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L. Chastine Bell-Parikh, ... , Muredach Reilly, Garret A. FitzGerald

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Biosynthesis of 15-deoxy- $\Delta^{12,14}$ -PGJ₂ and the ligation of PPAR γ

See the related Commentary beginning on page 828.

L. Chastine Bell-Parikh, Tomomi Ide, John A. Lawson, Peter McNamara, Muredach Reilly, and Garret A. FitzGerald

Center for Experimental Therapeutics, University of Pennsylvania School of Medicine, Philadelphia, Pennsylvania, USA

15-deoxy- $\Delta^{12,14}$ -PGJ₂ (15d-PGJ₂) has been identified as an endogenous ligand for PPAR γ , inducing adipogenesis in vitro. Additional roles for this molecule in the propagation and resolution of inflammation, ligation of NF- κ B, and mediation of apoptosis have been proposed. However, quantitative, physicochemical evidence for the formation of 15d-PGJ₂ in vivo is lacking. We report that 15d-PGJ₂ is detectable using liquid chromatography–mass spectrometry–mass spectrometry at low picomolar concentrations in the medium of 3T3-L1 preadipocytes. However, despite induction of COX-2, production of PGs, including 15d-PGJ₂, does not increase during adipocyte differentiation, a process unaltered by COX inhibition. 15d-PGJ₂ is detectable as a minor product of COX-2 in human urine. However, its biosynthesis is unaltered during or after COX activation in vivo by LPS. Furthermore, the biosynthesis of 15d-PGJ₂ is not augmented in the joint fluid of patients with arthritis, nor is its urinary excretion increased in patients with diabetes or obesity. 15d-PGJ₂ is not the endogenous mediator of PPAR γ -dependent adipocyte activation and is unaltered in clinical settings in which PPAR γ activation has been implicated.

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Introduction

The PPAR nuclear hormone transcription factors heterodimerize with retinoic acid receptors to regulate expression of distinct gene cassettes (1, 2). Although three PPAR isotypes (α , δ , and γ) have been well characterized, the identity of their natural ligands remains controversial (3–6). PPAR γ is expressed predominantly in adipose tissue, colon and macrophages, and to a lesser extent, in vascular smooth muscle cells (1–3). Among the genes regulated by PPAR γ , many are involved in lipid metabolism,

including lipoprotein lipase (LPL), adipocyte fatty acid binding protein (aP2), acyl-CoA synthase, and fatty acid transport protein (2, 3). Synthetic PPAR γ agonists promote the maturation of fibroblasts to adipocytes in vitro (2, 3) and one class of such agonists, the glitazones, has been used in the treatment of diabetes. In addition, synthetic PPAR γ agonists promote uptake of modified lipids by macrophages (7, 8) and may retard atherogenesis in mice (9). The increasing convergence of insulin resistance, obesity, and cardiovascular morbidity in the metabolic syndrome has further focused attention on the molecular mechanisms of PPAR γ activation.

PGs are formed through the sequential actions of COXs and a variety of PG synthases, which are expressed with some tissue specificity (10, 11). PGD₂ is formed abundantly in the brain (12) and in mast cells (13). It undergoes spontaneous dehydration to PGJ₂ in vitro, a reaction enhanced by albumin-induced catalysis, which yields a number of additional derivatives, including 15-deoxy- $\Delta^{12,14}$ -PGJ₂ (15d-PGJ₂) (14, 15). This compound, like other PGs (16), can be actively transported into cells (17). In contrast to conventional PGs, PGJ₂ and its derivatives possess a highly reactive cyclopentenone (CP) ring (18), which permits them to ligate nuclear receptors and to modify intracellular signaling molecules. CP-PGs exhibit biologic properties in vitro of potential relevance to inflammation, cellular proliferation, and differentiation (18, 19).

In addition to its effects on adipocyte maturation, 15d-PGJ₂ has been implicated in the PPAR γ -dependent propagation (20) and resolution (21) of inflammation, and in the fibrotic reaction to activation of protease-

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Address correspondence to: Garret A. FitzGerald, Center for Experimental Therapeutics, University of Pennsylvania School of Medicine, 153 Johnson Pavilion, 3620 Hamilton Walk, Philadelphia, Pennsylvania 19104-6084, USA. Phone: (215) 898-1184; Fax: (215) 573-9135; E-mail: garret@spirit.gcrc.upenn.edu. Peter McNamara's present address is: Phenomix Corp., La Jolla, California, USA.

L. Chastine Bell-Parikh and Tomomi Ide contributed equally to this work.

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Nonstandard abbreviations used: 15-deoxy- $\Delta^{12,14}$ -PGJ₂ (15d-PGJ₂); lipoprotein lipase (LPL); adipocyte fatty acid binding protein (aP2); cyclopentenone (CP); gas chromatography-mass spectrometry (GC-MS); 9-deoxy- $\Delta^{9,12}$ -(E)-PGD₂ (Δ^{12} -PGJ₂); bisphenol A diglycidyl ether (BADGE); triacylglyceride (TAG); 3-isobutyl-1-methylxanthine (IBMX); differentiation medium (DM); solid phase extraction (SPE); liquid chromatography-mass spectrometry-mass spectrometry (LC-MS-MS); multiple reaction monitoring (MRM); electrospray ionization (ESI); 2,3-dinor-6-keto-PGF_{1 α} (PGI-M); isoprostane (iP); thromboxane (Tx); methoxime (MO); negative ion (NI); General Clinical Research Center (GCRC); pentafluorobenzyl (PFB).

activated receptor 2 by mast cell tryptase (22). Alternatively, 15d-PGJ₂ induces apoptosis and growth inhibition *in vitro* via ligation of NF-κB (23) and/or IκB kinase (24) by its reactive CP moiety.

