Research article

Prostaglandin E2 synthesis in cartilage explants under compression: mPGES-1 is a mechanosensitive gene

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Abstract

Knee osteoarthritis (OA) results, at least in part, from overloading and inflammation leading to cartilage degradation. Prostaglandin E2 (PGE2) is one of the main catabolic factors involved in OA. Its synthesis is the result of cyclooxygenase (COX) and prostaglandin E synthase (PGES) activities whereas NAD+-dependent 15 hydroxy prostaglandin dehydrogenase (15-PGDH) is the key enzyme implicated in the catabolism of PGE2. For both COX and PGES, three isoforms have been described: in cartilage, COX-1 and cytosolic PGES are constitutively expressed whereas COX-2 and microsomal PGES type 1 (mPGES-1) are inducible in an inflammatory context. COX-3 (a variant of COX-1) and mPGES-2 have been recently cloned but little is known about their expression and regulation in cartilage, as is also the case for 15-PGDH. We investigated the regulation of the genes encoding COX and PGES isoforms during mechanical stress applied to cartilage explants. Mouse cartilage explants were subjected to compression (0.5 Hz, 1 MPa) for 2 to 24 hours. After determination of the amount of PGE2 released in the media (enzyme immunoassay), mRNA and proteins were extracted directly from the cartilage explants and analyzed by real-time RT-PCR and western blotting respectively. Mechanical compression of cartilage explants significantly increased PGE2 production in a time-dependent manner. This was not due to the synthesis of IL-1, since pretreatment with interleukin 1 receptor antagonist (IL1-Ra) did not alter the PGE2 synthesis. Interestingly, COX-2 and mPGES-1 mRNA expression significantly increased after 2 hours, in parallel with protein expression, whereas COX-3 and mPGES-2 mRNA expression was not modified. Moreover, we observed a delayed overexpression of 15-PGDH just before the decline of PGE2 synthesis after 18 hours, suggesting that PGE2 synthesis could be altered by the induction of 15-PGDH expression. We conclude that, along with COX-2, dynamic compression induces mPGES-1 mRNA and protein expression in cartilage explants. Thus, the mechanosensitive mPGES-1 enzyme represents a potential therapeutic target in osteoarthritis.

Introduction

Osteoarthritis (OA) is the leading cause of disability among the elderly population [1]. Traumatic joint injury and joint overload are two major causes of cartilage degradation leading to OA. Although the process of this disease is not yet fully understood, it results from an imbalance in the loss of cartilage caused by matrix degradation and the death of the unique cellular population of cartilage, the chondrocytes. Joints are physiologically exposed to mechanical stress, which triggers gene expression and metabolic activity of chondrocytes in order to turn over the extra cellular matrix and eventually adapt the tis-
sus to loading. The magnitude of the forces that are physiologically applied to cartilage is up to 20 MPa, according to the type of articulation, movement and weight of the individual [2]. Moreover, pressure that is applied on joints comprises a complex combination of strain, shear stress and compressive forces, the latter seemingly being more prevalent in cartilage. The duration of mechanical stress is less than 1 second and leads to cartilage deformation of only 1% to 3% [3]. Many biochemical changes are associated with cartilage degradation and OA progression. These include an increased production of matrix metalloproteinases, proinflammatory cytokines, proinflammatory lipid mediators, extracellular nucleotides, reactive oxygen species and reactive nitrated oxygen species as nitric oxide (NO). It is noteworthy that abnormal cartilage loading may trigger the synthesis of all of these mediators [4-6]. Notably, Fermor and colleagues [6] described that intermittent compression (0.5 Hz, 24 hours, 0.1 to 0.5 MPa) caused an increase in NO production and inducible NO synthase activity (P < 0.05). Different mechanoreceptors have been proven to be at the surface of chondrocytes [7], but the integrin α6β1 could be the major link between extracellular mobilization and intracellular events [8], which eventually promote the synthesis of the various mediators described above. Recent studies have focused on the intracellular events that promote these syntheses under mechanical stress. Among them are the extracellular signal regulated kinases 1/2 (ERK1/2), p38 mitogen-activated protein kinase (p38) and c-jun-N-terminal kinase (JNK) [9], known for their involvement in many biological events.

Prostaglandin E2 (PGE_{2}) is one of the major catabolic mediators involved in cartilage degradation and chondrocyte apoptosis [10-12]. OA cartilage spontaneously releases more PGE_{2} than normal cartilage [13] and in knock-out mice for EP4, a membrane receptor for PGE2, a decreased incidence and severity of cartilage degradation in the collagen-induced arthritis model is observable [14]. Several studies have examined the effects of physical forces on PGE_{2} release. On the one hand, cyclic tensile strain [15] and dynamic compression applied on chondrocytes cultured in agarose for 48 hours [16] inhibited the release of PGE_{2}. On the other hand, fluid-induced shear stress [17] as intermittent mechanical compression for 1 hour increased PGE_{2} release in chondrocytes [6]. So, depending on the type, the magnitude and duration of mechanical stress, different molecular events, such as PGE_{2} release, are triggered in chondrocytes.

