Osteoarthritis is the most common degenerative joint disorder, affecting nearly half of the elderly population. Osteoarthritis is characterized by degradation of articular cartilage and overgrowth of cartilage and bone, known as osteophytes, at the periphery of the articular surface, which results in pain and loss of joint function (1, 2). Microscopically, loss of proteoglycan and fibrillation of the articular surface are observed at the early stage of arthritis. At later stages, clefts are formed, and at the end stage, erosive changes in the articular cartilage appear. The high prevalence of this disease results in high costs for treating patients, and therefore the development of good therapeutics for osteoarthritis is a matter of great urgency. Because of the limited capacity of spontaneous healing, the regeneration of intact articular cartilage is one of the most challenging issues in the orthopedic field (3, 4). Transplantation of autologous chondrocytes or mesenchymal progenitor cells and autogenous osteochondral transplantation (mosaicplasty) have been successfully utilized for the repair of focal osteochondral defects (3, 5–11). However, the application of these technologies is limited to small defects due to the difficulty of obtaining a sufficient amount of cells or tissues.

Synovium is a thin tissue lining the nonarticular surfaces of diarthrodial joints (12). Synovial tissues contain various types of cells, including type A cells, macrophage lineage cells, and type B cells, which are specialized synovial fibroblasts (SFs). It is now widely recognized that synovial tissues are involved primarily in the pathogenesis of arthritic joint disorders such as rheumatoid arthritis by producing the matrix-degenerating enzymes cysteine proteases and matrix metalloproteinases (MMPs) and the proinflammatory cytokines interleukin-1 (IL-1) and tumor necrosis factor-α (TNF-α) (12). We previously reported that SFs express a high level of receptor activator of NF-κB ligand, the osteoclast differentiation factor belonging to the TNF-α superfamily (13). In contrast to such catabolic actions, there is accumulating evidence that synovial cells have anabolic effects, leading to bone and cartilage production. Hunziker and Rosenberg reported that synovial cells can migrate into partial-thickness articular cartilage defects, where they proliferate and subsequently deposit a scar-like tissue (14). Nishimura et al. (15) demonstrated SFs show chondrogenic differentiation after being cultured in the presence of TGF-β, and de Bari et al. recently demonstrated that multipotent mesenchymal stem cells could be isolated from human synovial tissues, which differentiated into chondrocytes as well as osteoblasts, adipocytes, and myotubes under proper culture conditions (16, 17). Another dramatic clinical manifestation of the chondrogenic potential of synovial tissues is synovial chondromatosis, a tumor-like disorder characterized by the formation of multiple cartilaginous nodules, which is believed to be benign reactive metaplasia of synovial cells (18). These observations
lead us to speculate that synovial tissues contain multipotent cells with chondrogenic potential that might be involved in the repair process of articular cartilage defects and therefore might provide a good source for engineering articular cartilage.

There is accumulating evidence that TGF-β superfamily cytokines play an essential role in bone and cartilage development. Wozney and coworkers (19) reported that bone morphogenetic proteins (BMPs) induce early cartilage formation, and various studies have shown that TGF-β induces chondrocytic differentiation of undifferentiated mesenchymal cells (20–22). In the present study, we analyzed the role of TGF-β/BMP signaling on chondrogenic differentiation of human SFs by using the adenovirus vector–mediated gene transduction system. The introduction of an activated mutant of ALK3 (constitutively active activin receptor–like kinase 3 [ALK3CA]), also known as BMP type IA receptor, induced chondrocyte-specific marker expression in the cells. ALK3 signaling involves two different downstream cascades, the Smad pathway and the p38 MAP kinase pathway. We used a combination of adenoviral gene delivery and chemical inhibition to analyze the role of these two signaling cascades in inducing differentiation of SF cells toward a chondrocyte phenotype and found that both pathways are essential for chondrogenic differentiation. Interestingly, activation of p38 pathways alone induced markers of terminal chondrocyte differentiation, type X collagen expression and mineralization, which was suppressed by Smad1 coexpression. These results suggest that both the Smad and p38 pathways are necessary for chondrogenic differentiation of SFs and that the balance between these two pathways determines the stage of differentiation.

