

Chondrocyte responses to neurovascular peptides, cytokines, and a 3D environment: focus on ADAMs

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Abstract: Chondrocyte exposure to inflammatory stimuli in several arthritic conditions, including osteoarthritis, results in the well-characterized induction of extracellular matrix degrading proteinases, notably members of a disintegrin and metalloproteinase (ADAM) with thrombospondin domains and matrix metalloproteinase families. Here we briefly review the less-studied ADAM family of proteinases in chondrocyte and cartilage biology. Following damage, cartilage is exposed to neurovascular peptides, and in this study we hypothesized that substance P and bradykinin, alongside inflammatory cytokines, may modulate chondrocyte steady-state messenger RNA levels for the proteolytic ADAM family members as well as for key cytokines and neuropeptides. We compared chondrocytes cultured in both two-dimensional and three-dimensional (3D) environments and found that 3D culture generally resulted in repression of expression of the genes under investigation, with the exception of anti-inflammatory interleukin 10 which was markedly upregulated in a 3D environment. Substance P and bradykinin had little effect on ADAM family expression, but further investigation revealed that a combination of bradykinin and cytokines led to enhanced expression of ADAM28 and a synergistic upregulation of interleukin 6, also observed under hypoxic conditions. Overall these data reveal wider chondrocyte responses to neurovascular peptides which may have an impact in an osteoarthritis context.

Keywords: chondrocyte, ADAM, metalloproteinase, cytokine; bradykinin

Introduction

The cartilage environment

Chondrocytes are surrounded by an extensive extracellular matrix comprising type II collagen and aggrecan as well as a large number of other collagens, proteoglycans, and proteins together providing a remarkable, unique environment. Chondrocytes lack the capacity to drive efficient repair and thus in osteoarthritis (OA) articular cartilage is irreversibly damaged. While earlier studies viewed OA as a wear-and-tear condition, there is now a consensus, reviewed by Konitten et al,¹ that chondrocytes both respond to and may themselves produce proinflammatory cytokines which can result in cartilage breakdown through modulating the expression of metalloproteinases including a disintegrin and metalloproteinase with thrombospondin domains (ADAMTS) enzymes, matrix metalloproteinases (MMPs), and serine proteinases.²

Interplay of pain-related neurovascular peptides with cartilage biology

OA is increasingly viewed as a disease of the whole joint, including cartilage, synovium, bone, and, in knee OA, the infrapatellar fat pad.^{3,4} Adult cartilage is avascular and the

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resulting hypoxic environment promotes chondrocyte differentiation,⁵⁻⁷ in addition to maintaining the chondrocyte phenotype.⁸ Cartilage is also aneural, but damaged cartilage is recognized as a potentially permissive environment for neurovascular infiltration.⁹⁻¹¹ Sensory nerve fibers thus accompany new blood microvessels that grow into damaged cartilage in OA.

While effects of several cytokines and chemokines on neuronal excitation have been explored,¹² as yet the effects of neurovascular factors on chondrocyte gene expression and function remain less studied. Several neuropeptides, which play key roles in transmission of pain signals from the periphery to the central nervous system, are upregulated in joint disease. For example, substance P is elevated in the synovial fluid in both OA and rheumatoid arthritis (RA) joints, and the expression of substance P is particularly associated with more active OA.¹³ Im et al¹³ also demonstrated that substance P and its receptor NK1-R are expressed at higher levels in human articular cartilage from OA patients than controls. Substance P is produced in nerve fibers of the intracellular fat pad where a role in inflammation and cartilage destruction has been suggested.⁴

Kinins, including bradykinin, are often associated with acute and chronic inflammation, and, more recently, attention has turned to potential roles in OA. Bradykinin is a vasodilator with established roles in inflammation and a number of studies show that it is elevated in OA synovial fluid.¹⁴ An antagonist of bradykinin has been shown to reduce pain levels in a small study of patients with very painful OA.¹⁵ Functional bradykinin receptors are expressed in human chondrocytes with evidence of release of inflammatory cytokines interleukin (IL)6 and IL8 following bradykinin stimulation.¹⁶ The synovial compartment, which is itself innervated, may also respond to neuropeptide release and bradykinin is important in synovitis in OA.¹⁷ A more recent study showed that an antagonist of bradykinin receptor B (BDKRB)1 provides protection against joint injury in the rat anterior cruciate ligament model.¹⁸ In addition, a polymorphism in BDKRB2 is associated with both higher risk and higher severity of OA in one study.¹⁹

As well as roles in regulation of gene expression, neuropeptides may also modify aspects of chondrocyte biology including cell proliferation and migration. Chondrocytes can migrate *in vitro* and *in vivo* in development and possibly in OA.²⁰ Opolka et al²¹ demonstrated that substance P can induce proliferation of murine newborn costal chondrocytes and elevate cell adhesion and focal contact formation, indicating novel roles in cell-matrix interactions.