PGE_{2} is a prostanoid derived from arachidonic acid released from membranes by phospholipase A_{2}. Arachidonic acid is metabolized by cyclooxygenase (COX) activity to form the prostaglandin endoperoxide H_{2}. Three isoforms of COX (COX-1, COX-2 and COX-3) have been cloned. Whereas COX-1 is constitutively expressed in various cell types to maintain homeostasis, COX-2 is inducible in an inflammatory environment. COX-3 is a recently described derivative of COX-1 that occurs as the result of conservation of the first intron and is also called COX-1 V1. At this time, its expression is described in both canine and human cortex and aorta, and in the rodent heart, kidney and neuronal tissues [18]. Prostaglandin endoperoxide H_{2} is subsequently metabolized by PGE synthase (PGES) to form PGE_{2}. Three types of PGES have been cloned. The cytosolic form (cPGES) is ubiquitous and non-inducible, whereas the microsomal PGES type 1 (mPGES-1) is involved in PGE_{2} synthesis during inflammation. mPGES-1-deficient mice exhibit a significant reduction in disease severity and cartilage degradation in the collagen-induced arthritis model [19,20]. mPGES-1 belongs to the MAPEG family (membrane associated proteins in eicosanoid and glutathion metabolism) and is glutathion dependent. We and others have recently shown that IL-1β upregulates mPGES-1 expression in OA chondrocytes [21,22]. A third form of PGES, called microsomal PGES type 2 (mPGES-2), has recently been cloned. This glutathion-independent enzyme is expressed in various cells and seems to be poorly regulated by inflammation [23]; however, its expression and regulation have not yet been elucidated in cartilage.

Our investigation sought to explore the activation of the arachidonic acid cascade. We hypothesized that mechanical compression in certain conditions would lead to PGE_{2} synthesis by chondrocytes. Furthermore, we wanted to determine whether genes encoding COX and PGES isoforms are mechanosensitive or not.

**Materials and methods**

**Materials**

All of the reagents were purchased from Sigma-Aldrich (St Quentin Fallavier, France), unless stated otherwise. Collagenase D and a Complete protease inhibitor mixture were from Roche Diagnostics (Meylan, France). Antibodies used were: anti-mouse COX-2 polyclonal antibody (Santa Cruz Biotechnology from Tebu, Le Perray-en-Yvelines, France); anti-mouse COX-3 polyclonal antibody (Alpha Diagnostic International, San Antonio, Texas, USA); anti-mouse COX-1 polyclonal antibody; anti-mouse mPGES-1 polyclonal antibody; anti-mouse mPGES-2 polyclonal antibody; anti-mouse cPGES polyclonal antibody (Cayman from SPI-BIO, Massy, France); and anti-mouse β-actin monoclonal antibody. The ECL western-blot analysis system was purchased from Amersham Pharmacia Biotech (Orsay, France). The Immuno-Blot polyvinylidene difluoride (PVDF) membranes for western-blotting and kaleidoscope prestained standards were obtained from Bio-Rad (Ivry-sur-Seine, France). Inteleukin 1 receptor antagonist (IL1-Ra) was obtained from R&D Systems (Minneapolis, MN, USA). Anti-goat fibronectin receptor (integrin α5β1) blocking polyclonal antibody (AB1950) was purchased from Euromedex for Chemicon Inc. (Strasbourg, France) and rat anti-mouse β1 subunit of VLA1 integrins non-blocking monoclonal antibody (VMA 1997), was purchased from AbCys SA for Chemicon Inc. (Souffelweyersheim, France).
Compression experiments
All of the experiments were performed according to the protocols approved by the French/European ethics committee. Compression was applied either on costal cartilage or on articular cartilage. For each experiment, all of the rib cages and all of the knees and the hips were harvested from 6-day-old newborns from one Swiss mouse litter according to the procedure described in [24,25] (Figure 1).

For costal cartilage, explants were cleaned in PBS to eliminate soft tissues and bone sternum parts were discarded. The costal cartilage was cut and divided into segments, which were pooled. Each sample consisted of 50 mg of costal cartilage explants. For articular cartilage, cartilage of two femoral heads and two knees constitute one sample.