**Methods**

**Chemicals and antibodies.** Alpha-modified minimum essential medium (α-MEM) was purchased from Gibco BRL, Life Technologies Inc. (Rockville, Maryland, USA), and fetal bovine serum (FBS), from Sigma-Aldrich (St. Louis, Missouri, USA). Anti–p38 MAPK and anti–phospho-p38 MAPK (Thr180/Tyr182) were obtained from Cell Signaling Inc. (Cummings Center, Beverly, Massachusetts, USA). Anti-Flag was purchased from Sigma-Aldrich, and anti–hemagglutinin (anti-HA) was from Santa Cruz Biotechnology Inc. (Santa Cruz, California, USA). Anti–phospho-Smad1/5/8, which recognizes the phosphorylated form of Smad1, Smad 5, and Smad8, and anti–phospho-Smad2 were from Cell Signaling Inc. Anti–type II collagen was purchased from Oncogen (Boston, Massachusetts, USA) and anti–type X collagen was from LSL Co. (Cosmo Bio, Tokyo, Japan). Other chemicals and reagents used in this study were of analytical grade.

**Isolation of SFs from human synovial tissues.** Synovial cells were obtained as previously described (13, 23, 24). In brief, with enzymatic digestion, human synovial cells were isolated from synovial tissues of the knee joints of ten rheumatoid arthritis patients (37–75 years of age; mean, 60.3 years of age) at the time of total knee arthroplasty operations. Written informed consent for subsequent experiments was obtained from each patient. Cells were suspended in α-MEM containing 10% FBS and were cultured in monolayers. After three to five passages, subcultured cells were composed of morphologically uniform fibroblastic cells (SFs) that were free of macrophages. They were infected with adenovirus vectors and cultured in pellets (“pellet culture”). Primary chondrocytes were obtained from articular cartilage resected during the surgeries. Cartilage was minced finely in phosphate-buffered saline (PBS), and chondrocytes were isolated by sequential digestion at 37°C with 0.25% (weight/volume) trypsin for 30 minutes and with 2 mg/ml of clostridial collagenase in α-MEM containing 10% FBS and antibiotics (penicillin at 100 μg/ml and streptomycin at 100 μg/ml) overnight on an orbital shaker. Cells were isolated by centrifugation and were resuspended in α-MEM with 10% FBS. Cells were cultured in monolayers for 1 day and then subjected to RNA isolation.

**Constructs and gene transduction.** The recombinant adenovirus vectors carrying various molecules that modulate TGF-β superfamily signaling pathways, that is, HA-tagged constitutively active TGF-β/BMP type I receptors (ALK3CA, ALK5CA, and ALK66α), constitutively active MKK6 (MKK6CA), Flag-tagged Smad1 and Smad6 with CAG [cytomegalovirus IE enhancer + chicken β-actin promoter + rabbit β-globin poly(A) signal] promoter, were generated by the DNA–terminal protein complex method (25–27). SFs were infected with adenovirus vectors following a method previously described (13). In short, subconfluent SFs were incubated with a small amount of medium (α-MEM without serum) that contained the recombinant adenoviruses for 2 hours at 37°C at the indicated multiplicity of infection (MOI) and then with 10 times more medium to which 10% FBS had been added. Infected cells were cultured for additional 3 days for assessment of chondrogenic gene expression or were subjected to pellet culture 24 hours after the infection for histological examination.