Despite much discussion of the biologic activities of 15d-PGJ₂, there is actually scant physicochemical evidence for its formation *in vivo* (25). A commonly used immunoassay for the compound was withdrawn from the market (26). Dehydration and isomerization products of PGD₂ metabolism have been measured in humans (27). However, the concentrations of 15d-PGJ₂ that activate PPARγ or ligate proteins in the NF-κB signaling pathway *in vitro* typically range from 2.5 to 100 μM (5), whereas biologically active concentrations of conventional exogenous PGs are generally in the low picomolar range (28, 29). Thus, although *in vitro* albumin-catalyzed dehydration of PGD₂ yields levels of 15d-PGJ₂ detectable by gas chromatography-mass spectrometry (GC-MS), 15d-PGJ₂ has not been detected by physicochemical methods *in vivo* (27, 30). Similarly, isolated cells exposed to PGD₂ *in vitro* form detectable amounts of both PGJ₂ and a distinct CP derivative, 9-deoxy-Δ^{9,12}-(E)-PGD₂ (Δ¹²-PGJ₂), but not 15d-PGJ₂ (26, 31). Δ¹²-PGJ₂ has been detected in the urine by immunoassay (31), but PGJ₂ has never been detected as a urinary metabolite of PGD₂. Thus, discordance exists between the commonly reported role of 15d-PGJ₂ as a PPAR agonist and evidence for its formation *in vivo*. We addressed the hypothesis that it was formed *in vivo* by developing a specific quantitative mass spectrometric assay for 15d-PGJ₂. We investigated its role in adipocyte maturation and determined whether its biosynthesis was altered in clinical settings in which PPAR activation has been implicated.

Methods

Materials. Arachidonic acid (peroxide free), all unlabeled and [²H]-labeled PG standards, [³H]15d-PGJ₂ and COX antibodies (polyclonal, rabbit), indomethacin, NS-398 ciglitazone, and bisphenol A diglycidyl ether (BADGE), a synthetic PPARγ antagonist (32), were obtained from Cayman Chemical (Ann Arbor, Michigan, USA). The details of the various isomers of the 15d-PGJ₂ were described previously (26). [¹⁸O₂]-labeled PGJ₂ standard was synthesized from PGJ₂ using the method of Wescott et al. (33). Murine monoclonal PPARγ (E-8) antibody was purchased from Santa Cruz Biotechnology Inc. (Santa Cruz, California, USA). Reagents for quantification of triacylglycerol (TAG) were provided in an L-Type TG H Enzyme Color A Kit purchased from Wako Chemicals USA Inc. (Richmond, Virginia, USA). Oil red O, insulin, dexamethazone, 3-isobutyl-1-methylxanthine (IBMX), and ibuprofen were obtained from Sigma-Aldrich (St. Louis, Missouri, USA).

Cellular incorporation of 15d-PGJ₂. One micromolar of 15d-PGJ₂ containing 0.5 μCi [³H]15d-PGJ₂ was incubated with confluent 3T3-L1 cells cultured in six-well plates (2 ml per well). The persistence of 15d-PGJ₂ in the

medium was estimated by liquid scintillation counting. The cells were washed at 48 hours with PBS and harvested with 0.05 % trypsin. The 15d-PGJ₂ in the wash liquid and cell pellet was estimated in the same manner.

Cell culture and adipocyte differentiation assay. The 3T3-L1 cells (ATCC, Manassas, Virginia, USA) were maintained in DMEM with 10% fetal calf serum under nondifferentiating and differentiating conditions as described previously (34). Two days after cell confluence, cells were cultured in differentiation medium (DM) (day 0), supplemented with 10 μg/ml insulin, 1 μM dexamethasone, and 0.5 mM IBMX for 2 days and switched to a post-DM, which was supplemented only with 10 μg/ml insulin. Medium was collected from the cells at sequential 24-hour intervals for determinations of PGs. Cells were collected from parallel experiments for protein and mRNA analysis. The 3T3-F442A cells (obtained from Howard Green, Harvard School of Medicine, Boston, Massachusetts, USA) were maintained and induced to differentiate as previously described (35).

PG extraction. Samples (5.0 ml urine or cell-free tissue culture medium or 500 μl synovial fluid) were spiked with 1.0 ng [²H₄]15d-PGJ₂ and extracted on C18 solid phase extraction (SPE) cartridges. The cartridges were washed with 25% ethanol in water and eluted with ethyl acetate. Some samples for 15d-PGJ₂ were subjected to silica gel TLC after SPE, using a mobile phase of methanol/ethyl acetate/formic acid (95:5:0.1). Silica bands (10 mm) were extracted with ethyl acetate. The cells were washed twice with PBS for extraction of intracellular 15d-PGJ₂ (36), harvested in 0.6 ml PBS with 1 ng [²H₄]15d-PGJ₂, and then freeze-thawed twice in liquid N₂. Chloroform (1 ml) and methanol (2 ml) were added, followed by vigorous vortexing and further addition of 1 ml each chloroform and water. The organic phase was taken for the analysis.

PG analysis. Liquid chromatography-mass spectrometry-mass spectrometry (LC-MS-MS) analyses were performed on a Micromass Quattro Ultima triple quadrupole analyzer (Macromass, Beverly, Massachusetts, USA) operated in multiple reaction monitoring (MRM) mode. The collision cell was maintained at 2 × 10⁻³ T with argon. A coaxial electrospray ionization (ESI) probe interfaced the LC-MS-MS with dual LC-10ADVP syringe pumps (Shimadzu, Kyoto, Japan). HPLC separations used Hypersil C18-BD, 3 μm, 150 × 2-mm columns (Thermo Hypersil-Keystone, Bellefonte, Pennsylvania, USA). Linear gradients were performed from water (solvent A) and acetonitrile/methanol, 95:5 (solvent B), both containing 0.005% acetic acid adjusted to pH 5.7 with ammonium hydroxide. A flow rate of 200 μl/min was used for all assays.

A gradient of 20% to 60% B in 30 minutes was used for 15d-PGJ₂ analysis. Transitions monitored were *m/z* 315.2 → 271.0 for 15d-PGJ₂ and *m/z* 319.2 → 275.0 for [²H₄]15d-PGJ₂. MS-MS settings were as follows: collision energy 13 eV, capillary 3 kV, source 70°C, and desolvation 250°C.

