Corticosterone suppresses IL-1β-induced mPGE2 expression through regulation of the 11β-HSD1 bioactivity of synovial fibroblasts in vitro

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Abstract. The aim of the present study was to investigate the correlation between glucocorticoid activity regulation, prostaglandin E2 (PGE2) synthesis, and synovial inflammation inhibition activity, through microsomal prostaglandin E synthase-1 (mPGES-1) expression regulated by the glucocorticoid pre-receptor regulator, 11β-hydroxysteroid dehydrogenase-1 (11β-HSD1). In the present study, fibroblast-like synovial cells of rats were studied as a cell model. Cells were stimulated with 10 ng/ml interleukin (IL)-1β for 24 h, and were subsequently, within the next 24 h, treated with or without 10⁻⁴ mmol/l corticosterone alone or with 100 nmol/l PF915275. At the end of the second 24 h, PGE2 levels in culture supernatants were assayed. Cells were harvested for mRNA evaluation of 11β-HSD1, mPGES-1, IL-1β and tumor necrosis factor (TNF)-α, and protein detection of 11β-HSD1 and mPGES-1 using reverse transcription-qualitative polymerase chain reaction and western blot analysis, respectively. Corticosterone was demonstrated to suppress the mRNA expression levels of inflammatory factors, such as TNF-α and PGE2, induced by IL-1β in vitro. Simultaneously, expression levels of 11β-HSD1 decreased significantly at the mRNA and protein levels (P<0.05). Cortisol concentration in the medium of the group treated with corticosterone was significantly increased (P<0.05) compared with that of the control group; however, the cortisol concentration was decreased in the medium when the conversion bioactivity of 11β-HSD1 was inhibited by PF915275, while the changes in 11β-HSD1 and mPGES-1 mRNA expression levels and PGE2 content were reversed in the medium. These results indicated that a significant positive correlation (P<0.01) may exist between mRNA and protein expression levels. To conclude, 11β-HSD1 is a key regulator for the synthesis of mPGES-1 and PGE2 in the inflammatory synovial cells in vitro, suggesting a potential interference target for osteoarthritis.

Introduction

Osteoarthritis (OA) was once considered as a non-inflammatory form of arthritis, but studies have demonstrated that synovitis is associated with major symptoms of OA, such as pain and the degree of joint dysfunction, and may promote more rapid cartilage degeneration (1-3). Synovial inflammation is an important factor involved in the acceleration of cartilage degeneration (4,5). It is likely that multiple joint tissues contribute to joint inflammation (6). Fibroblast-like cells (FLS) are the main functional synovial cells that produce cytokines, such as tumor necrosis factor (TNF)-α, interleukin (IL)-1β, IL-1β and IL-6, which are involved in the degradation of cartilage (7). FLS can also affect the natural course of arthritis by releasing prostaglandin E2 (PGE2) (8).

It is now known that several cell types are able to generate active glucocorticoids within their cytoplasm through expression of the 11β-hydroxysteroid dehydrogenase type 1 (11β-HSD1) enzyme. The generation of active glucocorticoids in the synovium is strongly linked to the level of inflammation. Hardy et al (9) reported that 11β-HSD1 mRNA was highly expressed in synovial tissues affected by OA, and its activity was increased, as indicated by IL-1β and TNF-α. Synovial fibroblasts have been demonstrated to maintain the balance between intracellular glucocorticoid activation and inactivation, and execute biological effects by producing 11β-HSD1 and binding with its receptor (9,10). It has been reported that, in fetal membranes and adipose tissues, 11β-HSD1 mRNA expression and protein levels are upregulated by pro-inflammatory cytokines (11,12). Sun and Myatt reported a coordinated induction effect existed for the regulation of 11β-HSD1 by glucocorticoids and pro-inflammatory cytokines (11). Glucocorticoids usually play an opposing role to proinflammatory cytokines at sites of inflammation (13,14). However, how the

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glucocorticoid and pro-inflammatory mediators induce their effects on 11β-HSD1, or whether 11β-HSD1 correlates with PGE₂ expression in the synovial fibroblasts, remains unclear.