ADAMs in chondrocytes and cartilage

A disintegrin and metalloproteinase (ADAM) family members with proteolytic activity may play roles in chondrocyte biology and OA. While regulation of both MMPs and ADAMTS proteinases has been studied extensively in chondrocytes, less attention has been given to the ADAM family. Several ADAMTS proteinases can degrade proteoglycans²² and ADAMTS4 and 5 are known to play key roles in proteoglycan degradation in cartilage with human ADAMTS5 showing 1,000-fold greater aggrecan degradative capacity than ADAMTS4 in biochemical analyses.²³ Both enzymes can degrade aggrecan in human cartilage explants²² and recently developed antibodies specifically block ADAMTS5 aggrecan fragment release,²⁴ in one case to a far greater extent than possible through blockade of ADAMTS4.²⁵ In addition *adamts5* (but not *adamts4*) null mice are protected from experimentally induced OA.^{26,27}

ADAMs have a structure comprising a propeptide, catalytic, disintegrin, cysteine-rich, transmembrane, and cytoplasmic domains and are furin-activated (furin cleavage site located between pro- and catalytic domains).²⁸ As transmembrane enzymes, ADAMs are well placed for their extensive shedding functions, where cell adhesion molecules, growth factors, and their receptors among many other proteins are cleaved and thus shed from the cell surface.²⁹ Certain MMPs share shedding functions (including, for example, MMP9), but this seems to be a function mainly restricted to ADAMs. Through their disintegrin domains, several ADAMs can bind to integrins, a function also observed with certain MMPs (even though in MMPs disintegrin domains are absent),³⁰ but ADAMTS enzymes, despite the presence of a disintegrin domain, seem to lack this function.²⁸ Several proteolytic ADAMs are expressed in chondrocytes or OA cartilage.^{31,32} Here, we briefly review findings regarding ADAMs with proteolytic activity where expression and/or function relating to chondrocyte biology and/or OA have been reported.

ADAM8

Expression of ADAM8 has been observed in cartilage during mouse development,³³ and expression of an inactive ADAM8 mutant results in protection against murine experimental inflammatory arthritis.³⁴ In addition, ADAM8 is expressed in OA cartilage, and when isolated from chondrocyte culture medium, this protease is able to cleave fibronectin.³⁵ This is an important finding, since fibronectin fragments are found in damaged joint tissues and are able to induce MMP gene expression and aggrecan degradation in cartilage.³⁶

ADAM9

Flannery et al³⁷ showed the downregulation of ADAM9 expression in IL1-treated porcine chondrocytes in three-dimensional (3D) compared to two-dimensional (2D) culture,³⁷ and this proteinase is expressed during human mesenchymal cell differentiation along the chondrocyte lineage.³⁸ ADAM9 expression is upregulated in end-stage OA cartilage.²³

ADAM10

Earlier studies revealed increased ADAM10 mRNA levels in IL1-treated bovine chondrocytes,³⁷ and enhanced ADAM10 protein levels were observed in OA compared to healthy cartilage and in IL1-stimulated bovine nasal cartilage.³⁹ More recent observations indicate that ADAM10 cleavage of the cell adhesion molecule, N-cadherin, has an important role in chondroprogenitor cell–cell interactions leading to condensation and differentiation.⁴⁰

ADAM12

This ADAM has been associated with OA in genetic studies,^{41,42} and ADAM12 mRNA and protein (by immunocytochemistry) levels are upregulated in end-stage OA cartilage.^{22,23} ADAM12 plays a role in chondrocyte proliferation through cleavage of insulin-like growth factor binding protein (IGFBP)-5,²² which may be important in attempted repair by chondrocytes in OA.

ADAM15

IL1 was shown to enhance the expression of ADAM15 in monolayer cultures of porcine chondrocytes³⁷ and ADAM15 (also known as metargidin) is upregulated in end-stage OA cartilage.⁴³ Interestingly, ADAM15 null mice developed a more accelerated OA,⁴⁴ suggesting that ADAM15 is protective against the development of OA. Later studies revealed that ADAM15 is anti-apoptotic in OA chondrocytes and promotes cell adhesion to type II collagen when overexpressed in chondrocytes.⁴⁵ The anti-apoptotic mechanism involves ADAM15 cytoplasmic tail interaction with focal adhesion kinase (FAK) resulting in enhanced FAK phosphorylation and promotion of cell survival under genotoxic conditions.⁴⁶

ADAM17

Due to its role in shedding of tumor necrosis factor alpha (TNF α), along with cleavage of other relevant molecules, ADAM17 has been highly associated with RA.⁴⁷ Soluble IL6-receptor (sIL6-R) is often generated through ADAM17 cell surface cleavage, enabling sIL6-R to increase IL6-mediated cartilage damage. A very recent study has shown that

ADAM17-mediated shedding of IL6-R is blocked by secreted frizzled-related protein 3 (sFRP3) and the rare OA double-mutant variant of sFRP3 loses this inhibitory capacity.⁴⁸ This could result in greater release of sIL6-R and hence enhanced IL6 signaling leading to further cartilage destruction.

ADAM19

Expression of this ADAM is elevated during in vitro differentiation of human chondrocytes.³⁸ ADAM19 can shed TNF α from the cell surface,⁴⁹ though this has not as yet been demonstrated in chondrocytes.