All other PGs were prepared similarly, except that the SPE columns were washed with 5% methanol in water. GC-MS analysis for PGE₂ (37), 6-keto-PGF_{1α} (38), 2,3-dinor-6-keto-PGF_{1α} (PGI-M) (39), and 2,3-dinor TxB₂ (40), and LC-MS analysis of 8,12-*iso*-iPF_{2α}-VI (39) were performed as previously described. A gradient of 20% to 40% B in 15 minutes at a flow rate of 200 μl/min was used for ESI-LC-MS-MS detection of PGE₂. MS-MS data were generated using the following settings: cone 1.0 V, collision energy 18 eV, capillary 2.5 kV, source 150°C, and desolvation 150°C. Transitions monitored were *m/z* 351.2 → 271.0 for PGE₂ and *m/z* 355.2 → 275.0 for the [²H₄]PGE₂. The methoxime (MO) derivative of 6-keto-PGF_{1α}, was formed by drying the sample after SPE and incubating it overnight with 20 μl of a 0.5% solution of MO-HCl in pyridine. The HPLC gradient was 10–50% B in 18 minutes. The MS settings were as follows: collision energy 22 eV, capillary 2.5 kV, source 70°C, and desolvation 200°C. Transitions monitored were *m/z* 398.2 > 306.0 for 6-keto-PGF_{1α}-MO and *m/z* 402.2 > 310.0 for [²H₄]6-keto-PGF_{1α}-MO.

Oil red O staining and TAG assay. Cells were washed twice with PBS, fixed with 10% formaldehyde, and stained with Oil red O. Cellular TAG was extracted as previously described (41) with an additional overnight incubation in isopropanol. Total TAG was determined using a TAG assay kit (Wako Chemicals USA Inc.).

Preparation of whole cell extracts, gel electrophoresis, and immunoblotting. Cell contents were extracted by sonication into lysis buffer (50 mM Tris-HCl, pH 7.4, 100 μM EDTA, 100 μM EGTA, 1% NP-40, 0.1% SDS, 0.1% deoxycholic acid) containing protease inhibitor cocktail tablets (Complete; Roche, Nutley, New Jersey, USA). Total cellular protein (20 μg) was separated by electrophoresis on NuPAGE 4–12% Bis-Tris Gel (Invitrogen, Carlsbad, California, USA) and transferred to nitrocellulose membrane.

Antibodies for COX-1 and COX-2 (Cayman Chemical) were used at 1:500 and 1:1000 dilutions, respectively. Reagents for blocking, incubation, washing, and chemiluminescent-detection (WesternBreeze; Invitrogen) were used.

RNA preparation and analysis. RNA isolation and reverse transcription were performed with TRIzol Reagent (Invitrogen) and Superscript RT (Invitrogen). Twenty micrograms of total RNA were separated by electrophoresis in a 1% agarose gel and transferred to a nylon membrane, Hybond N+ (Amersham Biosciences, Piscataway, New Jersey, USA) for Northern analysis. The cDNA probes for αP2, PPARγ, and actin were radiolabeled with [α-³²P]dCTP using a random prime-labeling system, Rediprime II (Amersham) and hybridization was performed at 42°C in Rapid-hyb buffer (Amersham).

Clinical studies. The Institutional Review Board and the General Clinical Research Center (GCRC) Advisory Board of the University of Pennsylvania approved the clinical study protocols. Urine samples

for the inhibitor studies were collected before and after 6 days of treatment with aspirin (81 mg), ibuprofen (400 mg) three times a day, or celecoxib (200 mg twice daily), each in six healthy volunteers. Volunteers were 18–35 years of age, within 30% of ideal body weight, and had an unremarkable medical history and physical examination and routine hematology and biochemistry. Smokers and subjects with any bleeding disorders, allergy to aspirin or other NSAIDs, and history of any gastrointestinal or cerebrovascular disease were excluded. Subjects abstained from aspirin or other NSAIDs for at least 2 weeks before enrollment. The samples for urinary analysis of PGs were obtained in a study of bacterial LPS administration to healthy volunteers performed in the GCRC (39). Samples for the diabetic studies were obtained from a group of 200 maturity-onset diabetic patients (ages 35–75), without cardiovascular disease or symptoms, who were recruited to a cross-sectional observational study of risk factors for coronary atherosclerosis as measured by coronary artery calcification at electron beam computed tomography. Age, sex, and ethnically matched nonsmoking diabetic patients in the lowest and highest weight quartiles were selected for study. All abstained from aspirin, NSAIDs, COX-2 inhibitors, and PPAR agonists for at least 4 weeks before and during the study.

Statistics. Data are expressed as the mean ± SD of the mean. Data were subject to ANOVA and subsequent pairwise analysis, using two-tailed *t* tests, as appropriate.

Results

Incorporation of 15d-PGJ₂ into cells. We first examined the stability of 15d-PGJ₂ in aqueous phase. When 1 μM or 10 μM of [²H₄]15d-PGJ₂ was incubated in a 24-hour “spent” cell-free medium, the spike was 100% recoverable from the medium at 48 hours (data not shown). When 3T3-L1 fibroblast cells were incubated with 1 μM of [³H]15d-PGJ₂, no decrease in ³H in the medium was observed over 48 hours and only 0.5% of the initial counts were recovered from the cell pellet (Figure 1a).

Generation of 15d-PGJ₂ by 3T3-L1 preadipocytes. The 3T3-L1 fibroblast cells may be induced to differentiate to adipocytes by PPARγ ligands or by transient exposure to DM. We sought initially 15d-PGJ₂ in lipid extracts from the medium of these cells before their maturation to adipocytes, using negative ion (NI)-ESI-LC-MS-MS. A mass chromatogram of the cell medium revealed peaks corresponding to a parent ion (*m/z* 315.2) and an associated collision-induced transition ion (*m/z* 271.0) characteristic of 15d-PGJ₂ (Figure 1b).

15d-PGJ₂ was detectable (0.64 ± 0.05 pg/ml per 3 hours; ~2 pM) in the culture medium without serum under subconfluent conditions. Synthesis of substantially greater amounts of PGE₂ was also evident (0.98 ± 0.13 ng/ml per 3 hours; ~3 nM) in the same medium. Supplementation of the cells with substrate arachi-

donic acid (15 μ M) further stimulated formation of both PGs (22.5 \pm 3.1 pg/ml per 3 hours; \sim 71 pM for 15d-PGJ₂, and 18.3 \pm 1.8 ng/ml per 3 hours; \sim 52 nM for PGE₂) (Figure 1c) and induced a transient upregulation of COX-2 protein expression (Figure 1d). COX-1 protein was unaltered by arachidonate treatment (Figure 1d). The nonselective COX inhibitor, indomethacin (3 μ M), depressed arachidonate-induced stimulation of 15d-PGJ₂ by 84% and PGE₂ by 99%. The COX-2 selective inhibitor, NS-398 (IC₅₀s for COX-2 and COX-1 are 1.0 μ M and 65 μ M, respectively) depressed these PGs in a dose-dependent manner (Figure 1c), establishing that substrate enriched 3T3-L1 cells can form picomolar concentrations of 15d-PGJ₂ predominantly via the COX-2 isoform.