Therefore, we hypothesized that glucocorticoid activity correlated with PGE₂ synthesis, and that the glucocorticoid pre-receptor regulator, 11β-HSD1, may have an effect on relieving synovial inflammation by inhibiting microsomal prostaglandin E synthase-1 (mPGES-1) and PGE₂ expression. In the present study, a model cell, fibroblast-like synovial cell, derived from rats, was stimulated with IL-1β and the effect of treatment with corticosterone and 4'-cyano-biphenyl-4-sulfonic acid (6-amino-pyridin-2-yl)-amide was evaluated. PGE₂ levels in culture supernatants were assayed, the mRNA expression of 11β-HSD1, mPGES-1, IL-1β and TNF-α by the cells was analyzed and reverse transcription-quantitative polymerase chain reaction and western blot analysis were used to detect protein expression of 11β-HSD1 and mPGES-1. The anti-inflammatory mechanism of glucocorticoid in suppressing IL-β induced mPGES-1 expression through regulation of 11β-HSD1 bioactivity in synovial fibroblasts in vitro was explored.

Materials and methods

Isolation and culture of Sprague-Dawley (SD) rat synoviocytes. Synovial fibroblasts were isolated from the knee synovial tissue of 10 female healthy 3-month SD rats (200±30 g; Laboratory Animal Centre, Guangdong Medical College, Zhanjiang, China). Rats were kept under regular conditions with a temperature of 25°C, humidity of 50% and a natural day and night cycle. All rats were able to access food and water freely. Rats were euthanatized with 4% formaldehyde for 30 min. Then, the cells were washed in PBS for 5 min, incubated in 3% H₂O₂, and in 0.3% Triton X-100 solution for 30 min, respectively, and then blocked with BSA at room temperature for 10 min. The cells were incubated with primary antibody vimentin mouse anti-rat antibody (1:300 dilution; cat. no. BM0135; Wuhan Boster Biological Technology, Ltd., Wuhan, China). Instead of primary antibody, PBS was added as a negative control. The cells were incubated at 4°C overnight. The next day, the slices were washed in PBS for 5 min, 3 times, and then incubated with secondary antibody (polymeric HRP-Conjugated Anti-Goat IgG Super Vision Assay kit; cat. no. SV0003-1; Wuhan Boster Biological Technology, Ltd.) at room temperature for 30 min. DAB chromogen was used according to the manufacturer's guidance (Wuhan Boster Biological Technology, Ltd.).

Study design. Synovial fibroblasts, preserved in DMEM, were removed of glucocorticoid using 10% active carbon and subsequently stimulated with 10 ng/ml IL-1β (R&D Systems, Inc., Minneapolis, MN, USA) for 24 h. Synovial fibroblast cells were washed with PBS at 37°C for 5 min, three times. After 24 h, cells were treated with different components depending on the allocated group: Group A, treated with DMEM without glucocorticoid; group B, treated with 10 ng/ml IL-1β; group C, treated with 10 ng/ml IL-1β and 10⁻⁶ mmol/l corticosterone (Sigma-Aldrich); and group D, treated with 10 ng/ml IL-1β, 10⁻⁶ mmol/l corticosterone and 100 nmol/l PF915275 (Tocris Bioscience, Bristol, UK). Following a further 24 h, PGE₂ levels were assayed in culture supernatants by ELISA. Cells were harvested for mRNA evaluation of 11β-HSD1, mPGES-1, IL-1β and TNF-α levels; and protein expression level detection of 11β-HSD1 and mPGES-1.

RNA extraction and reverse transcription-quantitative real-time polymerase chain reaction (RT-qPCR) analysis. Total RNA was isolated from synovial fibroblast layers using TRIzol reagent (Takara Biotechnology Co., Ltd., Dalian, China) following the manufacturer's protocol. RNA quality was detected by agarose gel electrophoresis. Reverse transcription was carried out using the PrimeScript RT Reagent kit (Takara Biotechnology Co., Ltd.). The mRNA expression of IL-1β, TNF-α, 11β-HSD1 and mPGES-1 was determined using the SYBR Premix Ex Taq™ kit (Takara Biotechnology Co., Ltd.). The volume of the reaction system was 15 µl, and the reaction system contained 7.5 µl SYBR Premix Ex Taq II, 0.6 µl each upstream and downstream primer, 1.5 µl DNA template and 4.8 µl sterilized distilled water. Target mRNA levels were normalized to β-actin expression levels. Primers