**Pellet cultures of isolated SFs.** After 24 hours of viral infection, adherent cells were trypsinized and cells numbers were ascertained. Aliquots of 5 × 10⁶cells were spun down at 500 g in 15-ml polypropylene conical tubes in 5 ml of α-MEM with ascorbate-2-phosphate (0.1 mM) and 10% FBS. The cells were incubated at 37°C in 5% CO₂. Within 24 hours after incubation, the cells formed a single, free-floating pellet. The medium was changed every 2–3 days, and duplicate pellets were harvested after 3 and 7 days for real-time-PCR and Northern blotting and after 3 and 5 weeks for histological and immunohistochemical analysis. For visualization of the chondrogenic differentiation in vivo pellets were transplanted subcutaneously into nude BALB mice (nude mice) after 3 days of pellet culture. Mice were sacrificed 5 weeks after transplantation and the pellets were recovered and subjected to toluidine blue staining as well as immunostaining with anti–type II collagen.

**Immunoblotting.** All the extraction procedures were performed at 4°C or on ice. Cells were washed with PBS and then lysed by the addition of TNE buffer (1% NP-40, 10 mM Tris-HCl, pH 7.8, 150 mM NaCl, 1 mM EDTA, 2 mM NaVO₄, 10 mM NaF, and 10 μg/ml aprotinin). Lysates were prepared by centrifugation at 10,000 g for 20 minutes. An equal amount (15 μg) of proteins was separated by electrophoresis on 10% SDS-polyacrylamide gels. After electrophoresis, proteins were electronically transferred onto a nitrocellulose membrane. Immunoblotting with specific antibodies was performed with ECL Western blotting reagents (Amersham Co., Arlington Heights, Illinois, USA) according to the conditions recommended by the supplier.

**Histology and immunostaining.** Pellet cultures were fixed with 3.7% formaldehyde, embedded in paraffin, and cut into sections 4 μm in thickness. Representative sections were subjected to Alcian blue staining, Alizarin red staining, and immunohistochemistry. Alcian blue staining was performed according to the protocol described previously (28). Briefly, after deparaffinization, sections were stained with 0.5% Alcian blue 8GX (Wako, Osaka, Japan) in 0.1 N HCl for 1 hour. Mineralization was assessed by Alizarin red staining. In brief, sections were immersed in Alizarin red solution (40 mM, at pH 4.0) for 8 minutes at room temperature, and nonspecific staining was removed by several washes in distilled water. For immunostaining with anti–type
II collagen or anti–type X collagen, we utilized a CSA Kit (DAKO, Carpinteria, California, USA) following the manufacturer’s protocol.

**Total RNA extraction and real-time PCR.** Total RNA was isolated from SFs with ISOGEN (Wako) following the supplier’s protocol. Complementary DNA (cDNA) was synthesized from 1 μg of total RNA with the Superscript II reverse transcriptase kit (Invitrogen, Carlsbad, California, USA). For real-time PCR, the ABI Prism Sequence Detection System 7000 was used. Primers were designed based on sequences obtained from GenBank and amplicons of 50–250 base pairs with a melting temperature of between 55°C and 60°C were selected. Aliquots of first-strand cDNA (1 μg) were amplified with the Quantitect SYBER Green PCR Kit (Qiagen, Valencia, California, USA) under the following conditions: initial denaturation for 10 minutes at 94°C followed by 40 cycles consisting of 15 seconds at 94°C and 1 minute at 60°C. Data analysis consisted of fold induction, and the expression ratio was calculated from the differences in threshold cycles at which an increase in reporter fluorescence above a baseline signal could first be detected among three samples and was averaged for duplicate experiments. The primers we utilized in real-time PCR to detect sox9, type II collagen, osteocalcin, and GAPDH were as follows: sox9, 5′-AGAGGACACCAACCGGATTAC-3′ and 5′-AAGTGCTAGAGGGGTGTCT-3′; type II collagen, 5′-GGTTGGCTTCTTCATTCAGTA-3′ and 5′-TACCCGATGTTTTCTGTGCAG-3′; type X collagen, 5′-AGAGATGCGGTTGCTGCTTT-3′ and 5′-ACAGGCCTACCACAAAAGATGTA-3′; osteocalcin, 5′-GTGCAGAGTCCAGCAAAGGT-3′ and 5′-CGATAGGCCTCTCCTGAAGC-3′; osteopontin, 5′-ACAGGCGAGCTCATTGGAC-3′ and 5′-ACACTATCACCTCGGCCATC-3′; and GAPDH, 5′-GAAGGTGAAGGTCGGAGTCA-3′ and 5′-GAAGATGTGTGTGATGGGATTTC-3′.