Immediately after the dissection, each sample was placed into individual compression wells of Biopress culture plates (Flexercell International, Hillsborough, NC, USA) in 1.5 ml of culture medium (DMEM, containing penicillin-streptomycin 1% v/v, glutamin 2% v/v, albumin 0.1% v/v and Hepes 30 mM) (Figure 1). All of the experiments were performed at 37°C, in air. The compressive stress was applied to individual samples by the Biopress system (Flexercell International) described by Fermor and colleagues [26], whereas the control explants were kept in unloaded conditions. At each time point (2 h, 4 h, 18 h and 24 h), we analyzed compressed and uncompressed explants supplemented or not with effectors. Our results are expressed as fold-induction in comparison to controls. After the application of the mechanical regimen, supernatants and cartilage explants were collected and stored immediately at -20°C and -80°C, respectively.

Intermittent compression was applied using a sinusoidal waveform at 0.5 Hz and 1.0 MPa of magnitude.

Cell viability assay
Immediately after compression, cartilage was first incubated with collagenase D solution (3 mg/ml) for 90 minutes at 37°C,
and then incubated with collagenase D (0.5 mg/ml) overnight at 37°C. The cell suspension obtained was mixed to disperse any cell aggregates, producing a suspension of isolated cells. Cells suspended in a culture medium were colored with Trypan blue (0.04%) and counted in a hemocytometer. This cell viability assay was carried out on one uncompressed and two compressed explants, at 4 hours and 24 hours, and on one explant immediately after the dissection, in two independent experiments.

PGE<br/> and NO assays

Absolute concentrations of nitrite, a stable end-product of NO metabolism, were determined in the media of the cartilage explants using a spectrophotometric method based on the Griess assay (Griess Reagent System, Promega, Charbonnières, France). Absorbance was measured at 550 nm and nitrite concentration was determined by comparison with standard solutions of sodium nitrite.

PGE<sub>2</sub> production was measured in the media by a high sensitivity commercially available enzyme immunoassay kit (Cayman Chemical, Ann Arbor, MI, USA), as previously described [27]. The cross-reactivity of the antibody with other prostanoids is 43% PGE<sub>2</sub>, 37.4% 8-iso PGE<sub>2</sub>, 18.7% PGE<sub>1</sub>, 1% PGF<sub>xα</sub> and 0.25% 8-iso PGF<sub>xα</sub>. The limit of detection was 9 pg/ml. PGE<sub>2</sub> concentration was analyzed at serial dilutions in duplicate and was read against a standard curve.

RNA extraction, reverse transcription and quantitative real-time PCR

Frozen cartilage explants (50 mg) were milled in 600 µL of RLT buffer (from RNeasy Mini Kit, Qiagen GmbH, Hilden, Germany) using a Mixer Mill MM 300 apparatus (Qiagen). Disruption was achieved through the beating and grinding effect of beads on the cartilage samples as they were shaken together in the grinding vessels. One steel ball (diameter 5 mm) was added to each sample and they were mixed, at a cool temperature, for two cycles of 2 minutes at 25 pulses/second. Then, after removing the beads, the total RNA was extracted from each sample using the RNeasy Kit (Qiagen) according to the manufacturer's instructions. A proteinase K (Qiagen) digestion step was performed after the lysis of cartilage explants and a DNAse digestion step (RNAse free DNAse set, Qiagen) was added. RNA concentration was then measured using a spectrophotometer. The migration in an agarose gel enabled quality control.

Total RNA (1 µg) was reverse transcribed with Omniscript (Qiagen) in a final volume of 20 µL containing 50 ng of oligos dT. The enzyme was then inactivated by heating and the interesting mRNAs (COX-1, genbank BC005573; COX-2, NM_011198; COX-3, AY547265; mPGES-1, NM_022415; mPGES-2, BC004846; cPGES, NM_008278; 15-PGDH, NM_008278) were quantified by real-time quantitative reverse transcription RT-PCR using the iCycler iQ Real Time PCR (Bio-Rad) and QuantiTect SYBR PCR kits (Qiagen). Sense and antisense PCR primers were designed based on mouse sequence information for the amplification of genes of interest (Table 1). The PCR reactions were performed in a 25 µl final volume using 0.06 to 0.25 µl of cDNA, 600 ng of specific primers and 1× QuantiTect SYBR Green PCR master mixture, including HotStar Taq DNA Polymerase, QuantiTect SYBR Green PCR buffer, SYBR Green I, and ROX in which there was 5 mM MgCl<sub>2</sub>. PCR amplification conditions were: initial denaturation for 13 minutes at 95°C followed by 50 cycles consisting of 30 seconds at 95°C and 30 seconds at 58°C. Product formation was detected at 72°C in the fluorescein isothiocyanate channel. The generation of specific PCR products was confirmed by melting-curve analysis. For each real-time RT-PCR run, cDNA were run in quadruplicate in parallel with serial dilutions of a cDNA mixture tested for each primer pair to generate a linear standard curve, which was used to estimate relative quantities of COX, PGES and 15-PGDH mRNA.