ADAM28

ADAM28 is not observed in undamaged human cartilage; however, it is expressed in end-stage OA cartilage.²³ More recently, a proteomic approach found ADAM28 in the pericellular matrix of chondrocytes in articular cartilage.⁵⁰ Since ADAM28 can degrade IGFBP-3,⁵¹ it is possible that this proteinase plays a role in chondrocyte proliferation, as has been demonstrated for breast cancer cells.⁵² Another function of ADAM28 appears to be in the shedding of TNF α from the cell surface, demonstrated in monocytes and which may be of relevance in arthritic conditions.⁵³ ADAM28 can be regulated by 9-cis retinoic acid and peroxisome proliferator-activated receptor gamma ligands in differentiated monocytes,⁵⁴ as well as by 9-cis retinoic acid in chondrocytes,⁵⁵ indicating the involvement of potentially similar pathways across cell types. ADAM28 also degrades human chondrocyte proteoglycans,⁵⁵ such that co-culture of bovine nasal cartilage discs with ADAM28-expressing COS-7 cells results in greater proteoglycan release.⁵⁵

Neurovascular peptides and metalloproteinase expression

Studies on neurovascular peptides substance P and bradykinin in an arthritic context have to date focused on MMP or ADAMTS expression. Miller et al¹² showed that treatment of cartilage fragments with substance P in combination with fibroblast growth factor 2 results in cartilage degradation. In addition, exposure of human chondrocytes to increasing concentrations of substance P leads to enhanced expression of several MMPs.¹³ Taken together, these data suggest that substance P may be an important regulator of catabolism of articular cartilage in OA. Bradykinin levels correlate with glycosaminoglycan fragments in synovial fluid of OA patients,⁵⁶ which could suggest a link to matrix degradation. Bradykinin is also known for its ability to induce the activation of the ADAM17-EGFR transactivation pathway⁵⁷

and to indirectly enhance MMP production in a number of other cell types, but this has not been explored in chondrocytes.

The potential interplay between neurovascular peptides released in the OA joint and ADAM proteinase expression remains little explored. We hypothesized that a 3D culture environment may regulate chondrocyte responses to cytokines and neurovascular peptides. Specifically in the experiments reported here, we have surveyed the expression of genes encoding proteolytic ADAM family members, tissue inhibitors of metalloproteinases (TIMPs) and selected cytokines/neuropeptides and receptors in chondrocytes in response to the cytokines IL1 α and oncostatin M (OSM), substance P, and bradykinin in both 2D and 3D culture environments.

Materials and methods

Materials

Unless otherwise stated chemical reagents were purchased from Sigma Aldrich, Gillingham, UK, culture media from Gibco Thermo Fisher Scientific, UK, and tissue culture plastics from Nunc Thermo Scientific, UK.

Cell culture model

C28I/2 chondrocytes were cultured essentially as previously described⁵⁸ in DMEM with 10% fetal calf serum (FCS) at a cell density of 190,000 cells/24 well in 2D monolayer culture or in a 3D model with culture of chondrocytes over poly-HEMA,⁵⁸ for a total of 96 hours. During the final 24 hours, the cells were stimulated with a combination of IL1 α (10 ng/mL) and OSM (5 ng/mL), bradykinin (1 μ M), or substance P (100 μ M) in DMEM with 1% FCS. Where indicated cells were also exposed to a hypoxic environment (1% oxygen in a Billins chamber) for 24 hours prior to and during treatments.

RNA extraction, TaqMan low-density array, and qRT-PCR analysis

Following cell lysis, RNA was extracted using the RNeasy plus mini kit (Qiagen) according to the manufacturer's instructions. Total RNA (1 μ g) was reverse transcribed with 1 μ g random primer oligonucleotides (Invitrogen, Paisley, UK) and 200 units of M-MLV reverse transcriptase (Promega, Southampton, UK). cDNA samples were analyzed on custom-designed TaqMan low-density array (TLDA) plates (Applied Biosystems, Warrington, UK) to detect the expression of the genes encoding ADAMs 8, 9, 10, 12, 15, 17, 19, 20, 21, 22, 28, 33, TIMPs 1–4, the cytokines IL6 and IL10, the neurotrophins nerve growth factor (NGF) and brain-derived neurotrophic factor (BDNF), neuropeptide Y (NPY), and bradykinin receptors and the house-keeping gene 18S. (ADAM22 was included in this initial TLDA screen as, while it lacks proteolytic activity, ADAM22 expres-

sion is highly upregulated in OA cartilage compared to healthy cartilage, at least at the mRNA level²³). Prior to TLDA analysis, samples were analyzed on geNorm (Primer Design; Qiagen, Manchester, UK) where 18S ribosomal RNA was found to be the least-regulated housekeeping gene (of the 12 explored) for all treatment regimens (data not shown). TLDA analysis was performed in triplicate for each condition. A total of 150 ng of RNA was loaded into each port of the TLDA plates and analyzed in an ABI 7900HT Instrument (Applied Biosystems).

QRT-PCR using an ABI prism 7500 sequence detection system (Applied Biosystems) was performed as previously described⁵⁹ with primer probe sets from Applied Biosystems. Genes were analyzed using predesigned Applied Biosystems inventorized primer probe sets (Table 1) or primer probe sets designed using Primer Express 1.0 Software (Applied Biosystems; Table 1). Primers were designed spanning exon–intron boundaries to avoid amplification of genomic DNA and were tested for specificity by BLAST analysis. 18S ribosomal RNA was used as an endogenous control to normalize for differences