**Northern blotting.** Equal amounts (15 μg) of RNA were denatured in formaldehyde, separated by 1% agarose gel electrophoresis and transferred to a nitrocellulose membrane (Hybond N+) (Amersham Pharmacia, Piscataway, New Jersey, USA), followed by ultraviolet cross-linking. ULTRAHyb hybridization solution (Ambion, Austin, Texas, USA) was used according to the manufacturer’s protocol. The blots were hybridized with a cDNA probe labeled with [32P]dCTP using Ready-To-Go DNA Labeling Beads (Amersham Pharmacia). Rabbit type II collagen and aggrecan probes were generously provided by Yoshiyasu Iwamoto (Thomas Jefferson University, Philadelphia, Pennsylvania, USA). Membranes were washed in 2× SSC for 15 minutes at 42°C and then in 0.1× SSC for 30 minutes at 65°C for visualization. X-ray film was exposed to membranes overnight at –80°C.

**Osteoarthritis model mice.** Osteoarthritic changes were developed in the knee joint by transection of the anterior cruciate ligament (ACL) and medial meniscus (MM) in C57BL/6 mice (mean age, 8 weeks) (29, 30). Briefly, after mice were anesthetized with ketamine and xylazine, a medial parapatellar skin incision was made. The subcutaneous tissues were incised and retracted, along with the articular capsule. The medial compartment of the knee joint was visualized and the ACL and MM were transected with a scalpel, and thereafter the capsule, medial retinaculum, and skin were sutured. Mice were housed in regular individual cages and allowed to exercise. Eight weeks after the surgery, the mice were sacrificed and paraffin-embedded sections of the affected joints were immunostained with anti–type X collagen and anti–phospho-p38 (Cell Signaling Technology Inc).

**Results**

**Adenovirus-mediated gene transduction modulates the Smad and p38 pathways in SFs.** We previously reported that adenovirus vectors can efficiently transduce foreign genes into synovial cells both in vitro and in vivo and that adenovirus infection itself does not affect the phenotypes of the cells (13). We constructed adenovirus vectors to analyze the role of ALK signaling as well as the Smad pathways and p38 pathways, which lie downstream of ALK signaling. SFs were infected with adenovirus vectors carrying various signaling molecules that modulate TGF-β superfamily signaling pathways, that is, HA-tagged constitutively active ALK3, ALK5, and ALK6 constitutively active MKK6, and Flag-tagged Smad1 and Smad6, as well as a control virus carrying the β-galactosidase gene (LacZ virus), and gene expression was determined by immunoblotting with specific antibodies. As shown in Figure 1, clear induction of the genes encoding ALK3CA, ALK5CA, and ALK6CA was observed by immunoblotting with anti–HA (Figure 1A), and Smad1 and 6, by anti-Flag (Figure 1B). ALK3CA or ALK6CA overexpression induced phosphorylation of Smad1, Smad5, and Smad8 in SFs, and ALK5CA-transduced cells showed Smad2 phosphorylation (Figure 1A). MKK6CA virus infection specifically activated p38 pathways in SFs, and the pathways were also activated in ALK3CA-transduced cells as determined by Western blotting with anti–phospho-p38 (Figure 1C). The increased p38 phosphorylation induced by either ALK3CA or MKK6CA overexpression was suppressed by the p38-selective inhibitor SB203580.