| Genes       | Temperature (°C) | Genbank ID  | Forward primer                  | Reverse primer                  | Amplicon length (bp) |
|-------------|------------------|-------------|---------------------------------|---------------------------------|----------------------|
| HPRT        | 58               | NM_013556   | 5'-gtggtggtaaagacacc-tct-3'     | 5'-cacaggactagaacacctgc-3'      | 249                  |
| COX-1       | 58               | BC005573    | 5'-ctttgcacaaacactccac-3'       | 5'-agcaacccaaacaccttg-3'        | 285                  |
| COX-2       | 58               | NM_011198   | 5'-gattctttgccagctct-3'         | 5'-agaccaggcagcaaaaga-3'        | 299                  |
| COX-3       | 58               | AY547265    | 5'-tgamgtcagctgtcact-3'         | 5'-ggttctggcacggatagtaac-3'     | 349                  |
| cPGES       | 58               | AY281130    | 5'-agctgtgctcagttcactaacc-3'    | 5'-tggtaatgcatcagttccc-3'       | 196                  |
| mPGES-1     | 58               | NM_022415   | 5'-cttctgtaaatcagttacgc-3'      | 5'-cccaggtagccagctg-3'          | 294                  |
| mPGES-2     | 58               | BC004846    | 5'-aagatgtcctcctcctgc-3'        | 5'-cccaagtgggacacttcc-3'        | 133                  |
| 15-PGDH     | 58               | NM_008278   | 5'-gctggtgcatcaagatgtacg-3'     | 5'-cttccgaagttcatgtcact-3'      | 164                  |

15-PGDH, NAD+-dependent 15 hydroxy prostaglandin dehydrogenase; COX, cyclooxygenase; HPRT, hypoxanthine-guanine phosphoribosyltransferase; cPGES, cytosolic PGES; mPGES, microsomal PGES; PGES, prostaglandin E synthase.
normalized for Hypoxanthine-guanine phosphoribosyltransferase (HPRT genbank NM_008278) in the samples.

**Protein extraction and western blotting**

Frozen cartilage explants were disrupted using a Mixer Mill MM 300 apparatus (Qiagen) in 500 µL of cold lysis buffer (20 mM Tris pH 7.6, 120 mM NaCl, 10 mM EDTA, 10% glycerol, 1% Nonidet P-40, 100 mM NaF; 10 mM Na2HPO4, 1 mM AEBSF (4-(2-Aminoethyl)benzenesulphonyl fluoride), 2 mM Na2VO3, 40 µg/ml leupeptin, 1 µM pepstatin A, 10 µg/ml aprotinin). One steel ball (diameter 5 mm) was added to each sample, which were mixed at a cool temperature for two cycles of 2 minutes at 25 pulses/second. Then, after removing the beads, the samples were shaken gently for 1 hour at 4°C and then centrifuged for 1 hour (13,000 g, 4°C). The supernatants were collected and protein concentrations were determined using the bichinchoninic acid assay kit (Perbio Science for Pierce, Bezons, France).

Cartilage explant lysates were separated by 8% or 15% SDS-PAGE and transferred to nitrocellulose membranes. The blots were incubated (then stripped and reprobed) by the appropriate primary polyclonal antibody to COX-2, COX-3, COX-1, mPGES-1, mPGES-2, cPGES and monoclonal antibody to β-actin. The blots were then incubated with horseradish peroxidase-conjugated secondary goat antibody. The membranes were washed repeatedly with Tris-buffered Saline containing Tween-20 0.1% (v/v) and the signals were detected using the enhanced chemiluminescence detection system and exposed to Kodak BioMax MR-1 film. We transfected Cos cells with plasmids encoding COX-2 and mPGES-1. Cells extracts containing COX-2 and mPGES-1 proteins were used as positive controls.

**Immunohistochemistry**

After compression for 18 hours, cartilage explants were immediately collected and fixed in 70% ethanol at 4°C for 48 hours. After dehydratation, the cartilage samples were embedded without demineralization in methyl methacrylate (Merck, Darmstadt, Germany). Transversal sections (4 µm thick) were cut parallel to the rib axis using a Polycut E microtome (Leica, Wetzlar, Germany). Sections mounted onto slides were deplastified in 2-methoxyethylacetate prior to further processing. The primary polyclonal antibodies used were the same as those used for western blotting, as previously described. For immunochemistry, the sections were incubated overnight with 0.1 M PBS supplemented with 0.05% Tween 20 (Sigma) and 1% BSA (Euromedex) and the primary polyclonal antibody (1:50) at room temperature. They were then treated with 3% hydrogen peroxide (10 minutes), and an avidin-biotin peroxidase complex (ABC Vectastain kit, Vector) for 60 minutes. PBS (0.1 M) was used for the washing steps between incubations. Diaminobenzidine tetrahydrochloride (Sigma) was used as the chromogen. The sections were lightly counterstained with toluidine blue (pH 3.8). Negative controls were prepared by omitting the primary antibody in the diluant solution (BSA 10% and milk 1% for COX-2). Immunohistological analysis was carried out on two uncompressed and two compressed costal cartilage explants. Images were obtained using an optical microscope and analysis for each enzyme utilized a blind test.