Table 1 Inventorized primer set codes for TLDA analysis and for further experimentation (ADAM28)

| Gene | ABI inventorized primer set |
|--------------|---|
| ADAM9 | Hs00177638_m1 |
| ADAM10 | Hs00153853_m1 |
| ADAM12 | Hs01106101_m1 |
| ADAM15 | Hs00984794_m1 |
| ADAM17 | Hs01041915_m1 |
| ADAM19 | Hs00224960_m1 |
| ADAM20 | Hs01083178_s1 |
| ADAM21 | Hs01652548_s1 |
| ADAM22 | Hs00244640_m1 |
| ADAM28 | Hs00248020_m1 |
| ADAM33 | Hs00905552_m1 |
| TIMP1 | Hs00171558_m1 |
| TIMP2 | Hs00234278_m1 |
| TIMP3 | Hs00165949_m1 |
| TIMP4 | Hs00162784_m1 |
| IL6 | Hs00985639_m1 |
| IL10 | Hs00961622_m1 |
| BDNF | Hs00380947_m1 |
| NGF | Hs00171458_m1 |
| NPY | Hs00173470_m1 |
| Tachykinin 3 | Hs00203109_m1 |
| BDKRB1 | Hs00664201_s1 |
| BDKRB2 | Hs00176121_m1 |
| ADAM28 | Forward: 5'-GGGCCACGATTGCA-3' Reverse: 5'-TGAACCTTCCTGTCTTTCAATTTACT-3' Probe: 5'FAM- AGAACATTGCCCTACCTGCCACCAAAC- TAMRA-3 |

Abbreviations: ADAM, a disintegrin and metalloproteinase; TLDA, TaqMan low-density array; TIMP, tissue inhibitors of metalloproteinase; IL, interleukin; BDNF, brain-derived neurotrophic factor; NGF, nerve growth factor; NPY, neuropeptide Y; BDKRB, bradykinin receptor B.

in the amount of total RNA in each sample (18S rRNA primers and probe were purchased from Applied Bioscience).

Results

Effects of culture in 3D and of cytokine stimulation

Of the 12 ADAMs whose steady-state mRNA expression was determined, eleven showed modulation of expression in chondrocytes by TLDA analysis (Figure 1; Table 2). 3D culture resulted in the downregulation of steady-state mRNA levels of seven of the ADAMs expressed in chondrocytes, with ADAM9, ADAM20, and ADAM33 mRNAs showing at least a twofold suppression (Figure 1; Table 2). TIMP2 mRNA was also significantly repressed, but other TIMPs were unaffected by 3D culture. Cytokine stimulation resulted in the upregulation of expression of several ADAM mRNAs in 2D, but in 3D culture, this was restricted to ADAMs12 and 28 (Table 2). Interestingly, ADAM10 and ADAM15 mRNAs were no longer modulated by cytokines when cells were cultured in 3D and that of ADAM19 and ADAM21 was now repressed (Table 2). TIMP1 mRNA was increased by cytokines in both 2D and 3D cultures, whereas that of TIMP3 was repressed by cytokines in both culture environments (Table 2).

We surveyed a small number of receptors for neuropeptides as well as neurotrophins and cytokines. It is interesting to note that cytokines upregulated the expression of bradykinin receptors 1 and 2 both in 2D and in 3D cultures of chondrocytes. In addition IL6 and IL10 mRNAs were strongly upregulated by cytokines in both 2D and 3D (Table 2). In contrast, cytokines repressed the expression of BDNF, NPY, and tachykinin 3 in both culture environments.

Effects of substance P and bradykinin

Substance P treatment slightly upregulated ADAM17 mRNA (1.4-fold) but bradykinin did not alter ADAM expression in 2D culture. In chondrocytes in 3D culture, substance P and bradykinin both significantly repressed the gene expression of several ADAMs (Figure 1; Table 2). Substance P repressed TIMPs 2 and 3 whereas bradykinin repressed TIMPs 2 and 4, respectively (Figure 1; Table 2). Steady-state mRNA levels of few other genes were modulated by either substance P or bradykinin (Figure 1; Table 2).

Effects of co-stimulation with neurovascular peptides and cytokines under normoxic and hypoxic conditions

In general, little overlap of induction of gene expression was observed under different stimuli. The small number of

genes regulated by substance P or bradykinin alone led us to explore the effects of co-stimulation with cytokines and these neurovascular peptides. We were particularly interested to explore further the regulation of ADAM28 since this proteinase has recently been found in the pericellular matrix of chondrocytes.⁵⁰ We pursued its expression alongside that of IL6, since this cytokine was upregulated by several stimuli. We exposed cells in 3D culture to cytokines or neuropeptides both alone and in combination with cytokines under normoxic and hypoxic conditions.

In 3D culture under normoxic conditions, we confirmed the upregulation of expression of ADAM28 in the presence of cytokines. Interestingly expression was further enhanced by co-stimulation with bradykinin (Figure 2A). A further TLDA survey of the other proteolytic ADAMs did not reveal any effect of bradykinin on their expression in the presence of cytokines (results not shown). Preliminary investigations revealed that exposure of chondrocytes to hypoxic conditions significantly upregulated ADAM28 mRNA by 3.7-fold and in the presence of cytokines alone no further regulation was observed. However, under hypoxic conditions co-stimulation with bradykinin led to further upregulation of ADAM28 mRNA (Figure 2A). Cytokine induction of IL6 mRNA was confirmed in 3D culture under normoxic conditions and an apparently synergistic induction with bradykinin co-stimulation was observed (Figure 2B). In contrast to ADAM28, the cytokine-induced expression of IL6 mRNA was suppressed under hypoxic conditions (2.4-fold) but synergistic induction with co-stimulation with bradykinin was again observed (Figure 2B). Expression of TNF α was also explored but no synergy with bradykinin was observed (data not shown).