**Induction of chondrocyte-specific gene expression by ALK3CA transduction in pellet cultures of SFs.** To determine the effects of these transduced gene products on chondrocyte-specific gene expression in SFs, we subjected infected cells to pellet culture. After 7 days of culture, clear induction of type II collagen and aggrecan genes was observed in ALK3CA-transduced cultures by both Northern blot analysis (Figure 2A) and real-time PCR (Figure 2, B and C). Expression of these genes was also observed in ALK6CA-transduced cultures, albeit less efficiently, as shown in Figure 2, B and C, by real-time PCR. Contrary to the strong chondrogenic effects of ALK3CA virus, expression...
of osteocalcin or osteopontin was hardly detectable in the cells (Figure 2, C and D), indicating that hypertrophic and osteogenic differentiation were somehow blocked in these cultures. In contrast, neither type II collagen nor aggrecan gene expression was observed in ALK5CA virus–infected cells (Figure 2, A–C). Type II collagen and aggrecan expression induced by ALK3CA transduction was completely suppressed by coexpression with Smad6 or by SB203580 (Figure 2, B and C).

**ALK3 gene transduction increases Alcian blue–positive matrix and type II collagen deposition in pellet cultures of SFs.** For histological analysis, cells were subjected to pellet culture 24 hours after viral infection. After 3 weeks of pellet culture, cells were fixed and examined by Alcian blue staining (Figure 3, A, D, G, and I) and Alizarin red staining and type II collagen immunostaining (Figure 3, B, E, G, and J). ALK3CA virus–infected cultures showed cartilaginous matrix production that was strongly positive for Alcian blue staining (Figure 3D), while no positive staining was observed in LacZ virus–infected cultures (Figure 3A) or ALK5CA virus–infected cultures (Figure 3G), and only weak staining was observed in ALK6CA virus–infected cultures (Figure 3H). No Alizarin red staining was observed in ALK3CA–infected cultures (not shown), indicating that mineralization associated with osteogenic differentiation was not induced. ALK3CA virus–infected SFs showed an oval shape, morphologically reminiscent of chondrocytes (Figure 3D). Immunostaining with anti–type II collagen showed positive staining in ALK3CA virus–infected pellet cultures (Figure 3E) and weak staining in ALK6CA virus–infected cultures (Figure 3H), while we failed to detect type X collagen in ALK3CA virus–infected cultures (Figure 3F), which suggests an absence of terminal differentiation to hypertrophic chondrocytes. No positive type II collagen immunostaining was detected in LacZ virus–infected cultures (Figure 3B) or ALK5CA virus–infected cultures (Figure 3H).

**ALK3CA-transduced SFs after pellet culture form cartilage matrix in vivo.** To study chondrogenic differentiation of SFs in vivo, we subcutaneously transplanted the pellets into nude mice. Mice were sacrificed 3 weeks after the transplantation and the pellets were recovered and subjected to histological analysis. The transplanted SF pellets expressing ALK3CA were positively stained for toluidine blue (Figure 4C), which detects proteoglycan components, as does Alcian blue staining. Type II collagen immunostaining was also positive (Figure 4D), indicating the cartilaginous differentiation of the cultures in vivo, while Alizarin red staining was almost undetectable (data not shown). ALK6CA expression also induced chondrogenesis, albeit much less prominently (not shown), while neither LacZ (Figure 4, A and B) or ALK5CA (not shown) expression could induce chondrogenic phenotypes in the cultures. The histological observation was further confirmed by real-time PCR; expression of type II collagen and aggrecan was significantly higher in ALK3CA–transduced pellets (Figure 4, E and F). These results suggest that ALK3CA...
overexpression was able to target cartilage formation without subsequent bone formation in vivo.

