Human articular chondrocytes express functional Leukotriene B4 receptors

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Running title: LTB4 receptors in chondrocytes

Summary

Leukotriene B4 (LTB₄) is a potent chemoattractant associated with the development of osteoarthritis (OA), while its receptors BLT1 and BLT2 have been found in synovium and subchondral bone. In this study, we have investigated whether these receptors are also expressed by human cartilage cells and their potential effects on cartilage cells. The expression of LTB₄ receptors in native tissue and cultured cells was assessed by immunohistochemistry, immunocytochemistry, PCR and electron microscopy. The functional significance of the LTB₄ receptor expression was studied by Western blotting, using phosphospecific antibodies in the presence or absence of receptor antagonists. In further studies, the secretion of pro-inflammatory cytokines, growth factors and metalloproteinases by LTB₄ - stimulated chondrocytes was measured by multiplex protein assays. The effects of LTB₄ in cartilage signature gene expression in cultured cells were assessed by qPCR, whereas the LTB₄-promoted matrix synthesis was determined using 3D pellet cultures.

Both receptors were present in cultured chondrocytes, as was confirmed by immunolabelling and PCR. The relative quantification by PCR demonstrated a higher expression of the receptors in cells from healthy joints compared with OA cases. The stimulation of cultured chondrocytes with LTB₄ resulted in a phosphorylation of downstream transcription factor Erk 1/2, which was reduced after blocking BLT1 signalling. No alteration in the secretion of cytokine and metalloproteinases was recorded after challenging cultured cells with LTB₄; likewise, cartilage matrix gene expression and 3D tissue synthesis were unaffected. Chondrocytes express BLT1 and BLT2 receptors, and LTB₄ activates the downstream Erk 1/2 pathway by engaging the high-affinity receptor BLT1. However, any putative role in cartilage biology could not be revealed, and remains to be clarified.

Keywords: leukotriene B₄, BLT1, BLT2, cartilage, chondrocytes, osteoarthritis, inflammation

Introduction

The metabolism of arachidonic acid by the cyclooxygenase (COX) or 5-lipoxygenase (5-LO) pathways generates eicosanoids, which have been implicated in the pathophysiology of a variety of human diseases, including arthritis of the joints (Molloy and McCarthy, 2005; Korotkova and Jakobsson, 2014).

Arachidonic acid released from cell membranes by events such as inflammation or trauma can be converted to unstable precursor leukotriene A₄ (LTA₄) via the 5-LO and 5-lipoxygenase activating protein (FLAP), and further into leukotriene B₄ (LTB₄) by LTA₄ hydrolase (Peters-Golden and Henderson, 2007). In the presence of COX-1 or COX-2, arachidonic acid is converted into prostaglandin H₂, and subsequently into prostaglandin E₂ by prostaglandin E synthase (Li et al., 2009). Leukotriene B₄ (LTB₄) is a powerful inflammatory mediator implicated in diseases such as asthma, atherosclerosis and rheumatoid arthritis by the recruitment and activation of leukocytes (Tager and Luster, 2003; Peters-Golden and Henderson, 2007; Di Gennaro and Haeggstrom, 2014). Importantly, LTB₄ levels are enhanced in synovial tissue and -fluid, cartilage and subchondral bone during both osteoarthritis (OA) and rheumatoid arthritis (RA) (Ahmadzadeh et al., 1991; Wittenberg et al., 1993; Paredes et al., 2002; Attur et al., 2012).

LTB₄ binds two G-coupled receptors: the high-affinity leukotriene B₄ receptor 1 (BLT1) and the low-affinity leukotriene B₄ receptor 2 (BLT2) (Yokomizo et al., 2001a). The BLT1 receptor was initially described in humans in myeloid cells (Yokomizo et al., 1997), though later studies have reported its presence in non-myeloid tissues as well, including subchondral osteoblasts and synovial cells (Paredes et al., 2002; Hashimoto et al., 2003). BLT1 activation in non-myeloid cells promotes the cell differentiation, migration, proliferation and chemokine production needed to recruit neutrophils (Kim and Luster, 2007). BLT2 is ubiquitously

3

expressed in humans (Yokomizo et al., 2000), while in synovial tissue the expression of BLT2 surpasses that of BLT1 (Hashimoto et al., 2003).