Discussion

Neurovascular mediators including substance P and bradykinin are released into the synovial fluid in both OA and RA and are involved in pain perception. Here we show that the cellular environment has profound effects on constitutive ADAM expression as well as on the modulating effects of soluble mediators, either cytokines or neurovascular peptides.

3D suspension culture results in downregulation of mRNA levels for many genes studied

Overall, we observed that the culture of chondrocytes in 3D significantly repressed steady-state mRNA levels of seven ADAMs including ADAM9, as well as IL6, TIMP2, BDNF, and NGF. Downregulation of ADAM9 mRNA in chondrocytes has been reported previously in 3D cultures.³⁷ Collagen I downregulation

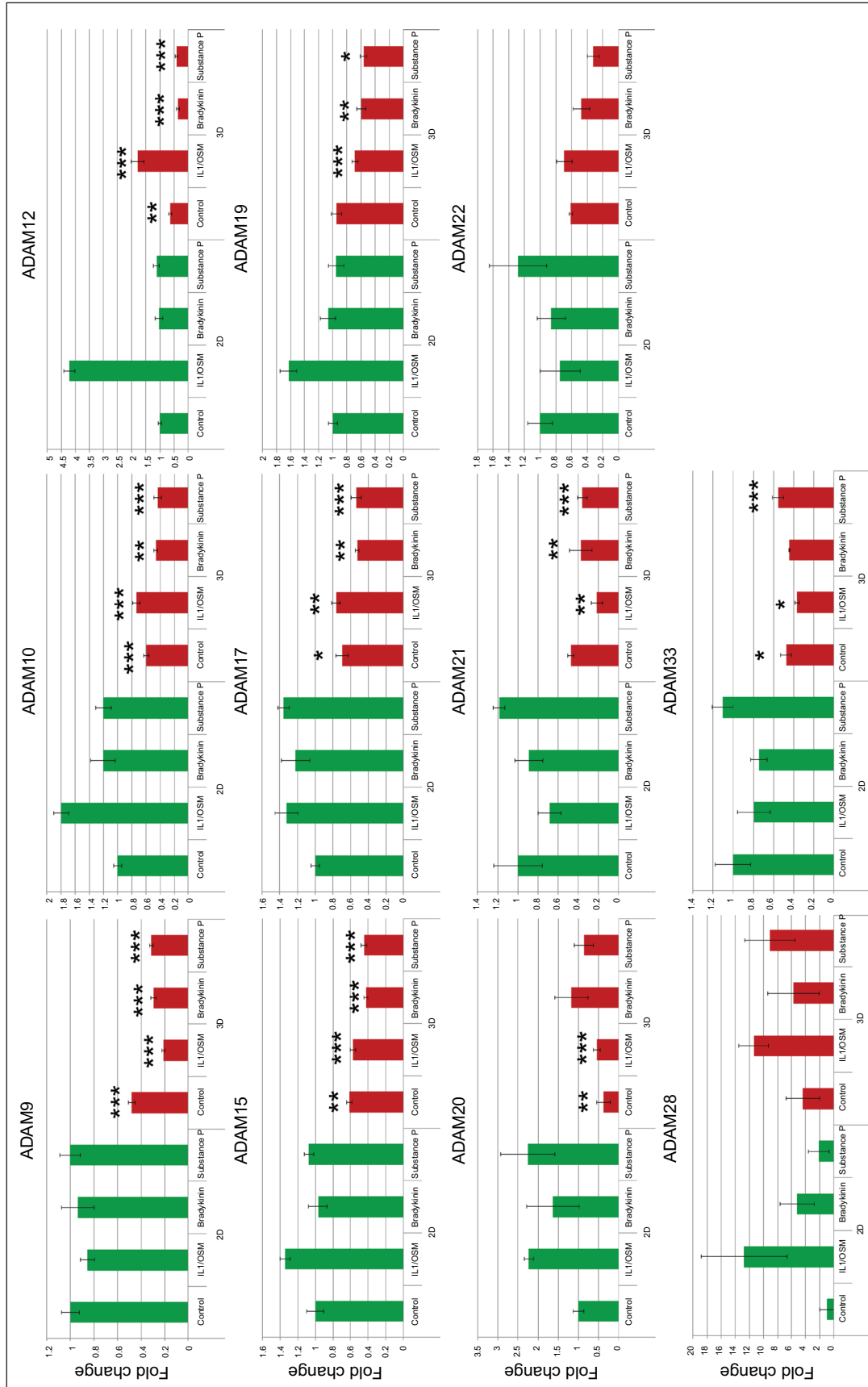


Figure 1 Culture in 3D in general suppresses ADAM steady-state mRNA levels, irrespective of the stimulus added.
Notes: C28I2 chondrocytes were cultured either in 2D (green) or 3D (red) with the indicated stimuli, and following RNA extraction and reverse transcription, TLDA analysis was performed. Each bar represents the mean \pm SEM for triplicate samples following normalization to 18S RNA. For this fold-change analysis, control (unstimulated) cells were set as 1. Comparisons between the levels in 2D and 3D for control conditions and for each stimulus are indicated. * $p < 0.05$; ** $p < 0.02$; *** $p < 0.01$.
Abbreviations: 2D, two dimensional; 3D, three dimensional; TLDA, TaqMan low-density array; SEM, standard error of mean; ADAM, a disintegrin and metalloproteinase; IL, interleukin; OSM, oncostatin M.