Segregation of ALK signaling pathways. ALK signaling is known to be mediated by both the Smad pathways and MAP kinase pathways, especially the p38 pathways (31–33). We therefore attempted to distinguish the roles of the Smad pathways and p38 pathways from each other using a specific p38 inhibitor or adenovirus vectors. Smad6 coexpression or treatment of the cultures with the p38 inhibitor SB203580 completely abrogated the chondrogenic gene expression induced by ALK3CA (Figure 2, B and C). These results indicate that both the Smad pathways and the p38 MAP kinase pathways are required for the differentiation. Although Smad1 expression alone (MOI = 20) or a small amount of ALK3CA virus (MOI = 2) failed to induce type II collagen expression in SFs, both had synergistic effects, and robust upregulation of type II collagen gene was observed by coinfection of Smad1 virus (MOI = 20) and ALK3CA virus (MOI = 2) (Figure 5A). Interestingly, activation of p38 pathways alone by MKK6CA expression in SFs induced rapid induction of Sox9 and type II collagen, which declined rapidly, however, and type X collagen expression was subsequently increased (Figure 5B). Coexpression of Smad1 together with MKK6CA not only reduced type X collagen expression but also maintained type II collagen expression in the cells (Figure 5B). Pellet cultures infected

**Figure 3**
ALK3CA gene transduction increases Alcian blue–positive matrix and type II collagen deposition in pellet cultures of SFs. (A–J) Adenovirus–infect ed SF pellets were fixed with 3.7% formaldehyde after 3 weeks of culture and then were subjected to Alcian blue staining (A, D, G, and I) or immunostaining with anti–type II collagen (B, E, H, and J) or anti–type X collagen (Col X) (C and F). Distinct Alcian blue (D) and type II collagen (E) staining was observed in ALK3CA-expressing cultures. ALK6CA-expressing cultures showed weaker staining (I and J), and no positive staining was observed in ALK6CA–virus–infected (G and H) or LacZ virus–infected (A and B) cultures. No type X collagen immunostaining was observed in cultures expressing LacZ or ALK3CA (C and F). Scale bar: 100 µm.

**Figure 4**
ALK3CA-transduced SFs form cartilage matrix in vivo. (A–D) Three weeks after transplantation into nude mice, pellets were recovered and stained with toluidine blue (A and C) and immunostained with anti–type II collagen (B and D). Type II collagen immunohistochemistry was shown in the enlarged features of the rectangular area in the toluidine blue staining. Distinct positive staining was observed in ALK3CA-expressing cultures (B and D) in contrast to LacZ virus–infected cultures (A and C). Scale bars: 100 µm. (E and F) Real-time PCR analysis of type II collagen and aggrecan. Their expression was significantly higher in ALK3CA-expressing pellets than in LacZ-expressing pellets. *P < 0.001 (significantly different).
with MKK6CA virus were positively stained by type X collagen immunostaining as well as Alizarin red staining, which was suppressed by Smad1 virus coinfection (Figure 6).

**Type X collagen expression and p38 activation in synovial cells in osteoarthritic joints.** To examine the role of p38 activation in the development of degenerative changes in the articular cartilage, we next analyzed synovial tissues in the mouse model of osteoarthritis. After ACL and MM resection, the animals developed degenerative joint changes mimicking osteoarthritis. Osteochondrocytes were formed at the posterior edge of the femoral condyle and they were positively stained by anti-type X collagen as well as toluidine blue (rectangular areas in Figure 7, A and C) 4 weeks after the operation (corresponding to the stage of moderate osteoarthritis).

Clusters of migrating synovial cells were observed adjacent to the osteochondrocytes (Figure 7B, arrowheads), where future osteochondrocytes will develop, and they were weakly stained by toluidine blue and anti-type X collagen at the marginal area between synovium and osteophytes (rectangular areas in Figure 7, B and D). This region was also positively stained by anti–phospho-p38 (Figure 7F). No positive staining was observed in the normal synovium, however (data not shown).