Prostaglandins are also involved in OA development (Amin et al., 1997; Li et al., 2009), with the current first-line treatment of OA including COX inhibitors (McAlindon et al., 2014). However, blocking COX-regulated arachidonic acid pathways leads to a shunting of the system to the 5-LO pathway, as well as an undesired elevation of LTB₄ levels (Marcouiller et al., 2005; Maxis et al., 2006; Attur et al., 2012).

Despite studies arguing for the importance of LTB_4 and its receptors on RA and OA disease development, the expression and function of LTB_4 receptors in chondrocytes has not yet been reported. The aim of this study was to assess the expression of LTB_4 receptors BLT1 and BLT2 in cartilage and the biological effects of LTB_4 on cultured chondrocytes.

Material and method

Human material and cell culturing

Macroscopically healthy looking cartilage was harvested from the lateral femoral condyle of 15 patients (age 55-82) undergoing total knee replacement (TKR), and the overall grading of cartilage according to the ICRS/Outerbridge score (Kleemann et al., 2005) was 2-4. In addition, we obtained surplus cells from the autologous chondrocyte implantation (ACI) of 10 patients (aged 31-65); these cells were harvested and expanded as previously described (Brittberg et al., 1994; Peterson et al., 2002), with the procedure also demanding a normal joint on X-ray and arthroscopy (Knutsen et al., 2004). The patients participated by informed consent, and the Regional Ethical Committee of North Norway approved the study. Cartilage from TKR was kept in a sterile saline and processed within two hours from harvest. Specimens were processed directly as described for immunohistochemistry and electron microscopy, enzymatically digested for the determination of BLT receptors and signature genes by qPCR or processed for expansion in culture by mincing into ~1mm³ pieces, followed

by digestion in 0.8 mg/mL of collagenase XI solution (Cat. no. C9407, Sigma-Aldrich) for four hours at 37°C during gentle agitation. Chondrocytes were plated in 25cm² culture flasks containing Dulbecco's Modified Eagle's Medium/Ham's F-12 Mixture (DMEM/F-12), and supplemented with ascorbic acid (62 mg/L), penicillin (2500 U/L), streptomycin (2.5 mg/L) and 20% fetal bovine serum (FBS, Cat. no. S0115, Biochrome). After cell attachment, cultures were further propagated in a medium supplemented with 10% serum and the cell number was serially expanded in appropriate vessels, and subsequently used for experimentation at passages 3-4.

Immunohistochemistry and immunocytochemistry

The immunolabelling of BLT1 and BLT2 receptors in native cartilage and cultured chondrocytes was performed using polyclonal rabbit anti-human BLT1 antibody (Cat. no. 120114, Cayman Chemicals), and/or polyclonal rabbit anti-human BLT2 antibody (Cat. no. 120124, Cayman Chemicals) or no primary antibody for control. For immunohistochemistry, biopsies from five TKR donors were fixed in paraformaldehyde and prepared as previously described (Berg et al., 2010). Sections were incubated overnight at 4°C in BLT1 or BLT2 antibody diluted at 1:100. Slides were stained using an Anti-rabbit HRP SuperPicTure[™] Polymer detection kit (Cat. no. 87-9263, Invitrogen), and counterstained with haematoxylin. Isotype control was used to assess non-specific binding. For immunocytochemistry, an appropriate number of expanded chondrocytes from four TKR donors were seeded on fibronectin-coated slides as previously described (Berg et al., 2010). Slides were incubated overnight at 4°C in BLT1 or BLT2 antibody diluted at 1:250. After washing in PBS, the slides were incubated with secondary antibody Alexa Fluor® Goat Anti-Rabbit IgG (Cat. no. A11008, Invitrogen) at 4°C for four hours until nuclear staining and mounting using DAPI-Fluoromount-GTM (Cat. no. 0100-20, Southern Biotech). The slides were examined in a Zeiss Axiophot photomicroscope (Carl Zeiss, Oberkochen, Germany).