Vimentin immunocytochemical staining. The cells were fixed with 4% formaldehyde for 30 min. Then, the cells were washed in PBS for 5 min, incubated in 3% H₂O₂, and in 0.3% Triton X-100 solution for 30 min, respectively, and then blocked with BSA at room temperature for 10 min. The cells were incubated with primary antibody vimentin mouse anti-rat antibody (1:300 dilution; cat. no. BM0135; Wuhan Boster Biological Technology, Ltd., Wuhan, China). Instead of primary antibody, PBS was added as a negative control. The cells were incubated at 4°C overnight. The next day, the slices were washed in PBS for 5 min, 3 times, and then incubated with secondary antibody (polymeric HRP-Conjugated Anti-Goat IgG Super Vision Assay kit; cat. no. SV0003-1; Wuhan Boster Biological Technology, Ltd.) at room temperature for 30 min. DAB chromogen was used according to the manufacturer's guidance (Wuhan Boster Biological Technology, Ltd.).
used in the experiment are listed below. The following cycling conditions were used: 95°C for 2 min, followed by 40 cycles of 95°C for 15 sec and 60°C for 30 sec. Three replications of this experiment were performed for the same reaction. A Real-time Quantitative PCR instrument (Bio-Rad Laboratories, Inc., Hercules, CA, USA) and the 2−ΔΔCt method (15) with LERTPA-V1.0 software (http://www.biostatistic.net/portal.php) was used to analyze the relative gene expression data: Experimental group (Cq target gene-Cq housekeeping gene)-blank group (Cq target gene- Cq housekeeping gene). The Cq value in this formula represented the number of amplification cycles required for a fluorescent signal in each well to reach a specific threshold. The Cq value was automatically calculated by the computer based on the amplification curve. Gene-specific primer pairs used were as follows: IL-1β, forward 5'-CTTCAATCTCACGCAGCAGC-3' and reverse 5'-GCTGTCATATGGGAACATACA-3'; TNF-α, forward 5'-GTGCCACTAGCTCTTCTCATT-3' and reverse 5'-CTCTGCTTGGTGTGTTGCTAC-3'; 11β-HSD1, forward 5'-AAA ATGACCCAGCTATGATTG-3' and reverse 5'-GGACACAGAGAGTGATGGACAC-3'; mPGES-1, forward 5'-GTGATTGGACACCCAGGT-3' and reverse 5'-CAAGAGGAGGAAGGGTAGATG-3'; and β-actin, forward 5'-CCCATCT ATGAGGGTTACGC-3' and reverse 5'-TTTAATGTGACACGAGATTCC-3'.

Assays for PGE$_2$ and cortisol. Levels of PGE$_2$ in culture supernatants were determined using an ELISA kit (cat. no. KGE004B; R&D Systems, Inc.) in accordance with the manufacturer's instructions. Assays were based on the combined use of a monoclonal antibody against PGE$_2$ and an alkaline phosphatase-conjugated polyclonal antibody; 5-nitrophenyl phosphate substrate was added, and the absorbance at 405 nm was analyzed using a micro Multiskan plate reader. The limits of detection were 10 and 1.4 pg/ml for PGE$_2$. Positive controls were used in each experiment.

Cortisol levels were also determined in culture supernatants using an ELISA kit (cat. no. KGE008B; R&D Systems, Inc.), in accordance with manufacturer's instructions to indicate the conversion rate of cortisol. The assay required microtiter plates coated with purified antibody and the standards or samples, anti-cortisol antibody and horseradish peroxidase-labeled avidin were subsequently added. Absorbance (optical density) was read using a Multiskan microplate reader at 450 nm, following a substrate 3,3',5,5'-tetramethylbenzidine color reaction. Sample concentrations were calculated using a standard curve.