**Statistical analysis**

All data are reported as mean ± SEM, unless stated otherwise. Unpaired Students’ t-tests were used to compare the mean values between groups with the GraphPad InStat version.
to compression exhibit an increase in PGE2 release that was sustained up to 4 hours (6-fold increase; p < 0.001) and then decreased at 24 hours (Figure 2a), as described by Fermor and colleagues [6]. Also, PGE2 synthesis in the media significantly increased, with a peak at 2 hours that was sustained up to 4 hours (6-fold increase; p < 0.001) and then decreased at 24 hours (Figure 2b).

To determine the effects of compressive stress on chondrocyte activation, we assessed NO and PGE2 release in the media of compressed and uncompressed costal cartilage explants. Different magnitudes and lengths of stress were applied in order to define the optimum conditions (data not shown). At a sinusoidal waveform frequency of 0.5 Hz and a magnitude of 1 MPa, NO release significantly increased (6-fold increase; p < 0.001) within 2 hours compared to uncompressed explants, and lasted 24 hours (Figure 2a), as described by Fermor and colleagues [6]. Also, PGE2 synthesis in the media significantly increased, with a peak at 2 hours that was sustained up to 4 hours (6-fold increase; p < 0.001) and then decreased at 24 hours (Figure 2b).

Compressive stress was also applied to mouse articular cartilage explants in order to avoid a bias due to the origin of cartilage. As in costal cartilage, articular cartilage explants submitted to compression exhibit an increase in PGE2 release after 2 hours (16-fold increase; p < 0.01), which was sustained from 4 hours to 18 hours (6-fold increase) before declining to control levels. Even though PGE2 release in articular cartilage (16-fold increase, p < 0.01) was stronger at 2 hours than in costal cartilage (6-fold increase, p < 0.001) and was sustained for longer, only minor differences in kinetics were observed (Figure 3).

Viability of the chondrocytes in the mouse costal cartilage explants was tested using Blue Trypan coloration. No alteration of cell viability was seen between compressed and uncompressed samples within 24 hours (data not shown).

To confirm the validity of our compressive model on mouse costal cartilage, we wanted to highlight the implication of integrin α5β1 in the PGE2 release triggered by mechanical stress. Cartilage explants treated with anti-integrin α5β1 blocking antibody (AB1950) at 25 μg/ml induced a 50% decrease of compression-induced NO (4.16 ± 0.57 versus 2.68 ± 0.43 μM; data obtained from 2 independent experiments with n = 2/group/experiments; p < 0.001, data not shown). Moreover, a decrease in PGE2 release of approximately 50% in compressed cartilage treated with the blocking antibody was observed (Figure 4a). No modification in NO (4.01 ± 0.1 versus 4.4 ± 1.43 μM; data obtained from 2 independent experiments with n = 2/group/experiments, data not shown) or PGE2 release (Figure 4a) was detected in media of compressed cartilage treated with the non-blocking anti-β1 subunit antibody at 2.5 μg/ml (VMA1997).

We previously reported that the pro-inflammatory cytokine IL-1 triggers the expression of COX-2 and mPGES-1 [21,28]. Since the integrin antibody did not fully inhibit a compression-induced PGE2 release, even if other mechanoreceptors have been described on chondrocytes, we hypothesized that compression could indirectly act on cartilage by inducing the synthesis of IL-1. When the IL-1 receptor antagonist IL1-Ra was added at a concentration of 100 ng/ml prior to compression, no variation in PGE2 release was observed (Figure 4b), suggesting that compression-induced PGE2 release is not mediated by IL-1.

Expression of COX and PGES enzymes in uncompressed and compressed cartilage explants

We subsequently focused our study on the enzymes involved in PGE2 synthesis, cyclooxygenases and prostaglandin E synthases. Mouse costal cartilage explants subjected to compressive stress for 18 hours were fixed in ethanol, embedded in methyl methacrylate and cut into serial sections that were immunostained with antibody against COX-2, mPGES-1, mPGES-2 and cPGES. Toluidine blue counterstaining colors the extracellular matrix and nuclei of cells. In uncompressed cartilage, a few peripheral cells presented positive immunostaining (brown) for COX-2 and none did so for mPGES-1. After compression, an increased brown staining for COX-2 and mPGES-1 in cells was visible around the nuclei, suggesting a colocalization of these enzymes in the perinuclear region in loaded chondrocytes. For cPGES and mPGES-2, no differences appeared in chondrocytes from compressed cartilage explants compared to uncompressed explants (Figure 5).