**Discussion**

The signaling events leading to chondrogenesis still remain elusive, although there is accumulating evidence that TGF-β superfamily cytokines may play an important role (19–22). The receptors of TGF-β family members are composed of two different types of serine/threonine kinase receptors, known as type I and type II (31, 34, 35). Type II receptors are constitutively active kinases and phosphorylate type I receptors, also called ALKs. Type I receptors in turn mediate specific intracellular signaling pathways and therefore determine the specificity of the downstream signaling. So far, seven type I receptors have been identified, ALKs 1–7. ALK3 (BMPR-IA) and ALK6 (BMPR-IB) are structurally similar to each other and function as BMP receptors, while ALK5 and ALK4 work as type I TGF-β receptors. Using the adenovirus vector system, Fujii et al. reported that ALK1 CA, ALK2 CA, ALK3 CA, and ALK6 CA induced osteoblastic differentiation of C2C12 myoblasts and that ALK3 CA or ALK6 CA introduction induced chondrocytic differentiation of ATDC5 teratocarcinoma cells (27).

In the present study, we focused on the regulation of chondrogenic differentiation of primary SFs obtained from rheumatoid arthritis patients. SFs have chondrogenic potential (15, 16) and can migrate into articular cartilage defects, where they deposit a scar-like tissue as Hunziker et al. pointed out (14), suggesting that SFs have anabolic effects on joint homeostasis and are involved in the restoration process of articular cartilage. We demonstrated that adenovirus vector–mediated ALK3CA gene expression induced robust induction of chondrocyte-specific gene expression in SFs in a ligand-independent manner. Clear induction of Sox9, a key transcription factor regulating chondrogenesis (36, 37), followed by type II collagen and aggrecan expression, was observed in the ALK3CA-expressing cultures, while type X collagen was only weakly induced in the cultures and no osteocalcin expression could be found (Figures 2 and 5). Induction of these chondrocyte-specific genes through ALK3CA expression was not observed in skin fibroblasts, suggesting the cell specificity of the events (data not shown).

**Figure 5**

Segregation of downstream signaling pathways of ALK3. (A) Synergistic effect of Smad1 expression on the chondrogenic effects of the ALK3CA virus. Expression of Smad1 (MOI = 20) together with ALK3CA virus (MOI = 2) strongly induced expression of type II collagen in SFs. While bars indicate type II collagen expression on day 1 of cultures, and black bars indicate that on day 3. **P < 0.005 (significantly different). (B) MKK6-p38 pathways promote terminal chondrocytic differentiation of SFs. Mandatory activation of p38 pathways by expression of MKK6CA using adenovirus vectors rapidly activated expression of the Sox9 and type II collagen genes, which rapidly declined, while expression of a terminal chondrocytic differentiation marker, type X collagen, was gradually increased. Adenovirus vector–mediated overexpression of Smad1 together with MKK6CA suppressed type X collagen expression and maintained type II collagen expression in SFs. *P < 0.001; **P < 0.005 (significantly different).
shown). The chondrogenic effect of ALK3\textsuperscript{CA} virus was further confirmed histologically by pellet cultures performed in vitro and in vivo (Figures 3 and 4). Induction of neither the osteoblast markers osteopontin and osteocalcin nor the terminal chondrocyte differentiation markers type X collagen and mineralization was observed in ALK3\textsuperscript{CA}-expressing cells (Figures 2, 3, 5, and 6). These results suggest that ALK3 signaling, that is, BMP signaling, has both stimulatory and regulatory roles in chondrogenesis: to induce the chondrogenic differentiation of SFs and at the same time to block their osteoblastic or hypertrophic differentiation. Despite the structural similarity between ALK3 and ALK6, the ALK6\textsuperscript{CA} virus was much less efficient in chondrogenesis, the reason for which remains to be clarified. Although many studies have demonstrated a prochondrogenic effect for TGF-\(\beta\) (15, 16, 20–22), we failed to find an anabolic effect for ALK5\textsuperscript{CA} which is expected to mimic TGF-\(\beta\) signaling, on the chondrogenic differentiation of SFs. We cannot fully explain the discrepancy between our results and those of previous studies, but Robbins and coworkers recently reported that adenovirus vector–mediated TGF-\(\beta\) gene transduction into arthritic joints in fact exacerbated cartilage degradation (38), raising the possibility that sustained activation of TGF-\(\beta\) signaling, via ALK5, has instead a negative effect on chondrogenesis. Further study will be required to elucidate the difference between TGF-\(\beta\) and BMP signaling.