**Western blot analysis.** Synovial fibroblasts, seeded in 6-well plates and grown to 80% confluency, were washed twice with ice-cold PBS and scraped off the wells in TRIzol lysates, containing 1% protease inhibitors. Cells were collected in 1.5 ml Eppendorf tubes and centrifuged 19,419 at x g for 10 min at 4°C, prior to determination of protein concentration, using a Bradford-based assay (Beyotime institute of Biotechnology, Shanghai, China). Protein samples (100 µg) were separated by 30% SDS-PAGE, and electroblotted on a polyvinylchloride membrane. Following 1-h incubation in blocking buffer [Tris-buffered saline-Tween (TBS-T) with 5% non-fat dried milk], membranes (Millipore Saint-Quentin-en-Yvelines, France) were incubated overnight at 4°C with β-actin (1:2,000; cat. no. 4967; Cell Signaling Technology, Inc., Danvers, MA, USA), 11β-HSD1 (cat. no. sc-20175; Santa Cruz Biotechnology, Dallas, TX, USA) and mPGES-1 (both 1:500; cat. no. 13035; Cell Signaling Technology, Inc) primary antibodies diluted in TBS-T with 5% bovine serum albumin. Following three washes with TBS-T, the membrane was incubated for 2 h at room temperature with anti-rabbit IgG conjugated with horse-radish peroxidase (cat. no. sc-2374; Santa Cruz Biotechnology, Inc.) at 1:2,000 dilution in TBS-T containing 5% non-fat dried milk. Cells were washed three times for 10 min with TBS-T and protein bands were detected by chemiluminescence, using a Phototope Detection system in accordance with the manufacturer's instructions (Beyotime Institute of Biotechnology, Shanghai, China). The gray values of protein bands were analyzed using Image J software (ver. 1.46; National Institutes of Health, Bethesda, MD, USA). The relative expression of the target protein was calculated according the following formula: Relative expression of target protein = target protein gray value/contingent internal reference value.

**Statistical analysis.** Results are expressed as the mean ± standard deviation. A minimum of three assays were performed. Comparisons were made using analysis of variance analysis, followed by a least significant difference test and subsequent homogeneity of variance test. Correlation analysis between the groups was performed using Pearson correlation analysis. SPSS 21.0 software (IBM SPSS, Armonk, NY, USA) was used for statistical analyses. P<0.05 was considered to be statistically significant difference.

**Results**

**Evaluation of the isolated primary synovial fibroblasts.** Primary synovial fibroblasts, derived from the fibrous membrane of SD rat knees, were used as the model cells for the present study. Cells were isolated using the tissue culture (Fig. 1A) method and were passaged to the third generation (Fig. 1B) and subsequently evaluated by immunochemical staining and flow cytometry. Type B synoviocytes, or FLS, are mesenchymal cells that display various characteristics of fibroblasts, including expression of type IV and V collagens, vimentin, and CD90. Cells that were stained brown indicated the presence of vimentin (Fig. 1C), compared with the cells in negative control group (Fig. 1D). Using flow cytometry, while cells marked with homotype-negative phycoerythin-labeled IgG antibodies showed negative result (Fig. 1E), CD90-positive cells were detected and confirmed the phenotype of type B synovial fibroblasts (Fig. 1F).

**IL-1β induces the expression of fibroblast inflammatory cytokines.** Primary synovial fibroblasts derived from the fibrous membrane of SD rat knees were used as model cells to examine the expression levels of inflammatory factors induced by IL-1β. Expression levels of the inflammatory cytokines, IL-1β and TNF-α were significantly increased following induction with 10 ng/ml IL-1β for 24 h, when compared with the control, group A (P<0.05; Fig. 2), which indicated that IL-1β leads to an inflammatory state in a cell. However, the expression levels of inflammatory cytokines significantly
Figure 1. Evaluation of isolated synovial fibroblasts from passage 3 using vimentin staining and CD90 detection by flow cytometry. (A and B) Isolated cells from synovial membranes of Sprague-Dawley rat knees (magnification, x40). (C) Vimentin in the synovial cells was stained brown (magnification, x100; immunocytochemical staining with diaminobenzidine chromogen). (D) PBS was used instead of vimentin staining to act as a negative control (magnification, x100). (E) Cells were marked with homotype-negative phycoerythrin-labeled IgG antibody (negative control). (F) Positive cells marked with CD90 were detected by flow cytometry and indicated that the phenotype conformed to synovial fibroblasts. CD90, cluster of differentiation-90; IgG, immunoglobulin G.

Figure 2. IL-1β inducible inflammatory factor expression in synovial fibroblasts. Synovial fibroblasts were stimulated with 10 ng/ml IL-1β for 24 h. Following a further 24 h, cells in group A were treated with Dulbecco's modified Eagle's medium medium without glucocorticoid; group B, with 10 ng/ml IL-1β; group C, with 10 ng/ml IL-1β and 10^-6 mmol/l corticosterone; and group D, as group C plus 100 nmol/l PF. Expression of (A) IL-1β and (B) TNF-α mRNA increased significantly in the IL-1β-induced synovial fibroblasts. When corticosterone was applied, the expression of inflammatory cytokines significantly decreased. The 11β-HSD1 inhibitor, PF, reversed the reduction in expression of inflammatory factors caused by corticosterone. Results are expressed as the mean ± standard deviation and a minimum of three assays were performed.