COX expression in cartilage explants subjected to compression

To study the effects of mechanical loading on COX gene expression, we used real-time RT-PCR quantitative analysis...
and immunoblotting to evaluate, respectively, the transcrip-
tional and translational expression of COX-1, COX-2 and 
COX-3 genes. We extracted the total RNA and proteins 
directly from costal cartilage explants. An increased expres-
sion of COX-2 mRNA, but not COX-1, was observed after 2 
hours with a maximal effect after 4 hours of compression (Fig-
ure 6a,b). Interestingly, COX-3 mRNA was expressed in carti-
lage but compressive stress had no effect on its transcriptional 
expression in cartilage explants (Figure 6c).

We then assessed COX protein levels by immunoblotting 
using polyclonal antibodies raised against COX-1, COX-2 and 
COX-3. As expected, compression induced COX-2 protein 
expression after 4 hours and peaked at 18 hours, whereas 
COX-1 expression remained unchanged. Both the COX-3 
protein and its mRNA were expressed in cartilage; however, 
compression did not modify its expression (Figure 6d).
Expression of mPGES-1 but not cPGES mRNA increased after 2 hours of compression with a peak at 4 hours. Similarly, the amount of mPGES-1 protein increased in compressed explants after 2 hours, peaking at 18 hours (Figure 7a,b,d). The increased expression over the time of mPGES-1 protein in uncompressed samples, which was also observed for COX-2, could be triggered by mediators release during explantation and cutting of cartilage.

Interestingly, mPGES-2 was not regulated by compressive stress, both at the mRNA and protein levels (Figure 7c,d).

The gene encoding 15-prostaglandin dehydrogenase is mechanosensitive
Because our results highlighted a discrepancy between the kinetics of PGE2 release and COX-2 and mPGES-1 expression (compare Figure 2b with Figures 6 and 7), we hypothesized that the decrease in PGE2 production observed after 4 hours was due, at least in part, to the activation of a catabolic pathway of PGE2.

NAD+-dependent 15 hydroxy prostaglandin dehydrogenase (15-PGDH) is considered to be the key enzyme in the catabolism of PGE2. Interestingly, the gene encoding 15-PGDH is mechanosensitive and the kinetics of its expression is in agreement with our hypothesis since a peak of expression is observed at 4 hours (Figure 8).

Discussion
Our findings demonstrate that dynamic mechanical loading of costal cartilage can significantly increase PGE2 release. Moreover, we describe here for the first time that COX-2 and mPGES-1 expression is increased in mouse costal cartilage explants under compression but not their constitutive isoforms COX-1 and cPGES. Therefore, it appears that COX-2 and mPGES-1 are encoded by mechanosensitive genes implicated in the compression-induced PGE2 release. PGE2 is the pivotal eicosanoid involved in the initiation and the development of inflammatory disease, such as rheumatoid arthritis [29]. Notably, it is thought to be a key regulator of cartilage degradation during OA [30]. An increase in PGE2 release induced by mechanical stress has already been described in various tissues [31,32] and particularly in articular cartilage subjected to dynamic compression representative of the physiological range [6].

Regulation of COX-2 mRNA expression in cartilage by mechanical stress has already been reported in the literature [6]. Notably, elements including AP-1 sites, cyclic AMP response elements (CREs) and shear stress response elements (SSRE) are found in the promoter region of mechanical stress-response genes, such as those encoding COX-2 and inducible NO synthase. Shear stress response elements contain a TPA response element to which NFκB, which is part of a main mechanical pathway, binds [33]. Ogasawara and colleagues [34] have described the role of C/EBP beta, AP-1 sites and CREB in shear stress-induced COX-2 expression in osteoblasts. Moreover, post-transcriptional regulation by mRNA stabilization seems to be involved in COX-2 gene expression in vascular endothelial cells subjected to fluid...
shear stress [35]. In addition to these studies at the mRNA level, we show here, for the first time in cartilage, that COX-2 is also increased at the protein level.

Interestingly, our data indicate that COX-3, also named COX-1 V1, is expressed in mouse costal cartilage. COX-3, which was cloned in 2002, was derived from COX-1 through retention of intron 1 in its mRNA. This probably resulted in the modification of the active site conformation of the enzyme. COX-3 expression has actually been found in several canine, human and rodent tissues, but never in cartilage, whatever the species [18]. In this present study, we report for the first time the expression of COX-3 (mRNA and protein) in mouse cartilage. Moreover, we show that mechanical loading did not modify its expression. As COX-1 and COX-3 are derived from the same gene, these enzymes share the same promoter. However, no sites that are regulated through mechanical stress or pro-inflammatory cytokines have been found so far in the COX-1 promoter, which is consistent with the fact that COX-1 is constitutively and ubiquitously expressed. Thus, this might explain the lack of COX-3 regulation by compression.