Low concentrations (1.0 \pm 0.4 nM) of intracellular 15d-PGJ₂ were also detectable in the lipid extract under subconfluent conditions. Intracellular 15d-PGJ₂ was evaluated after glutathione depletion to assess the possibility that thiol conjugation might account for the apparently low levels of 15d-PGJ₂. Glutathione depletion was achieved by exposure of cells to the γ -glutamylcysteine synthetase, DL-buthionine-sulfoximine 6 mM (42) for 24 hours. This treatment failed to increase detectably intracellular 15d-PGJ₂ when compared with vehicle (1.1 \pm 0.3 nM vs. 1.0 \pm 0.4 nM).

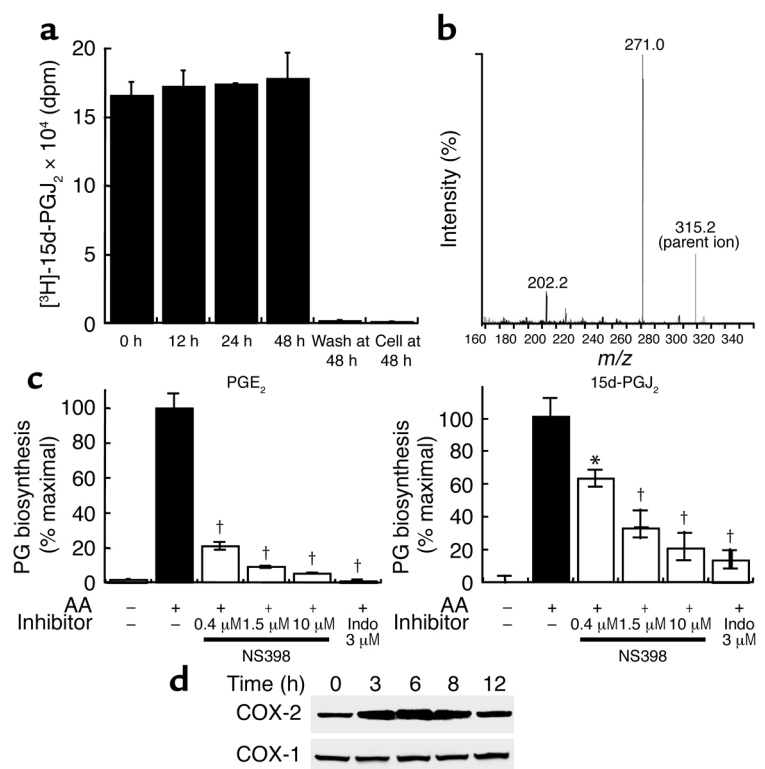
The role of 15d-PGJ₂ in PPAR γ -mediated adipocyte differentiation was investigated using 3T3-L1 fibroblasts induced to differentiate by transient exposure to DM or a PPAR γ agonist. The extent of 3T3-L1 differentiation to mature adipocytes was assessed by measurement of cellular TAG (Figure 2a), by appear-

ance of adipocyte-specific markers such as PPAR γ and aP-2 (Figure 2b and c), and by Oil red O staining of accumulated lipids (Figure 2d). Comparable induction was achieved by treatment with DM or the PPAR γ agonist, ciglitazone (15 μ M), as reflected by TAG accumulation (1.9 \pm 0.1 vs. 1.9 \pm 0.2 mg/10⁶ cells; Figure 2a) and by PPAR γ and aP2 expression (Figure 2b and c) by day 10.

The expression of COX-1 mRNA and protein are unaltered during the course of DM-induced differentiation of 3T3-L1 cells to mature adipocytes (Figure 2b). COX-2 mRNA, however, is depressed initially on addition of DM (Figure 2b). After removal of DM (on day 2), COX-2 mRNA is progressively upregulated, and marked protein expression is evident by day 10 (Figure 2c), coincident with adipocyte maturation. Analysis of medium in differentiated adipocytes for PGE₂, 6-keto PGF_{1 α} (the hydrolysis product of PGI₂), PGF_{2 α} , and 15d-PGJ₂ established that the predominant products were PGE₂ and PGI₂ (Table 1). PGJ₂ was undetectable. The concentrations of PGE₂, 6-keto PGF_{1 α} , and 15d-PGJ₂ all declined sharply after the cells reached confluence on day -2 (data not shown). Addition of DM resulted in a further detectable reduction in formation of these PGs, coincident with the transient reduction in expression of COX-2 (Figure 2b; 6-keto PGF_{1 α} not shown). However, synthesis of all three products remained depressed, despite consequent upregulation of COX-2 during adipocyte differentiation. The relative amounts of the three PGs, PGE₂, 6-keto-PGF_{1 α} , and 15d-PGJ₂, in mature adipocytes in excess of the background levels in the

Figure 1

COX-dependent biosynthesis of 15d-PGJ₂ by 3T3-L1 preadipocytes. (a) Cellular incorporation of 15d-PGJ₂. 3T3-L1 cells were incubated with 15d-PGJ₂ mixed with [³H]15d-PGJ₂ for 48 hours. Vertical bars represent dpm in the medium at the times indicated and in the cell wash and cell pellet after 48 hours. (b) Mass spectrum of standard 15d-PGJ₂. Spectrum obtained by NI-ESI-LC-MS-MS. For LC-MS-MS conditions, see Methods section. (c) Substrate enhanced, COX-dependent formation of PGs. Stimulation of 15d-PGJ₂ and PGE₂ formation in 3T3-L1, 1-day preconfluent cells, pretreated with 15 μ M arachidonic acid (AA) alone for 1 hour and in the presence of the nonselective COX inhibitor, indomethacin (Indo) (3 μ M), or the COX-2 selective inhibitor, NS-398 (0.4–10 μ M) in serum-free medium. Formation of PGE₂ and 15d-PGJ₂ were analyzed 3 hours after the stimulation. **P* < 0.05 versus no inhibitors; †*P* < 0.01. (d) Substrate-enhanced COX proteins. Western blot analysis for COX-2 and COX-1 in 3T3-L1 1-day preconfluent cells after pretreatment with 15 μ M AA.