Figure 3. mPGES-1 mRNA and PGE2 expression in the inflammatory synovial fibroblasts. Total RNA was isolated from synovial fibroblasts layers, and mRNA was quantified using reverse transcription-quantitative polymerase chain reaction. PGE2 was detected by ELISA. (A) mPGES-1 mRNA and (B) PGE2 expression levels were significantly increased in the synoviocytes induced by IL-1β when compared with the control, group A. However, IL-1β-induced expression was inhibited significantly by corticosterone, and the inhibitory effect of corticosterone was significantly reversed by the 11β-HSD1 inhibitor PF. Group A, synovial fibroblasts treated with medium without glucocorticoid (control); group B, synovial fibroblasts treated with 10 ng/ml IL-1β; group C, synovial fibroblasts treated with 10 ng/ml IL-1β and 10^-6 mmol/l corticosterone; group D, synovial fibroblasts treated as group C plus 100 nmol/l PF. *P<0.05 vs. control; ##P<0.05 vs. group C. C, corticosterone; PF, PF915275 [4'-cyano-biphenyl-4-sulfonic acid (6-amino-pyridin-2-yl)-amide]; mPGES-1, microsomal prostaglandin E synthase-1; PGE2, prostaglandin E2.
decreased compared with those in group B following the administration of corticosterone (P<0.05; Fig. 2), suggesting an inhibitory effect of corticosterone on IL-1β-induced inflammation. However, when cells were treated with PF915275 (group D), an 11β-HSD1 inhibitor, the decreased expression levels of inflammatory factors (suppressed by corticosterone) exhibited a reversal effect, with a significant increase (P<0.05) in the expression levels of IL-1β and TNF-α compared with those in group C (Fig. 2). In comparison with group B, a relatively low expression level of the inflammatory factors IL-1β and TNF-α was observed in group D (P<0.05).

PGE₂ is a key factor involved in the development and perpetuation of inflammation in disease, such as with rheumatoid arthritis. In this disease, local inflammation of synovial tissue is characterized, in part, by increased local levels of PG, predominantly PGE₂ (5). Inducible mPGES-1 has an essential role in the localized increase of PGE₂ during inflammatory arthritis. Expression notably increased when cells were induced by pro-inflammatory cytokines (IL-1β and TNF-α) in vitro (6). In the present study, mPGES-1 mRNA expression and PGE₂ production of synovial fibroblasts were significantly increased following treatment with 10 ng/ml IL-1β for 24 h, when compared with the control, group A (P<0.05; Fig. 3).

**IL-1β induces 11β-HSD1 expression in fibroblasts by suppressing corticosterone activation.** 11β-HSD1 is capable of converting inactive glucocorticoids (cortisone and prednisone) into the active counterparts, cortisol and prednisolone. Synovial fibroblasts and osteoblasts generate active glucocorticoids through the expression of 11β-HSD1 (7). Such activity increases, *in vitro*, in response to pro-inflammatory cytokines or glucocorticoids. The present study indicated that IL-1β significantly increased 11β-HSD1 expression levels in synoviocytes (P<0.05), whereas these expression levels markedly decreased when corticosterone was added to the media (Fig. 4), which may be due to an increase in the production of active cortisol (Fig. 5) and the inhibition of the synoviocyte inflammation. When the activity of 11β-HSD1 was inhibited by PF915275, the 11β-HSD1 mRNA and protein expression levels that had previously been suppressed by corticosterone were reversed, and the expression levels were higher than those of cells stimulated with IL-1β (P<0.05; Fig. 4).

Corticosterone suppresses synoviocyte PGE₂ production by promoting the activity of 11β-HSD1. Synovial fibroblasts expressed high levels of mPGES-1 mRNA when treated with IL-1β, and PGE₂ concentration was higher in the media when compared with the control, group A, as detected by ELISA (P<0.05; Fig. 3). Corticosterone appeared to counteract the effects of IL-1β, as indicated by the reduction in expression of mPGES-1 mRNA and PGE₂ (P<0.05; Fig. 3). This inhibitory effect may result from a low level of active corticosterone that inhibited 11β-HSD1 expression; thus corticosterone may be
effectively converted by synoviocytes (Fig. 5). In the present study, PF915275 was used to suppress the conversion activity of $11\beta$-HSD1. Suppression by PF915275 caused a significant increase in the levels of mPGES-1 mRNA (Fig. 3A) expression in synoviocytes and the concentration of PGE$_2$ (Fig. 3B) in the media, when compared with the control and corticosterone groups (groups A and C, respectively; $P<0.05$). mPGES-1 mRNA and PGE$_2$ were not inhibited when the conversion activity of enzyme $11\beta$-HSD1 was blocked ($P>0.05$, compared with the control group; Fig. 3). These results suggest that PGE$_2$ production by the synoviocytes may be suppressed by activating the $11\beta$-HSD1 pathway.