RT-PCR of BLT1 and BLT2 mRNAs

As previously described, a reverse-transcription PCR was performed (Berg et al., 2010). In brief, total RNA was extracted from two different TKR cell cultures using an RNeasy® Mini Kit (Cat. no. 74106, Qiagen), and converted to cDNA using a Superscript® III Reverse Transcriptase (Cat. no. 18080-093, Invitrogen). The PCR reaction volume of 50 μ L included 5 μ L cDNA, 150 nM of each specific primer, ultrapure water and MasterMix (Cat. no PRME2200100, 5 Prime). Sequences for the specific primers for human BLT1 (sense 5'-TATGTCTGCGGAGTCAGCATGTACGC-3', antisense 5'-

CCTGTAGCCGACGCCCTATGTCCG-3') yields a 345 bp product, while human BLT2 (sense 5'-AGCCTGGAGACTCTGACCGCTTTCG-3', antisense 5'-

GACGTAGAGCACCGGGTTGACGCTA-3') yields a 320 bp product. Adenine phosphoribosyltransferase (APRT) (sense 5'-CCCGAGGCTTCCTCTTTG GC-3', antisense 5'-CTCCCTGCCCTTAAGCGAGG-3') which yields a 300 bp product, was included for RNA quality control; in case of contaminating DNA, a 800 bp product would occur. The PCR was run on a GeneAmp® PCR System 9700 (Applied Biosystems), with an amplification profile of: 7 minutes at 94°C (denaturation), 30 seconds at 94°C (denaturation), 30 seconds at 55°C (annealing), 1 minute at 72°C (extension) and 30 cycles completed with a 10-minute extension at 72°C. RT-PCR products were separated on Novex® 6% TBE gel (Cat. no. EC6265BOX, Invitrogen) and stained with SYBR® Safe DNA gel stain (Cat. no. S33102, Invitrogen), while images were developed using a G-Box (Syngene). To help confirm that BLT1 and BLT2 transcripts matched those reported in GeneBank®, the PCR products were sequenced using a BigDye® Terminator v3.1 Cycle Sequencing Kit (Cat.no 4337455, Applied Biosystems) and GeneAmp® PCR Systems 9700 (Applied Biosystems).

Immuno Electron microscopy

Cartilage specimens from two different TKR donors were freshly collected, washed in icecold PBS and fixed in 4% formaldehyde in a 200 mM HEPES buffer, pH 7.5, for 24 hours. Cryosections were prepared, and immunolabelling was performed as described elsewhere (Tokuyasu, 1986). Antibodies against BLT1 (diluted 1:20) were detected by protein A-gold complexes, and the dried sections were then examined in a JEOL JEM 1010 transmission electron microscope (JEOL, Tokyo, Japan) operating at 80 kV. Additionally, negative controls were routinely included in parallel by the omission of primary antibodies.

Relative quantification of BLT1 and BLT2 receptor by qPCR

RNA was extracted from three sources: cartilage biopsies from total knee replacement biopsies (N=3), cultured cells from total knee replacement biopsies at passage 4 (N=5) and surplus cells from ACI at passage 4 (N=5). For RNA extraction from cartilage, the biopsies were minced to $\sim 1 \text{ mm}^3$ pieces, and chondrocytes were then rapidly released by digestion in a 0.25% trypsin solution (Cat. no. T3924, Sigma-Aldrich) for 60 minutes and then in a 0.8 mg/mL collagenase XI solution for six hours at 37°C during gentle agitation. The freshly isolated cells were lysed using Qiazol, followed by DNase I treatment and cleanup on RNeasy mini-spin columns (Cat. no. 74804, Qiagen) according to the manufacturer's instructions. Cultured cells were then extracted using a PerfectPure RNA Cultured Cell Kit (Cat. no. 2302340, 5 prime), including DNase I treatment according to the manufacturer's instructions. RNA was quantified by spectrophotometry (Nano Drop ND-1000), and diluted at 1:5 to a final concentration of 9.82 ng/µL before reverse transcription (RT) using a High Capacity RNA to cDNA kit (Cat. no. 4387406, Invitrogen) according to the manufacturer's instructions. The relative quantification experiment was run with hydrolysis probes targeting BLT1 (LTB4R, Hs01938704_s1) and BLT2 (LTB4R2, Hs01885851_s1). Based on previous validation experiments, ribosomal protein L13A (RPL13A, Hs04194366_g, all probes from

Life Technologies) was used as the reference gene. The reaction volume of 10 μ L included 5 μ L TaqMan® Fast Universal PCR Mastermix No AmpErase® UNG (Cat. no. 4366072, Applied Biosystems), 2.5 μ L water, 2 μ L cDNA (50 ng/ μ L) and 0.5 μ L of the respective assays. Three technical replicates of each sample were applied to a MicroAmp® Fast Optical 96-well reaction plate (Cat. no. 4346906, Applied Biosystems), and water and no-RT samples were added for the negative control. The plates were run on StepOnePlus, using the following cycling conditions: hold at 95°C for 20 seconds, 40 cycles at 95°C for one second and then at 60°C for 20 seconds. Data was analysed in Microsoft Excel, a t-test was used to test significance and the differences were considered significant at p < 0.05.