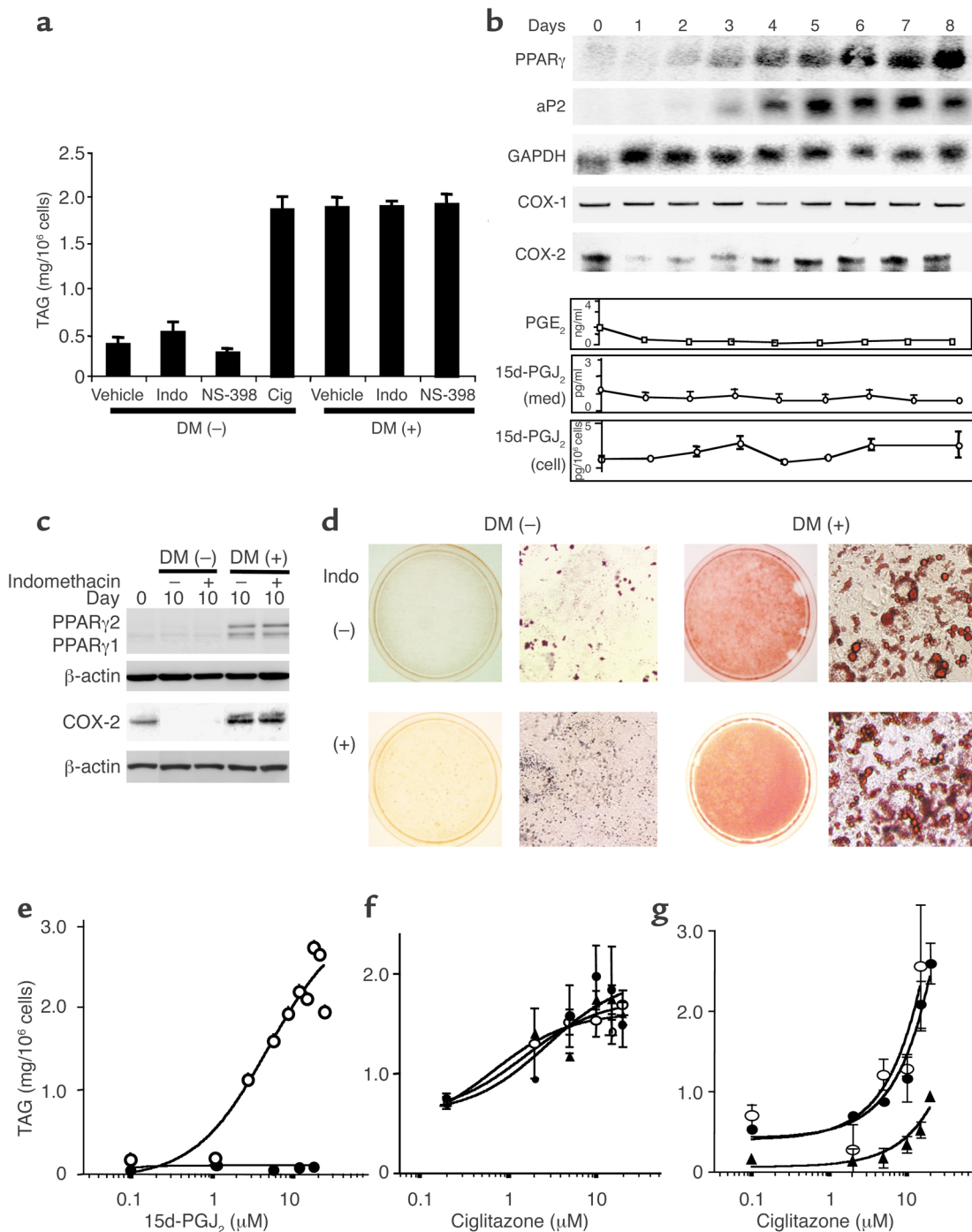


Figure 2

3T3-L1 cells were treated with DM or ciglitazone (15 μM) to initiate differentiation and were evaluated on days 8 or 10. (a) COX inhibition fails to modulate DM-induced adipocyte differentiation. The extent of TAG formation was estimated in cells differentiated with DM, ciglitazone (Cig), or DM in combination with indomethacin (3 μM) or NS-398 (10 μM). (b) PG formation is not enhanced during adipogenesis. Products during adipogenesis, as determined by Northern blotting (PPARγ, aP2, GAPDH), RT-PCR (COX-1), ribonuclease protection assay (COX-2) analysis, and LC-MS/MS (PGE₂, 15d-PGJ₂) in the medium (med) and cells. (c) Inhibition of PG formation by indomethacin fails to modulate the induction of adipocyte protein expression by DM. (d) Inhibition of PG formation fails to influence accumulation of lipid induced by DM. Oil red O staining of cells illustrates lipid accumulation in untreated cells versus cells treated with DM ± indomethacin. (e) 15d-PGJ₂ drives adipogenesis in a PPARγ-dependent manner. Cells were treated with 15d-PGJ₂ (0.1–15 μM) for 2 days in the absence (open circle) or presence (filled circle) of the PPARγ antagonist BADGE (100 μM). (f) Low concentrations of 15d-PGJ₂ do not amplify the effect of ciglitazone. Cells were differentiated with ciglitazone alone (filled circle) or in combination with 1 μM (open circle) or 0.1 μM 15d-PGJ₂ (filled triangle). (g) COX inhibition fails to modulate the enhancement of ciglitazone-dependent differentiation by DM. Cells were incubated to differentiate by adding ciglitazone alone (filled triangle) or in combination with DM (filled circle) or DM + indomethacin (open circle).