$11\beta$-HSD1 expression positively correlates with PGE$_2$ concentration in inflammatory synoviocytes. Synovial fibroblasts treated with IL-1$\beta$ exhibited significantly increased mRNA levels of $11\beta$-HSD1 (Fig. 4A) and mPGES-1 (Fig. 3A) when compared with the normal synoviocytes ($P<0.05$). $11\beta$-HSD1 and PGE$_2$ proteins were increasingly expressed in the inflammatory synoviocytes; however, the association between $11\beta$-HSD1 and PGE$_2$ was unclear. $11\beta$-HSD1 mRNA and protein expression levels in the inflammatory synoviocytes were found to be positively correlated with PGE$_2$ concentration ($r=0.74$, $P<0.01$ and $r=0.94$, $P<0.01$; Fig. 6). The positive correlation between $11\beta$-HSD1 and mPGES-1 mRNA expression levels was significant ($r=0.97$, $P<0.01$; Fig. 6).

Discussion

Synovial fibroblasts are important functional compartments that contribute to the production of lubricious synovial fluid and diffusion of the nutrients (16). Synovial fibroblasts are mesenchymal cells that display multiple characteristics of fibroblasts, including expression of type IV and V collagens, vimentin, and CD90 (Thy-1) (16). CD90, which is evolutionarily conserved and developmentally regulated, often has marked effects on cell phenotype and is a specific surface marker expressed by synovial fibroblasts (17,18). In isolated situations, the expression or lack of expression of CD90 has been used to separate fibroblast subsets (19). Vimentin is the specific protein marker expressed by cells derived from the mesoderm and is used for cell isolation (20). In the present study, purified cells derived from synovial membranes in SD rat knees were stained vimentin-positive, due to the expression of the surface marker CD90. These results indicated that the cells were confirmed to be synovial fibroblasts, which was consistent with the literature (21). However, the proliferative activity of synovial fibroblasts declined significantly in the seven generations of culturing. Therefore, FLS at passage three, with high purity and suitable activity were used in the present study.

Synovial inflammation is an important contributor to OA pathogenesis (22,23). Synovitis has been demonstrated to correlate with OA symptom severity, and hormonal factors, such as cytokines and chemokines, are important for crosstalk in joint tissues (23,24). These mediators have a critical role in the development of inflammation and induce catabolic changes in joint tissues (23,25,26). Studies have demonstrated that FLS produce inflammatory cytokines (such as TNF-$\alpha$ and IL-1$\beta$) and chondrolytic mediators, including matrix metalloproteinases (27). IL-1$\beta$ and TNF-$\alpha$, which are the two most extensively studied factors, have been implicated in the pathogenesis of OA (22,23). In the present study, when stimulated by IL-1$\beta$, FLS expressed increased levels of IL-1$\beta$, TNF-$\alpha$ and mPGES-1 and produced high concentrations of PGE$_2$ in the medium. The present data indicated that the inflammatory state in the FLS induced by IL-1$\beta$ was suitable for this study.

The tissue availability of active glucocorticoids is dependent on their rate of synthesis from cholesterol, downstream metabolism, excretion and interconversion (28). The latter is mediated by $11\beta$-HSDs. $11\beta$-HSD is a glucocorticoid pre-receptor regulating enzyme that regulates the local concentration of glucocorticoid through its oxidation effect (9). Non-oxidized corticosteroids, such as cortisol and 11-dehydrogenation of corticosterone, cannot combine with the glucocorticoid receptor and thus have an effective role in biological function. The only way to promote activation is by $11\beta$-HSD1 transformation (29). A prior study considered that the sustained inflammatory state of arthritis is due to the partial abnormal hormone metabolism in tissues (30). Synovitis is an important cause of chronic, persistent OA. In animal models of arthritis, $11\beta$-HSD1 gene knockout increases the risk of original joint inflammation (31), suggesting a positive effect of...
11β-HSD1 on controlling the severity of arthritis. However, the underlying mechanism remains unknown. 11β-HSD1-related genes and proteins of IL-1β and TNF-α are upregulated in various stromal cells (32). In this study, the cortisol concentration in the medium was detected by ELISA to determine the conversion activity of 11β-HSD1. 11β-HSD1 was highly expressed when stimulated by IL-1β, whereas cortisol concentration remained at a low level. As the substrate corticosterone was added, the concentration increased significantly. By contrast, 11β-HSD1 levels decreased, indicating 11β-HSD1 was blocked by PF915275. These results suggested that FLS had an active transformation ability in cortisol activation; and the cortisol/cortisone ratio in the medium and 11β-HSD1 expression were regulated under a feedback mechanism.