Table 2 Modulation of steady-state mRNA levels for ADAMs, TIMPs, cytokines, and neurovascular factors under the indicated conditions, determined by TLDA analysis

| Gene | 3D versus 2D | IL1/OSM (2D) | IL1/OSM (3D) | SubP (2D) | SubP (3D) | BK (2D) | BK (3D) |
|--------------|--------------|--------------|--------------|-----------|-----------|---------|---------|
| ADAM9 | -2.1 | ↔ | -2.5 | ↔ | -1.5 | ↔ | -1.7 |
| ADAM10 | -1.7 | 2 | ↔ | ↔ | ↔ | ↔ | ↔ |
| ADAM12 | -1.7 | 4 | 2 | ↔ | -1.7 | ↔ | -1.7 |
| ADAM15 | -1.6 | 1.4 | ↔ | ↔ | -1.4 | | -1.4 |
| ADAM17 | -1.4 | ↔ | ↔ | 1.4 | ↔ | ↔ | ↔ |
| ADAM19 | ↔ | 1.7 | -1.4 | ↔ | -1.7 | ↔ | ↔ |
| ADAM20 | -2.7 | 2 | ↔ | ↔ | ↔ | ↔ | ↔ |
| ADAM21 | ↔ | ↔ | -2 | ↔ | ↔ | ↔ | ↔ |
| ADAM22 | ↔ | ↔ | ↔ | ↔ | -1.9 | ↔ | ↔ |
| ADAM28 | ↔ | ↔ | (2; P=0.09) | ↔ | ↔ | ↔ | ↔ |
| ADAM33 | -2 | ↔ | ↔ | ↔ | ↔ | ↔ | ↔ |
| TIMP1 | ↔ | 2 | 2.5 | ↔ | ↔ | ↔ | ↔ |
| TIMP2 | -1.5 | ↔ | ↔ | ↔ | -1.5 | ↔ | -1.7 |
| TIMP3 | ↔ | -2 | -4 | ↔ | -1.6 | ↔ | ↔ |
| TIMP4 | ↔ | ↔ | ↔ | ↔ | ↔ | ↔ | -1.8 |
| IL6 | -2 | 77 | 55 | 2.7 | ↔ | ↔ | ↔ |
| IL10 | 5.9 | 65 | 13 | ↔ | ↔ | ↔ | ↔ |
| BDNF | -2.3 | -2.2 | -3.5 | ↔ | ↔ | ↔ | ↔ |
| NGF | -4 | ↔ | ↔ | ↔ | ↔ | ↔ | ↔ |
| NPY | ↔ | -4.2 | -3.5 | ↔ | -1.7 | ↔ | ↔ |
| Tachykinin 3 | ↔ | -2.5 | -2.1 | ↔ | ↔ | ↔ | ↔ |
| BDKRB1 | ↔ | 5.3 | 4.1 | 1.7 | ↔ | ↔ | ↔ |
| BDKRB2 | ↔ | 2.2 | 2 | ↔ | ↔ | ↔ | ↔ |

Notes: Significant fold changes are indicated compared to unstimulated cells in either 2D or 3D culture (or comparison between 2D and 3D for unstimulated cells only). ↔ indicates no change.

Abbreviations: IL1/OSM, interleukin 1/oncostatin M; subP, substance P; BK, bradykinin; ADAM, a disintegrin and metalloproteinase; TIMP, tissue inhibitor of metalloproteinase; IL6, interleukin 6; IL10, interleukin 10; BDNF, brain-derived neurotrophic factor; NGF, nerve growth factor; NPY, neuropeptide Y; BDKRB1, bradykinin receptor B1; BDKRB2, bradykinin receptor B2; TLDA, TaqMan low-density array.

was observed in primary chondrocytes in a 3D culture model similar to that used here⁶⁰ (indicative of chondrocyte redifferentiation) and we observed similar results in our cultures (data not shown). Repression of IL6 mRNA in 3D collagen gel culture (compared to culture on a thin coating of collagen) was observed previously in ovarian tumor cells.⁶¹ Future studies will reveal whether similar mechanisms are involved in downregulation of steady-state mRNA levels for the 11 genes affected by 3D culture in our studies. Uniquely among the genes explored in our study, the anti-inflammatory cytokine, IL10, was upregulated in cells cultured in 3D. In relation to this finding, it is interesting to note that fibroblasts cultured in a 3D matrix of enhanced stiffness in a wound healing context express greater levels of IL10.⁶² Taken together, our data on IL6 and IL10 suggest that chondrocytes in a 3D environment have an overall anti-inflammatory phenotype.