Table 1

LC-MS-MS determination of PG levels in the medium of 3T3-L1 cells differentiated with DM

Sample	PGE ₂ (ng/10 ⁶ cells)	6-keto-PGF _{1α} (ng/10 ⁶ cells)	PGF _{2α} (ng/10 ⁶ cells)	15d-PGJ ₂ (pg/10 ⁶ cells)
Preadipocytes	24.0 ± 3.7	2.4 ± 1.1	0.6 ± 0.3	77 ± 22
Mature adipocytes	3.2 ± 0.3	1.2 ± 0.3	0.8 ± 0.2	43 ± 6
DM alone	1.2 ± 0.4	0.1 ± 0.03	0.5 ± 0.1	33 ± 13

Determinations based on *n* = 3–8. Values for mature adipocytes reflect determinations made between days 7–10 of differentiation.

medium collected over 24 hours averaged 0.3 ng/ml (~0.9 nM), 0.16 ng/ml (~0.4 nM), and 1.4 pg/ml (~4 pM), respectively (Table 1).

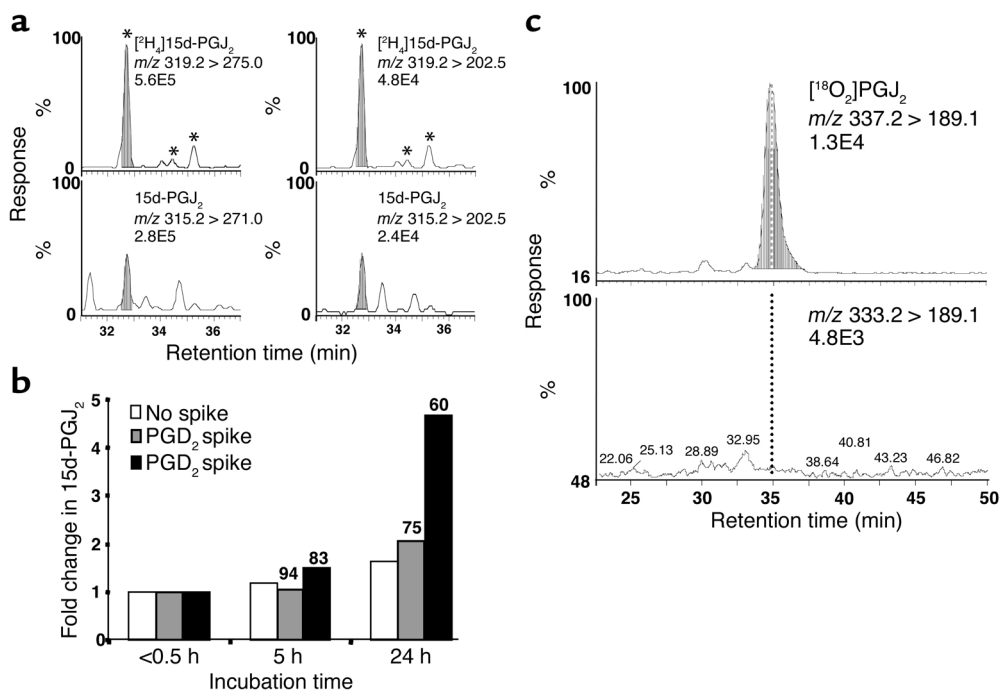
Indomethacin (3 μM) and NS-398 (10 μM), were added daily to 3T3-L1 cells along with DM and post-DM to suppress PG formation. Neither the extent nor the rate of their maturation to adipocytes induced by DM was affected, as evaluated by accumulation of TAG (Figure 2a), expression of PPARγ (Figure 2c), or total lipid accumulation (Figure 2d).

NSAIDs have been shown to bind and activate PPARα and PPARγ in vitro (43) at high concentrations. The EC₅₀ for indomethacin in a standard lipogenesis assay, using C3H10T1/2 mouse mesenchymal

stem cells, is 80 μM (43). Concentrations of indomethacin exceeding those necessary for maximal COX inhibition, drove adipocyte maturation, as reflected by TAG accumulation (data not shown). This effect was amplified in the presence of DM and was inhibited by the PPARγ specific antagonist, BADGE (not shown) (42). However, indomethacin, when used at a concentration sufficient to inhibit COX activity, did not induce significant differentiation

of 3T3-L1 cells as characterized by PPARγ expression (Figure 2c) or TAG accumulation (Figure 2d). Thus, although high concentrations of a COX inhibitor can drive differentiation in a PPARγ-dependent manner, concentrations of the same compound sufficient to inhibit completely PG production fail to induce differentiation or to modify the rate of differentiation induced by DM.

Given the very low levels of 15d-PGJ₂ that we identified in preadipocytes, we sought to characterize the concentration of added eicosanoid necessary to drive adipocyte maturation by acting as a PPARγ ligand. The EC₅₀ for response to 15d-PGJ₂, measured by TAG accumulation, was 4.5 μM and this response was antagonized by pre-

**Figure 3**

15d-PGJ₂ in human urine. (a) Mass chromatogram of 15d-PGJ₂ in human urine. Representative chromatogram obtained by NI-ESI⁺ MRM. Healthy human volunteers abstained from NSAID use for more than 5 days before urine collection. Asterisks denote isomers of [²H]₄15d-PGJ₂ present in the internal standard. The major detectable isomer is identified by shading. (b) Formation of 15d-PGJ₂ by dehydration of PGJ₂ or PGD₂. Unlabeled PGD₂ or PGJ₂ (0.2 ng/ml) was added to human urine at T = 0 hour. Aliquots (5.0 ml) were analyzed at the indicated time points for loss of spike and concomitant formation of 15d-PGJ₂. The numbers above bars represent amount (%) of added PG remaining at the corresponding time point. (c) Failure to detect endogenous PGJ₂ in freshly voided urine. Representative chromatogram of PGJ₂ in urine. NI-ESI⁺ MRM of urine samples (5.0 ml + 1.0 ng [¹⁸O₂]-PGJ₂) that were SPE extracted within 15 minutes of voiding. HPLC mobile phase was 30% CH₃CN (isocratic). Other parameters were identical to those used for 15d-PGJ₂. The PGJ₂ peak is shaded. Estimation of the threshold of PGJ₂ detection was based on the least integratable peak.