Western blotting of phosphorylated protein kinases

Cultured chondrocytes from three ACI donors were starved for 24 hours before LTB₄ (Cat. no. 20110, Cayman Chemicals) was added at a final concentration of 10^{-8} M for 3 and 5 minutes each. For inhibition studies, cultured chondrocytes were challenged with BLT1 receptor antagonist U-75302 at 3 x 10^{-7} M (Cat. no. 70705, Cayman Chemicals) or BLT2 receptor antagonist LY2552833 at 10^{-6} M (Cat. no. 70715, Cayman Chemicals) for 1 hour before being stimulated with LTB₄ for 5 minutes (Yokomizo et al., 2001b; Lundeen et al., 2006). Cells were lysed in a 150 µL radioimmunoprecipitation (RIPA) buffer (Cat. no. 20-188, Millipore), including protease- and phosphatase inhibitor, and an equal amount of protein was separated with gel electrophoresis and blotted to PVDF membranes. Membranes were incubated overnight at 4°C with Phospho-p44/42 MAPK (Erk 1/2) Rabbit mAb (Cat. no. 4370, Cell Signaling) and loading control β -actin (13E5) Rabbit mAb (Cat. no. 4970, Cell Signaling) at a final dilution of 1:2000. Next, the membranes were incubated for 1 hour with Anti-rabbit IgG HRP-linked Antibody (Cat. no. 7074, Cell Signaling), and bands were detected using chemiluminescence.

Multiplex protein analyses

The protein concentration in culture supernatants was analysed by antibody-based multiplex protein arrays. Chondrocytes from three different TKR cultures were seeded on 24-well plates at a concentration of 1.0×10^5 cells/well, and incubated overnight in a medium with a 10% FBS. Thereafter the cells were starved overnight in a medium containing 0.1% FBS before challenging with LTB₄ at a concentration of 10^{-12} M, 10^{-10} M, 10^{-8} M or vehicle only at 37°C for 18 hours. Vascular endothelial growth factor (VEGF), basic fibroblast growth factor (bFGF) and platelet-derived growth factor (PDGF), along with inflammatory cytokines IL-1 β , TNF- α , IL-6, IL-8 and MCP-1, were analysed in an 8-plex assay (Cat. no. Z50007ZORG, Bio Rad). Matrix metalloproteinases (MMP) of -1, -2, -3, -8, -9 and -13 were measured using a multiplex MMP assay (Cat .no. LMP000, R&D Systems). Samples were diluted at 1:4 and analysed in duplicate using the Bio-Plex 200 System (Bio-Rad). The results are the average of two independent experiments, and were analysed in Excel using a one-way ANOVA.

qPCR of cartilage signature genes

Surplus cultured chondrocytes from four ACI procedures were grown in parallel, in which one culture remained untreated for the control, and the other stimulated with LTB₄ at a final concentration of 10⁻⁸ M for 18 hours. Chondrocytes were harvested in a lysis solution and RNA extracted using the PerfectPure RNA Cultured Cell Kit (Cat. no. 2302340, 5 prime) according to the manufacturer's instructions, including DNase I treatment. RNA was quantified using Nano Drop ND-1000, and converted to cDNA with a High Capacity RNA to cDNA kit (Cat. no. 4387406, Invitrogen) according to the manufacturer's instructions. For the qPCR, the following pre-optimized hydrolysis probes from Applied Biosystems were used: Collagen 1A1 (COL1A1, Assay ID: Hs00164004_m1), Collagen 2A1 (COL2A1, Assay ID: Hs00264051_m1), Aggrecan (ACAN, Assay ID: Hs00153936_m1) and SOX9 (SOX9, Assay ID: Hs00165814_m1). Glyceraldehyde 3-phosphate dehydrogenase (GAPDH, Assay ID: Hs99999905_m1) was included as a reference gene after validating a stable expression in monolayer cultures. A reaction volume of 20 μ L contained 10 μ L of TaqMan® Fast Universal PCR Mastermix No AmpErase® UNG (Cat. no. 4366072, Applied Biosystems), 7 μ L of water, 2 μ L of cDNA (50 ng/ μ L) and 1 μ L of the respective assays. Samples in three technical replicates were applied to a MicroAmp® Fast Optical 96-well reaction plate (Cat. no. 4346906, Applied Biosystems); water for a negative control and cDNA from freshly isolated cartilage cells were added in triplicate for positive control and interplate calibration. Plates were analysed on an Applied Biosystems 7900HT Fast Real-Time PCR System with the following cycling conditions: hold at 95°C for 20 seconds, 40 cycles at 95°C for one second and then 60°C for 20 seconds.