Effects of cytokines and neurovascular peptides

We exposed chondrocytes to a standard combination of IL1 α and OSM since these cytokines have previously been shown to

be present in synovial fluid of OA patients⁶³ and have long been known to act synergistically to induce complete breakdown of both bovine nasal cartilage⁶⁴ and human OA cartilage.⁶⁵

ADAMs

In 2D culture, cytokine treatment upregulated the steady-state mRNA levels for six ADAMs, several of which were previously reported to be expressed in chondrocytes or OA cartilage.^{31,32} In our 3D cultures, we no longer saw cytokine-mediated upregulation of ADAM10, ADAM15, and ADAM19, and this could reflect the overall suppressive nature of the 3D environment. ADAM12 mRNA was elevated by cytokines in both 2D and 3D culture, with overall expression levels approximately halved in 3D compared to monolayer culture. Recently an ADAM12 SNP associated with severity of knee OA was reported.⁴² ADAM12 (among other ADAMs) can shed the cell surface receptor LRP-1 which mediates endocytosis of TIMP3 and aggrecan-degrading ADAMTS5.⁶⁶ Thus, alterations in ADAM12 levels could indirectly have consequences on cartilage degradation.

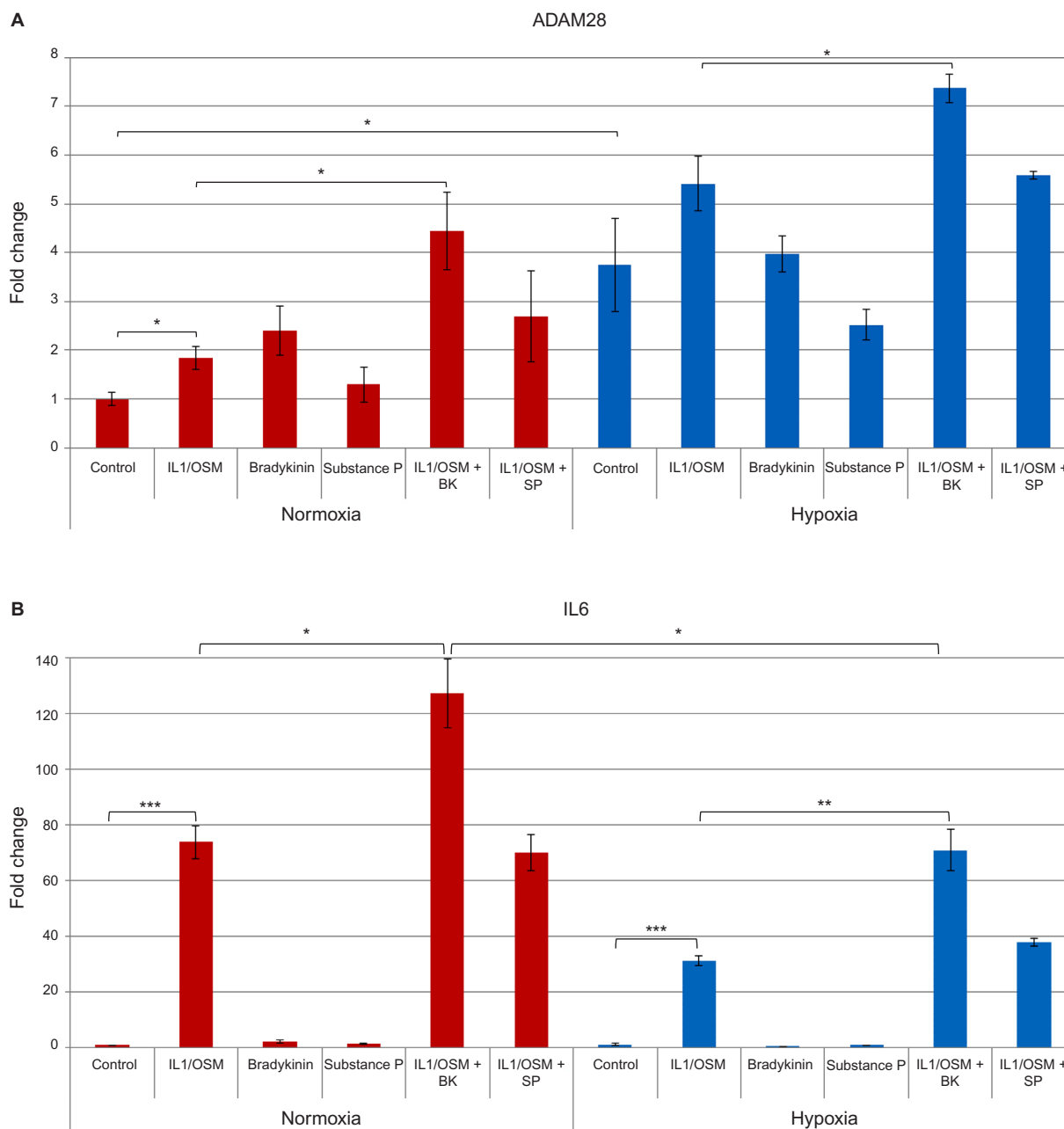


Figure 2 Confirmation of co-stimulatory effects of bradykinin and cytokines in chondrocytes cultured in 3D. **Notes:** (A) ADAM28 steady-state mRNA levels are elevated by a combination of cytokines and bradykinin under normoxic conditions (red bars) and are maintained under hypoxic conditions (blue). (B) IL6 expression is synergistically enhanced by the combination of cytokines and bradykinin under normoxic conditions (red) and overall responses are reduced under hypoxic conditions (blue). Steady-state mRNA levels for ADAM28 and IL6 were determined by qRT-PCR. For this fold-change analysis, control (unstimulated) cells were set as 1. Each bar represents the mean \pm SEM for triplicate samples following normalization to 18S RNA. * $P < 0.05$; ** $P < 0.02$; *** $P < 0.01$. **Abbreviations:** ADAM, a disintegrin and metalloproteinase; IL, interleukin; OSM, oncostatin M

In our initial survey, ADAM28 showed a trend toward increased expression following cytokine treatment in 3D culture (confirmed in our later experiment) and overall expression levels were not repressed in 3D, in contrast to other ADAMs. Cytokines regulate ADAM28 in monocytic cells⁵³ and ADAM28 expression is elevated in peripheral blood mononuclear cells in Kashin–Beck disease, characterized by chondrocyte necrosis.⁶⁷ We saw a trend toward increase

in expression with bradykinin treatment alone, but this did not reach significance.