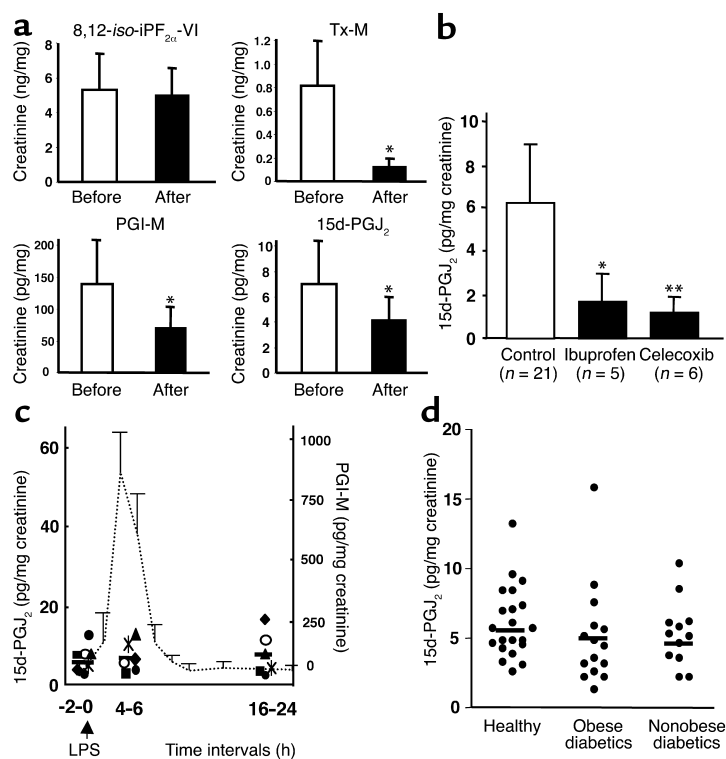


Figure 4
 Biosynthesis of 15d-PGJ₂ in humans. (a) Varied impact of COX inhibition on PG metabolites and a auto-oxidative product of arachidonic acid. Urine collections were obtained before and after aspirin (81 mg/d × 6 days) as described in the Methods section. The effects of a COX inhibitor on 15d-PGJ₂ formation are compared with markers of COX-dependent (Tx-M and PGI-M) and COX-independent, free radical catalyzed isoprostane (8,12-iso-iPF_{2α}-VI). Tx-M = 2,3-dinor TxB₂. **P* < 0.05. (b) COX-2 is the dominant source of 15d-PGJ₂ biosynthesis in humans. 15d-PGJ₂ levels in control volunteers (*n* = 21) were compared with those treated with the COX-2 selective NSAID celecoxib (*n* = 6) or nonselective ibuprofen (*n* = 5) (800 mg), administered acutely 30 minutes before urine collection. **P* < 0.05; ***P* < 0.01. (c) Biosynthesis of PGI₂, but not 15d-PGJ₂, is altered during an acute inflammatory response in humans. Volunteers (*n* = 6) received LPS (4 ng/kg). PGI-M (2,3-dinor 6-keto-PGF_{1α}) was measured at multiple time points and is plotted as the mean ± SD (dotted line). For 15d-PGJ₂ measurements, urine was collected before LPS treatment (−2 to 0 hours) and at time points corresponding to peak inflammatory (4–6 hours) and resolution (16–24 hours) phases of response. Each individual is represented by a unique symbol that is conserved across time points. Median levels are indicated by horizontal lines. (d) Biosynthesis of 15d-PGJ₂ is unaltered in diabetes or obesity. 15d-PGJ₂ levels in healthy individuals are shown in comparison to obese (BMI ≥ 30) and nonobese (BMI < 30) patients with noninsulin-dependent type 2 diabetes. Median levels are indicated by horizontal lines.

treatment with PPAR γ antagonist (Figure 2e). We also addressed the possibility that lower concentrations of 15d-PGJ₂ might enhance effects of an alternative PPAR γ agonist by ligating a coactivator molecule (44). The ability of ciglitazone to induce adipocyte maturation was unaltered by adding 15d-PGJ₂ (0.1 μ M and 1.0 μ M) (Figure 2f). Similarly, COX inhibition failed to modulate differentiation induced by ciglitazone, consistent with its failure to modulate a response to DM (Figure 2g).

In summary, 15d-PGJ₂ is formed by 3T3-L1 cells. However, despite upregulation of COX-2, synthesis of

this trivially abundant PG is unaltered during adipocyte differentiation, and suppression of its formation by COX inhibition fails to influence differentiation. Although addition of exogenous 15d-PGJ₂ can drive adipocyte maturation by ligating PPAR γ , the concentrations required (~5 μ M) exceed those of the endogenous PG released into the medium (~5 pM) and the intracellular levels (~1 nM) by orders of magnitude.

Endogenous 15d-PGJ₂ in human urine. Estimation of urinary metabolite excretion is a classical and sensitive approach to the study of PG biosynthesis (45). A representative chromatogram from human urine is shown in Figure 3a. A dominant endogenous peak at *m/z* 315.2 co-elutes from reversed phase separation with the internal standard [²H₄]15d-PGJ₂ and is associated with two independent transition ions (*m/z* 271.5 and 202.5), also characteristic of 15d-PGJ₂. Although four distinct isomers of 15d-PGJ₂ are identified in the spike, only the major isomer is detected in urine (additional urinary peaks at *m/z* 315.2 > 271.5 have not been identified, but are believed to be isoprostanes [iPs]). Urinary 15d-PGJ₂ in healthy volunteers was 6.3 ± 2.7 pg/mg creatinine (*n* = 21) with similar levels in males (6.1 ± 3.4 [*n* = 11]) and females (6.6 ± 1.6 [*n* = 10]) (*P* > 0.05). The stability of PGD₂ and PGJ₂, known to dehydrate partially to 15d-PGJ₂ in aqueous solutions, was examined by adding synthetic PGD₂ and PGJ₂ to urine and observing their disappearance over 24 hours with the concomitant formation of 15d-PGJ₂. Although both PGs are unstable in urine over 24 hours, only PGJ₂ is a significant source of 15d-PGJ₂ (Figure 3b). However, PGJ₂ was undetectable in urine, even when samples were collected and extracted into organic solvent within 15 minutes of voiding, thus excluding its rapid dehydration to 15d-PGJ₂ ex vivo (Figure 3c).