The present study also demonstrated that normal synovial fibroblasts, induced by IL-1β, highly expressed 11β-HSD1 gene and protein. Li et al (33) reported that IL-1β and other inflammatory mediators are able to jointly upregulate 11β-HSD1 mRNA in amnion; however, it has not yet been reported in the literature whether a similar regulating mechanism in synovial fibroblasts exists. In the present study, interactions between IL-1β and glucocorticoid were identified. When synovial fibroblasts were interfered with IL-1β and corticosterone, the expression of 11β-HSD1 decreased, which suggested that functional normal synovial cells can be activated by the enzymatic conversion of corticosterone and produce corticosteroids (cortisol) locally to exert anti-inflammatory effects. Therefore, high expression of 11β-HSD1 may attenuate the inflammatory state. By contrast, when enzyme activity and/or glucocorticoid receptor was inhibited, the concentration of local activated cortical hormone decreased and the probability of binding with its receptor also decreased, thus 11β-HSD1 failed to suppress the local inflammatory state effectively. However, when the involved inflammatory factors increased to high levels, expression of 11β-HSD1 was initiated and increased in response to inflammation. These data suggest that hormone levels regulated by 11β-HSD1 pre-receptor have a critical role in controlling synovitis, and such regulation may have a feedback mechanism.

In the pathological process of OA, the secretion of inflammatory factors, such as PGE₂, have an important role in the development of synovitis, cartilage matrix disintegration and bone destruction (8). PGE₂ dilates blood vessels and induces alterations in second messenger level via autocrine/paracrine signaling (34). Jia et al (35) identified that PGE₂ receptors present in articular tissue may combine with PGE₂, and subsequently induce cartilage degradation, glycosaminoglycan loss and collagen type II degradation in OA animal models. However, these pathological changes can be alleviated in PGE₂ receptor knockout mice. Sun and Myatt (11) reported that IL-1β was able to significantly improve 11β-HSD1 mRNA expression and activity. With prior induction of 11β-HSD1 expression by dexamethasone, cortisone induced more PGE₂ production in the amnion fibroblast. This study suggests that glucocorticoids are able to positively induce 11β-HSD1 expression in amnion fibroblasts and this effect was further strengthened by pro-inflammatory cytokines. However, in synovial B cells, whether glucocorticoids affect 11β-HSD1 expression and PGE₂ biosynthesis remains unclear. In the present study, IL-1β significantly induced mPGEs-1 and PGE₂ expression in synoviocytes, and corticosterone effectively inhibited this effect. However, when the transformation activity of 11β-HSD1 was blocked by PF915275, the activated corticosterone (cortisol) production decreased significantly and the inhibitive effect of PGE₂ was reversed. This indicated that the pre-receptor activity is able to effectively regulate the expression and biological effect of PGE₂ in FLS.

Both 11β-HSD1 and mPGEs are present in microsomes and co-exist in the complex pathogenesis of OA. PGE₂ is an important inflammatory factor in the human body. 11β-HSD1 has an important role in regulating anti-inflammatory substance levels, and thus activating glucocorticoid hormone mPGEs-1, which is an inducible key limited enzyme, to control PGE₂ production in inflammation process (25). In the present study, a preliminary analysis was constructed to elucidate the association between the expression and production of mPGEs-1, PGE₂, and 11β-HSD1 in synoviocytes under an inflammatory state. The results revealed significant positive linear correlations between them, both at the gene and protein expression levels. To date, no reports exist on whether PGE₂ and 11β-HSD1 promote each other in synovial B cells under osteoarthritic conditions. Subsequently, the underlying mechanisms of interactions between 11β-HSD1, mPGEs-1 and PGE₂ require further investigation.

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