The signaling of TGF-\(\beta\)/BMPs is transduced by Smad family members (31, 34, 35). Receptor-regulated Smads (R-Smads) are direct substrates of type I receptors and are phosphorylated at the C-terminal SSV/MS motif. R-Smads then form heteromeric complexes with common-mediator Smads and translocate into the nuclei, where they regulate transcription of target genes. In addition to Smad pathways, there is evidence that MAP kinase cascades are also implicated in ALK signaling, in which TGF-\(\beta\)-activating kinase (TAK1), a member of the MAP kinase kinase family, plays a key role. TAK1 activates MAP kinase kinase in combination with an adaptor molecule, TAB1, which leads to JNK and p38 activation (32). The role of p38 in chondrogenesis has recently attracted particular interest because p38 inhibitors such as SB203580 suppress the chondrogenic differentiation of ATDC5 cells induced by growth/differentiation factor-5 (33, 39). However, the exact roles of the Smad pathways and p38 pathways in chondrocyte differentiation are not yet fully clarified. We used a combination of adenoviral gene delivery and a chemical inhibitor to segregate the roles of these two pathways downstream of ALK3 activation and found that (a) inhibitory Smad (Smad6) expression or treatment with the p38 inhibitor SB203580 suppressed the effect of ALK3\textsuperscript{CA} expression (Figure 2) (b) Smad1 synergistically augmented the effect of ALK3\textsuperscript{CA} (Figure 5A), and (c) activation of p38 pathways alone by MKK6\textsuperscript{CA} expression induced the hypertrophic differentiation markers type X collagen and mineralization in SFs, which was suppressed by Smad1 coexpression (Figures 5B and 6). These results suggest that although both Smad and p38 activation is necessary for chondrogenic differentiation of SFs, sustained activation of p38 pathways alone prompts the terminal differentiation of the cells. Consistent with our results, Zhen et al. (40) reported that parathyroid hormone inhibits type X collagen expression in hypertrophic chondrocytes by suppressing p38 pathways. Von der Mark et al. (41) reported the focal appearance of type X collagen in osteoarthritic cartilage, which may be involved in the degenerative changes of the articular cartilage and in the pathogenesis of osteoarthritis. Using the mouse model of osteoarthritis, we found that activated p38 is associated with type X collar
be differentiated into chondrocytes via ALK3 activation, and that activation of the Smad pathway while controlling the degree of p38 activation may be a way to generate committed chondrocytes for the repair and/or replacement of cartilage.

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Figure 7

Histological analysis of knee joints in the mouse ACL and MM resection model. (A–F) Toluidine blue staining (A and B) and type X collagen immunostaining (C and D) at the marginal area between the articular cartilage and synovium. B and D present higher-magnification views of A and C, respectively. Osteochondrocytes were formed at the posterior edge of the femoral condyle, and they were positively stained by anti–type X collagen as well as toluidine blue (rectangular areas in A and C). Clusters of migrating synovial cells were observed adjacent to the osteochondrocytes (B, arrowheads) where future osteochondrocytes will develop, and they were positively stained by anti–type X collagen at the marginal area between synovium and osteophytes (rectangular area in D). This region was also positively stained by anti–phospho-p38 (F). E and F represent phase-contrast microscopy (E) and immunostaining with anti–phospho-p38 (F) of the rectangular area in D. Positive phospho-p38 staining was observed at the area of osteochondrocytes as well as the marginal synovium. Scale bars: 500 µm (A–D) and 50 µm (E and F).