The regulation of mPGES-2 expression has never been described in cartilage. mPGES-2 is ubiquitously expressed...
under basal conditions in many tissues and is activated by reducing agents, but its role in PGE2 release in both basal and inflammatory contexts remains unclear. In human rheumatoid synoviocytes, expression of mPGES-1 increased with severity of the disease, whereas that of mPGES-2 did not [23]. In COX-2-deficient mouse brains, a decreased release of PGE2 and a decreased expression of mPGES-2, but not of mPGES-1 or cPGES, was observed, suggesting that mPGES-2 could be functionally coupled to COX-2 [36]. In the present study, mPGES-2 expression was similar in compressed and uncompressed cartilage explants, suggesting that mPGES-2 is not encoded by a mechanosensitive gene.

The striking point of our study is the evidence that mPGES-1 is encoded by a mechanosensitive gene. In recent years, several studies have demonstrated that inflammation induces mPGES-1. In rat paws of the acute and chronic arthritis model, up-regulation of mPGES-1 mRNA and protein expression was observed. Moreover, levels of mPGES-1 mRNA and protein were markedly elevated in OA versus normal cartilage [37]. Additionally, we and others have previously reported an over-expression of mPGES-1 in OA chondrocytes in primary cultures stimulated by IL-1 [21,22]. Interestingly, our results identify an earlier significant transcriptional expression (as soon as 2 hours) after mechanical stress compared to the effect of IL-1 (after 12 hours). Moreover, its induction was higher with compression (five-fold) compared to IL-1 stimulation (three-fold). A structural comparison of COX-2 and mPGES-1 promoters revealed that the gene encoding mPGES-1 does not contain transcriptional elements that are
classically involved in response to mechanical stress (AP-1, CRE, SSRE), as described for the COX-2 promoter [38]. It suggests that divergent transcriptional mechanisms are responsible for inducible mPGES-1 regulation. Yokota and colleagues [39] have described the role of the nuclear regulator CITED2 (CBP/p300-interacting transactivator with ED-rich tail 2) in shear-induced down-regulation of matrix metalloproteinase 1 and matrix metalloproteinase 13. This mechanism could also occur in compression-induced mPGES-1 expression. But post-transcriptional regulation could also be present since a stabilization of mPGES-1 mRNA has been recently described in cardiomyocytes, leading to delayed protein expression via cJNK [40].

Our data suggest that loading triggers mPGES-1 and COX-2 colocalization in the perinuclear region of the chondrocytes. This result is in line with the study from Kojima and colleagues [22] that recently described COX-2 and mPGES-1 colocalization around the nuclei of chondrocytes after IL-1 stimulation.

Selective inhibitors of mPGES-1 have been developed recently [41]. Effectively, targeting the mPGES-1 enzyme in preference to COX-2 should represent a new therapeutic approach to treating joint diseases such as OA. Interestingly, Cheng and colleagues [42] reported that mPGES-1-null mice exhibit no alteration in thrombogenesis or blood pressure, whereas selective COX-2 inhibitors (Coxib) did. These results suggest that inhibitors of mPGES-1 may retain their anti-inflammatory efficacy by depressing PGE2, while avoiding the adverse cardiovascular consequences associated with COX-2 deletion [42].

To understand the weak PGE2 release at 18 and 24 hours after compression, we focused on 15-PGDH mRNA expression because 15-PGDH is one of the major PGE2-inactivating enzymes [43]. 15-PGDH expression decreases in response to IL-1β treatment in trophoblast cells in primary culture [44] as well as in response to lipopolysaccharide (LPS) treatment in liver and lungs of rats [45]. Moreover, 15-PGDH expression is reduced in inflammatory bowel disease [46]. To our knowledge, 15-PGDH expression in cartilage has never been reported and its expression under mechanical stress was unknown. The findings presented in this study first indicate that 15-PGDH is expressed in mouse cartilage and is induced after 4 hours of compression (three-fold induction). Genomic structural analysis revealed that the human 15-PGDH promoter contains binding sites for AP-1 and CREB, two transcriptional factors implicated in the regulation of gene expression subjected to mechanical stress [47]. We suggest that the 15-PGDH enzyme could be implicated in the catabolism of PGE2 after 18 hours. So, proinflammatory stimuli (LPS or cytokines) induce monophasic PGE2 synthesis whereas mechanical stress triggers a biphasic response in cartilage, PGE2 synthesis followed by PGE2 degradation. We hypothesize that this response corresponds to an adaptation of the tissue to this stress.