3D Pellet cultures

Surplus ACI chondrocytes expanded in monolayers were detached by an enzyme-free detachment solution (Cat. no. S-004-B, Millipore) and a short (30 seconds) trypsinization. Cells were then resuspended in a complete growth medium, and 150 µL aliquots (50000 cells) were added to NuncTM 96-well polystyrene conical bottom MicroWellTM plates (Cat. no. 249935, Thermo Scientific) and centrifuged at 400 x g for 10 minutes to make pellets. After two days, rounded shaped and consistent spheroids were transferred to a low attachment 24-well plate in a serum-free medium containing ITS (Insulin-Transferrin-Selenite, Cat. no. 354352, BD Biosciences). For the treated group, 10⁻⁸ M LTB₄ was added to the medium. Spheroids were further incubated for six days, and the medium with or without LTB₄ was changed every second day. Lastly, spheroids were collected and prepared for histology as described under immunohistochemistry. To evaluate the matrix synthesis, slides were stained with Alcian blue to detect glycosaminoglycans or a collagen type II antibody.

Results

Initial experiments were performed to assess the expression of LTB₄ receptors in cartilage tissue and cells, and native cartilage sections from five different donors were immunolabelled

10

using BLT1 or BLT2 antibodies. A positive brown staining confirmed the presence of BLT1 receptor (Figure 1a) and BLT2 receptor (Figure 1b), while immunocytochemistry using respective antibodies showed that cultured chondrocytes at passages 3-4 also expressed BLT1 (Figure 2a) and BLT2 receptors (Figure 2b). The expression of mRNA encoding these receptors was confirmed by RT-PCR using specific BLT1 and BLT2 primers, as shown in Figures 3a and 3b, respectively. The sequencing of the products revealed an identical sequence to the GeneBank sequence for BLT1 and BLT2 (data not shown). The subcellular distribution of the BLT1 receptor was assessed by immune electron microscopy, and the specific detection of BLT1 by immune-gold labelling exhibited a predominant expression of the receptor in the plasma membrane of chondrocytes from native tissue, and to a much lesser extent in cytoplasm (Figures 4 A-D). The relative qPCR revealed that the expression of BLT1 was significantly higher in cultured chondrocytes from non-OA joints (ACI) than cartilage biopsies from OA cases (TKR) (Figure 5 A). Likewise, the gene expression of BLT1 was significantly higher in cartilage compared to cultured cells from TKR (Figure 5 C). The expression of BLT2 was higher in cultured ACI cells compared to cultured TKR cells, but equally expressed in expanded TKR cells and cartilage (Figure 5 B, D).

In order to assess the BLT1 and BLT2 downstream signalling, cells were stimulated with 10^{-8} M LTB₄, followed by the Western blotting of Phospho-Erk 1/2 (p44/42 MAPK). Figure 6 shows a modest background phosphorylation of Erk 1/2 in untreated chondrocytes (Lane 1); however, an LTB₄ challenge for five minutes markedly increased phosphorylation (Lane 3). The blocking of the BLT1 receptor with the specific receptor antagonist U-75302 before the LTB₄ challenge reduced the signal to background phosphorylation (Lane 4). In contrast, a specific blockage of the BLT2 receptor with the antagonist LY2552833 had no major impact (Lane 5), whereas concurrent blocking of the receptors conferred the blocking of BLT1 alone (Lane 6).

Having confirmed the expression of functional BLT1 in chondrocytes, we assessed the potential effects of LTB₄ on major biological parameters in subsequent experiments. A multiplex analysis of supernatants of chondrocytes cultured in the presence of LTB₄ at increasing doses revealed no significant change in the secretion of growth factors, proinflammatory cytokines or MMPs. Levels of bFGF, PDGF, IL-1 β , TNF- α , MMPs 1, 2, 8 and 9 were undetectable, with values falling under the detection limit of the assay in both the treated and untreated groups. VEGF, IL-6, IL-8, MCP-1 and MMPs 3 and 13 were detected, although LTB₄ did not significantly alter the secretion of these molecules when analysed using ANOVA (Figure 7).