TIMP3

TIMP3 expression was repressed following cytokine treatment, an effect seen in synovial cells⁶⁸ and observed in endothelial cells by us in previous studies.⁶⁹ Substance P also repressed TIMP3 mRNA levels in 3D culture, suggesting a

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potentially important role for this neuropeptide in cartilage homeostasis. TIMP3 deletion in mice results in the degradation of collagen and aggrecan over time and development of OA-like symptoms,⁷⁰ and very recently, a TIMP3 polymorphism associated with the severity of knee OA was reported,⁷¹ although any functional consequences remain unknown. As well as inhibiting many MPs involved in cartilage degradation (for example, ADAMTS5 and MMP13), TIMP3 also inhibits ADAM17 and other sheddases. Thus the repression of TIMP3 mRNA observed here may also have profound consequences on the cell surface ecology of chondrocytes.

Cytokines

The increase in steady-state mRNA levels for anti-inflammatory IL10 seen in response to cytokines in both 2D and 3D may reflect an attempt to counteract the damaging effects of proinflammatory cytokine stimulation. Exogenous IL10 can protect against cartilage damage⁷² and IL10 levels are upregulated in OA and postulated to be a prorepair response.⁷³ Cytokine upregulation of IL6 is in agreement with previous findings in OSM-treated mouse articular chondrocytes, for example, where a hierarchy of responses suggests that OSM may be a better therapeutic target than IL6.⁷⁴ Cytokines repressed neurotrophin BDNF expression as well as that of NPY and tachykinin 3 (both in 2D and in 3D). Perhaps of relevance is the finding that levels of BDNF are enhanced after exercise in patients with knee OA and may have anti-inflammatory benefits with corresponding improved pain scores.⁷⁵

Bradykinin and cytokine co-stimulation: effects on ADAM28 and IL6 expression

Enhanced expression of ADAM28 in the presence of cytokines and bradykinin may be due to the increased bradykinin receptor expression observed in the presence of cytokines alone. The synergistic increase in IL6 expression with cytokine and bradykinin co-treatment in normoxia is reminiscent of that seen in an early study in gingival fibroblasts where IL6 protein production was elevated by stimulation with bradykinin and IL1 or TNF α .⁷⁶ Similar synergistic effects of bradykinin and IL1 β on the expression of COX-2 and PGE-2 in synoviocytes have been observed.⁷⁷ These authors demonstrated that this effect was mediated by bradykinin receptors, using the inhibitor fasitibant, and was dependent on the NF κ B pathway. Future studies will determine mechanisms underpinning the upregulation of steady-state mRNA levels for ADAM28 and IL6 seen with bradykinin and cytokine co-stimulation in chondrocytes.

In our preliminary study, hypoxia exerted a positive effect on both constitutive and cytokine-induced ADAM28 mRNA levels but was suppressive for IL6 (and for the expression of several other genes; data not shown). Several studies have explored IL6 gene expression and/or protein secretion under various levels of hypoxia with 3% oxygen resulting in upregulation of IL6 secretion in human synovial cells,⁷⁸ but 5% oxygen leading to downregulation in placental cells.⁷⁹ Previous work showed that hypoxia (1% oxygen as used in our study) promotes chondrocyte differentiation⁷ and perhaps the repression of IL6 under hypoxic conditions is anti-inflammatory and prorepair.

Overall our experiments demonstrate interesting and novel regulation of steady-state mRNA levels for ADAMs and cytokines in C28/I2 chondrocytes, a readily available immortalized human culture model. Future experiments will reveal mechanisms underpinning our observations as well as any parallels in primary human chondrocytes and cartilage.

Limitations of the study

While we have demonstrated novel effects of neurovascular peptides and hypoxic conditions on ADAM and cytokine steady-state mRNA levels, we have not confirmed effects at the protein level nor have we explored functional consequences of any changes in expression. Future studies following confirmation of protein levels would include functional assessment of enzyme activity with relevant substrates and gene silencing approaches to determine, for example, ADAM28's role in proteoglycan degradation/cell proliferation in chondrocytes under bradykinin and cytokine stimulation.

Conclusion

This study reveals the novel regulation of ADAM proteinases at the mRNA level by a 3D culture environment which also modulates cytokine responses in chondrocytes. In addition, steady-state mRNA levels of ADAM28 and IL6 are elevated by bradykinin, in the presence of cytokines, which may represent an important modulator of the chondrocyte milieu. These findings pave the way for future studies of the impact of bradykinin and downstream pathways on cartilage function under inflammatory stress.

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Disclosure

The authors report no conflicts of interest in this work.

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