The choice of an experimental model to study the physiological regulation of cartilage homeostasis by mechanical stress is challenging. Three types of mechanical stress have been described: shear stress, strain and compressive stress. Overloading of the joints, a major risk factor for OA, mainly induces compressive stress. However, shear stress and strain have also been described when overloading is experimentally applied on cartilage. The compression of chondrocytes embedded in agarose or collagen gels as well as the compression of ex vivo porcine cartilage explants represent the main strategies developed [6,16,26,48]. Among the literature, it appears obvious that the consequences to metabolism of chondrocytes vary according to the type of stress applied. Takahashi and colleagues [49] reported that static compression promotes type II collagen, whereas cyclic loading could denature type II collagen in articular chondrocytes [50]. Here, we used mouse cartilage explants subjected to cyclical uniaxial mechanical stress. Mouse cartilage was chosen because of the availability of a large choice of biomolecular tools suitable to study it, such as genetically modified mice. However, many labs work on discs of porcine or bovine cartilage, permitting a more homogeneous compressive stress. In our model, the compressive stress is less uniform because of the size of the explants. Enhancements to our system, in order to better define the stress applied, is ongoing.

We decided to work on costal cartilage rather than articular cartilage because of the extremely small quantity of the latter available. However, we performed comparative experiments in order to validate our model with costal cartilage explants. We observed an induction of PGE2 release in both types of cartilage, but with slight differences in kinetics.
The mechanoreceptor integrin α5β1 is considered by many authors to be the critical mechanoreceptor for transducing signal from the extracellular matrix to the chondrocyte [8]. Our data confirm the major role of this receptor in compression-induced PGE₂ release.

Among the mechanotransduction pathways recently described, Mitogen-Activated Protein Kinase (MAPK) plays a major role. In smooth muscle cells and fibroblasts, mechanical strain increases the activity of all three MAPKs, namely p38 MAPK, ERK 1/2 and Junk kinase [51,52]. In particular, ex vivo cartilage compression has been reported to activate the three MAPKs, but with different kinetics [9]. Preliminary results suggest that this is also the case in our model (personal communication).

Surprisingly, overexpression of COX-2 and mPGES-1 proteins was observed at 18 hours in uncompressed samples. Several authors have described the release of mediators, such as basic fibroblast growth factor (bFGF) and IL-1α, in cartilage after explantation and cutting [53,54]. Moreover, Fermor and colleagues [6] reported that uncompressed explants of articular cartilage exhibited a significant production of PGE₂ (from 14 pg/mg/ml at 24 hours after explantation to 1 pg/mg/ml at 72 hours). We have considered that dissection alone induces COX-2 and mPGES-1 protein expression in the uncompressed samples.

As mature articular cartilage is an avascular tissue, the oxygen supply to resident chondrocytes could rather limit and influence chondrocyte activation under a mechanical signal. Fermor and colleagues have shown that oxygen tension influences the endogenous production of NO and PGE₂ in porcine cartilage explants in response to mechanical stimulation. Under mechanical compression, PGE₂ production in cartilage at 20% O₂ increased 50-fold, but in cartilage at 5% O₂ it increased only 4-fold and in cartilage at 1% O₂ it did not increase at all [48]. Since previous studies have suggested that the oxygen tension in the superficial layer of articular cartilage is higher than in the deeper layer [55], our results could have occurred, at least in part, in the superficial zone of articular cartilage.

Conclusion
We have demonstrated that COX-2 and mPGES-1 are mechanosensitive enzymes after dynamic compression. Moreover, the key enzyme implicated in the catabolism of PGE₂, 15-PGDH, is expressed in cartilage and its expression is regulated by compression. As mPGES-1 is currently described as a potential therapeutic target for controlling PGE₂ release in OA, further research into the regulation of this mechanosensitive gene should be helpful for a better understanding of the key pathways implicated in its expression in order to counteract cartilage degradation in overloading joints.

Competing interests
The authors declare that they have no competing interests.

Authors' contributions
MG performed animal surgery, compression experiments, PGE₂ and NO assays, PCR experiments, western blotting experiments and immunohistochemistry experiments, analyzed the results and drafted the manuscript. FB was involved in the conception and design of the study and critically revised the manuscript for important intellectual content. AL performed animal surgery, compression experiments, PGE₂ and NO assays and western blotting experiments. AP performed animal surgery, compression experiments and PGE₂ and NO assays. ST initiated and participated in compression experiments. JLS performed immunohistochemistry experiments. CJ designed the study, participated in data analysis and helped draft the manuscript. All authors read and approved the final manuscript.

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