To assess the potential effect of LTB₄ on chondrocytes differentiation, the expression of some cartilage signature genes, including Aggrecan, Sox-9, Collagen 1A1 and Collagen 2A1, were analysed using qPCR. Four biological replicates were included, and data were normalized against GAPDH. Relative gene expression in chondrocytes stimulated with 10^{-8} M LTB₄ normalized to untreated controls is illustrated in Figure 8. Furthermore, stimulating cultured chondrocytes with LTB₄ did not alter the expression of specific cartilage signature matrix genes.

Lastly, the cartilage-forming capability of cultured chondrocytes was assessed in the presence or absence of 10^{-8} M LTB₄. After six days, tissue with the characteristics of "immature" cartilage was formed in both settings, revealing similar features regarding size, shape, the amount of extracellular matrix and cell density. Specific staining with Alcian blue for the detection of glycosaminoglycans, and with anti-collagen type II antibodies to evaluate the LTB₄ effect on matrix synthesis, demonstrated no observable differences (Figure 9).

Discussion

A potential role of LTB₄ in RA and OA has been suggested based on cell culture and tissue models, exhibiting an increased production of LTB₄ in RA and OA joints, along with reports

demonstrating the presence of BLT1 and BLT2 receptors in synovial tissue and subchondral bone (Paredes et al., 2002; Hashimoto et al., 2003; Attur et al., 2012). Moreover, LTB4 receptor knock-out mice were protected from rheumatoid arthritis development (Kim et al., 2006; Mathis et al., 2010). In the present study, we show for the first time that the LTB4 receptors BLT1 and BLT2 are expressed by human chondrocytes, both *in vivo* and *in vitro*. Additionally, subcellular localization studies of cartilage tissue using cryo-immuno electron microscopy unveiled a predominant expression of BLT1 at the plasma membrane in chondrocytes from native tissue.

One of the important findings of our study was the differential expression of BLT1 and BLT2 in chondrocytes from OA vs. healthy patients. A quantitative PCR analysis revealed that both BLT1 and BLT2 expression was higher in cartilage cells from non-diseased joints. This downregulation of receptor expression in diseased tissue could represent a protective mechanism to help ameliorate tissue damage caused by chronic inflammation. Likewise, the expression of the BLT1 receptor diminished during cell expansion in monolayers (Figure 5), with this outcome possibly being part of the overall cellular dedifferentiation process undergone by chondrocytes during expansion in monolayer cultures.

Previous studies with non-myeloid cells have reported that LTB₄ stimulation results in the downstream phosphorylation of signal transduction factors such as Akt and Erk 1/2, as well as the subsequent enhancement of migration and proliferation (Lundeen et al., 2006; Watanabe et al., 2009). In the present study, we could see evidence of an increased downstream phosphorylation of the BLT1 receptor when chondrocytes were stimulated with LTB₄. The signal appears to be mediated via the BLT1 receptor, as blocking with the BLT1 receptor antagonist U-75302 led to less phosphorylation of Erk 1/2, while blocking BLT2 did not (Figure 6).

13

Earlier studies have shown that chondrocytes secrete a wide range of growth factors, cytokines and MMPs (Berg et al., 2010; Polacek et al., 2010; Polacek et al., 2011). A critical phenomenon associated with the onset and progression of OA is the enhanced expression of MMPs within the joint tissue (Goldring and Marcu, 2009), with pro-inflammatory cytokines such as IL-1 β and TNF- α being strong inducers of MMP production and driving forces in cartilage deterioration (Martel-Pelletier, 2004). The endogenous synthesis of LTB_4 by synovium is a crucial element involved in the upregulation of IL-1 β and TNF α in OA (He et al., 2002). We therefore hypothesized that the secretion of MMPs, cytokines and/or growth factors would be affected by LTB₄ stimulation. In our analyses, MMP-3, MMP-13, VEGF, IL-6, IL-8 and MCP-1 were readily detected in culture supernatants, though their synthesis was not altered by LTB₄ at any dose tested (Figure 7). The secretion of other mediators, including TNF- α and IL-1 β , was below the detectable range. *In vivo*, the production of MMPs is promoted by IL-1 β and TNF- α (Martel-Pelletier et al., 2004). Thus, a reasonable assumption is that the levels of these pro-inflammatory cytokines were insufficient in our assays, or that other synergistic factors required were not present in our *in vitro* systems. The most prominent physiological role of chondrocytes is the homeostasis of the extracellular matrix and the maintenance of cartilage integrity (Goldring and Marcu, 2009). In this regard, the gene expression of specific cartilage matrix molecules is used as a functional marker of chondrocyte phenotype and differentiation status. It has previously been reported that in cultured chondrocytes stimulated with PGE₂, aggrecan gene expression was significantly downregulated, whereas the gene encoding collagen type II remained unchanged (Attur et al., 2008). Our data revealed an unaltered gene expression of collagen type 1A1, collagen type 2A1, aggrecan or Sox9 after 18 hours of incubation with LTB₄ (Figure 8). The decreased expression of BLT1 receptors observed after monolayer expansion might be the cause of the lack of effect upon LTB₄ challenge.

When chondrocytes are cultured in monolayers, they gradually dedifferentiate into a more fibroblast-like cell, a transformation characterized by an upregulation of collagen I and a downregulation of collagen II. In pellet cultures, the cells partially regain their original phenotypic traits in a process involving the downregulation of mitotic activity and the upregulation of matrix synthesis (Martinez et al., 2008). Intriguingly, the treatment of chondrocytes in 3D pellets with the PGE₂ has been shown to reduce collagen types I and II content (Jakob et al., 2004). We hypothesized that adding LTB₄ during 3D cultures could affect chondrocyte redifferentiation and matrix synthesis, but in our hands there was no marked difference in the morphological or immunohistochemical characteristics of the 3D constructs (Figure 9).

Concluding remarks

We have demonstrated that human articular chondrocytes express LTB₄ receptors BLT1 and BLT2, and that at least the BLT1 receptor is functional, as shown by its capability to activate intracellular signalling pathways upon ligand binding. The physiological role of these receptors in human chondrocytes, particularly during joint inflammation, requires further investigation.

List of abbreviations

- ACI Autologous chondrocyte implantation
- BLT1 Leukotriene B₄ receptor 1
- BLT2 Leukotriene B₄ receptor 2
- COX Cyclooxygenase
- COX-1 Cyclooxygenase-1, most common isoform of cyclooxygenase
- COX-2 Cyclooxygenase-2, isoform particularly found at sites of inflammation
- FLAP 5-lipoxygenase activating protein
- ICRS International Cartilage Repair Society
- LTA₄ Leukotriene A₄
- LTA₄H Leukotriene A₄ hydrolase
- LTB₄ Leukotriene B₄
- MMP Matrix metalloproteinase
- OA Osteoarthritis
- PGE₂ Prostaglandin E₂
- RA Rheumatoid arthritis

TKR Total knee replacement

5-LO 5-lipoxygenase

Conflict of interests

The authors have no competing interest to declare.

Authors contributions

AKH carried out the quantitative PCR, Western blotting, interpretation of data and the drafting of the manuscript. JTI performed the PCR, the immunoassays and the interpretation of the data. IM carried out the electron microscopy and 3D culture assay, interpreted the data and drafted the manuscript. YF participated in the study design and coordination, helped to draft the manuscript, interpret the results and revise the manuscript. BS participated in the design and coordination, helped to draft the manuscript, performed the immunohistochemistry and revised the manuscript. All authors read and approved the final manuscript.

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Figure legends

Figure 1: Expression of BLT1 and BLT2 in cartilage tissue (60x micrograph). Cartilage tissue sections stained with BLT1 antibody (1a), BLT2 antibody (1b) and isotype control (1c).

Figure 2: Expression of BLT1 and BLT2 in cultured chondrocytes.

Immunofluorescence microscopy of cultured chondrocytes stained with BLT1 antibody (2a), BLT2 antibody (2b). Secondary antibody conjugated with Alexa Fluor488 and DAPI stained nuclei. Isotype controls had no staining (not shown). Scalebar = $50\mu m$.

Figure 3: Expression of BLT1 and BLT2 mRNA in cultured chondrocytes by Reverse

Transcriptase PCR. **3a**: Lane L: DNA ladder, Lanes 1 and 2: mRNA control, APRT primer yields 300 bp bands and no trace of the 800 bp bands that appears when APRT is primed to genomic DNA (Lane 7). Lanes 4 and 5: cDNA from two donors and specific BLT1 primers yield 345 bp bands. Lanes 3 and 6: Negative control. **3b**: Lane L: DNA ladder. Lane 1: mRNA control. Lane 2: Negative control. Lanes 3 and 4: cDNA from two donors and specific BLT2 primers yield 320 bp bands.

Figure 4: Subcellular localization of BLT1 receptors in cartilage cells by immune-cryo electron microscopy. 4a and 4c: Two different cartilage sections stained with gold-conjugated anti-BLT1 antibody binding to BLT1 receptor in plasma membrane Panels 4b and 4d are enlarged windows of 4a and 4c, respectively. Gold particles are marked with arrow heads. CP = cytoplasm, N = nucleus, PM = plasma membrane, ECM = extracellular matrix. Scale bars = 1µm

Figure 5: Differential expression of BLT1 and BLT2 mRNA in cartilage cells from OA vs non-OA. Expression of BLT1 (5A) and BLT2 (5B) in cultured cells from five healthy ACI donors (Non-OA) compared to cultured cells from five TKR (OA) donors. Panels 5C and 5D

is a comparison of cartilage from TKR to cultured cells from TKR for BLT1 and BLT2, respectively.

Figure 6: Functional assessment of BLT receptors by Western blot. p-Erk 1/2 (44/42 kD). Lane L: Biotinylated ladder. Lanes 1-3: Erk 1/2 phosphorylation at 0, 3 and 5 minutes exposure to LTB₄. Lanes 4: Phosphorylation of Erk 1/2 in presence of BLT1 receptor antagonist (U-75302 at 3 x 10⁻⁷ M). Lane 5: Phosphorylation of Erk 1/2 in presence of BLT2 receptor antagonist (LY2552833 at 10⁻⁶ M). Lane 6: Phosphorylation of Erk 1/2 in presence of both receptor antagonists. Beta-actin (45kD) was used as loading control.

Figure 7: Secretion of growth factors, MMPs and inflammatory mediators after stimulation of chondrocytes by LTB4. Levels measured in supernatants from cultured chondrocytes after 24 hours of stimulation with LTB4 at concentrations of 10⁻¹²M, 10⁻¹⁰M, 10⁻⁸M and vehicle.

Figure 8: Cartilage signature gene expression by qPCR. Relative expression of cartilage signature genes in cultured chondrocytes challenged with LTB₄ for 18h. Results are normalized to untreated chondrocytes. A cartilage sample was included to visualize expression in cartilage vs. cultured cells. P-values are a result of t-testing cartilage vs. control and treated vs. control, respectively.

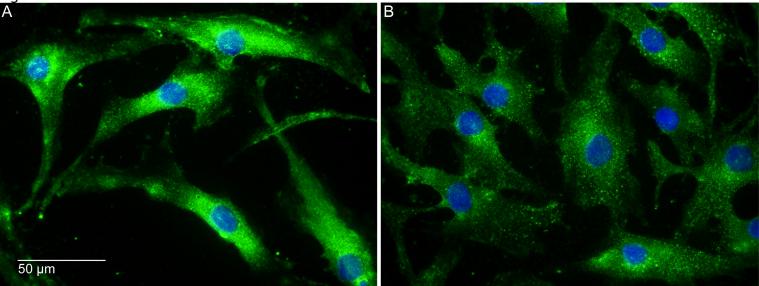
Figure 9: Histology and immunohistochemistry of scaffold-free 3D chondrocyte

spheroids. Chondrocyte pellets incubated for six days with LTB₄ at 10^{-7} M vs. unchallenged controls. Brown colour corresponds to expression of collagen type II, whereas the blue colour is indicative of matrix proteoglycans. The folds in the upper left- and lower right image are artefacts of preparation. Scalebar = 200 µm.

Figure 1



Figure 2



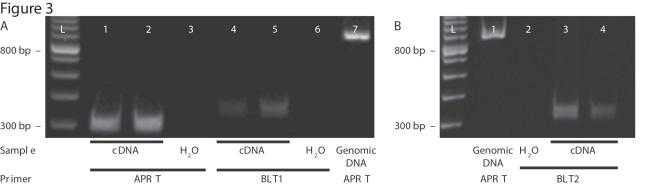


Figure 4