We studied the impact of a mixed COX inhibitor, aspirin (81 mg/d for 6 days), on excretion of 15d-PGJ₂, along with major urinary metabolites of thromboxane (Tx) (2,3-dinor-TxB₂) (40) and PGI₂ (PGI-M) (39) and the F₂ iP, 8,12-iso-iPF_{2α}-VI (46). Aspirin, as expected, had no effect on excretion of the free radical catalyzed iP, but depressed the Tx metabolite, which derives largely from COX-1, and to a lesser extent, PGI-M, which derives largely from COX-2. Excretion of 15d-PGJ₂ was also depressed by aspirin to an extent (mean 33%) similar to that observed with the PGI₂ (mean 44%) rather than with the Tx (mean 81%) metabolite (47) (Figure 4a). We assessed the contribution of the two COX isoforms to 15d-PGJ₂ biosynthesis by measuring the com-

Table 2

Clinical characteristic of patients and analyses of PGs in synovial fluid

Patient	Age	Sex	Diagnosis	Medication	6-keto-PGF _{1α} (pg/mg protein)	PGE ₂ (ng/mg protein)	15d-PGJ ₂ (pg/mg protein)
1	89	F	OA		ND	0.42	ND
2	89		RA		ND	4.41	ND
3	74	M	Psoriasis	Naproxen	ND	5.22	ND
4	62	M	OA	Naproxen, APAP,	ND	8.73	ND
5	58	M	Gout	APAP, allopurinol, colchicine	ND	24.20	ND
6	65	M	RA	MTX, FA	ND	24.92	ND
7	45	F	RA		ND	12.61	ND
8	61	M	Gout	APAP, allopurinol, colchicine, naproxen	3.2	46.57	1.3
9	69	M	RA	HCO, SSZ, MTX, FA	ND	7.45	ND
10	57	M	OA	Ibuprofen	ND	5.58	ND
11	48	F	RA	Codeine, MTX, HCO, APAP, etodolac	ND	9.26	ND

OA, osteoarthritis; RA, rheumatoid arthritis- APAP, acetaminophen; MTX, methotrexate; FA, folic acid; HCO, hydroxychloroquine; SSZ, sulfasalazine; ND, peak not distinguishable from background (15d-PGJ₂<5 pg/0.5 ml; 6-keto-PGF_{1α}<30 pg/ml).

parative impact of a nonspecific inhibitor (ibuprofen) with that of a selective COX-2 inhibitor (celecoxib), as described previously. Urinary excretion of 15d-PGJ₂ was depressed from 7.0 ± 2.7 to 1.7 ± 1.3 ($P < 0.05$) by ibuprofen and to 1.2 ± 0.7 ($P < 0.01$) by celecoxib. The similar effects of both drugs (Figure 4b) suggest that COX-2 is the dominant source of 15d-PGJ₂ biosynthesis under physiologic conditions in humans.

Biosynthesis of 15d-PGJ₂ in human inflammation. 15d-PGJ₂ has been proposed both as a propagator of the inflammatory response (20) and a contributor to the resolution of inflammation (21, 24). Administration of LPS to humans is associated with expression of both COX isoforms *ex vivo* and marked PG formation coincident with a flu-like response (39). This response is associated with a marked increase in urinary excretion of PGI-M, a Tx metabolite and distinct iPs, as we previously described (39). Although 15d-PGJ₂ was detectable in the urine of these individuals, there was no significant alteration in biosynthesis during or after the induction of an acute inflammatory response, despite associated expression of both COX isoforms *ex vivo* (39) and augmented biosynthesis of other PGs such as PGI-M (Figure 4c). We further assessed the role of 15d-PGJ₂ in inflammation by examining the profile of PGs present in synovial fluid of patients with arthritis (Table 2). Although PGE₂ was evident, 15d-PGJ₂ fell below the limit of detection. Finally, we sought altered biosynthesis of 15d-PGJ₂ in two conditions, type 2 diabetes and obesity, in which its role as an endogenous ligand for PPAR γ has been suggested (3, 48). Although 15d-PGJ₂ was detectable in the urine of all patients, biosynthesis of this PG was unaltered in patients with diabetes, irrespective of coincident obesity compared with healthy individuals (Figure 4d).

In summary, despite its detection at low levels in urine, alterations in the biosynthesis of 15d-PGJ₂ were not observed in diverse settings in which its role as an endogenous ligand for PPAR has been implicated.

Discussion

Interest has centered on oxidized lipids as potential agonists for nuclear receptors (49). In particular, several features of 15d-PGJ₂ have rendered it attractive as a candidate endogenous ligand for PPAR γ . First, it activates PPAR γ -dependent signaling systems more potently than other fatty acids that have been studied. Second, a representative synthetic CP-PG, Δ^{12} -PGJ₂, is actively transported into the nucleus in a time- and temperature-dependent manner with Michaelis-Menten kinetics, suggestive of carrier-mediated active transport (16, 17). Finally, formation of immunoreactive 15d-PGJ₂ has been detected during the propagation and resolution of inflammation (21) and in settings associated with PPAR γ activation (22). However, the reported concentrations of 15d-PGJ₂ necessary to activate PPAR γ (2.5–100 μ M) in such studies greatly exceed those (μ M) associated with the biologic activity of conventional PGs (28), the molecular characteristics of discrete transporters or PG-like receptors with high affinity for PGJ₂ or its 15-deoxy metabolite remain to be identified, the reported immunoreactive levels are high and the immunoassay used most commonly for detection of 15d-PGJ₂ (21, 24) has been withdrawn.

We sought to investigate the actual biosynthesis of 15d-PGJ₂ by developing a specific and sensitive approach based on mass spectrometry. We detected very small amounts of 15d-PGJ₂ formed within and released by fibroblasts *in vitro*. This was augmented by provision of arachidonate substrate, which itself upregulated expression of COX-2 (50), apparently the dominant source of 15d-PGJ₂ formation. However, the quantities of 15d-PGJ₂ formed are very small by comparison with conventional PGs (26, 27).

We focused on its role in 3T3-L1 fibroblasts, where its role as a PPAR γ ligand was first suggested (51). 3T3-L1 fibroblasts are stimulated to differentiate into adipocytes by the addition of DM or a synthetic PPAR γ agonist, such as ciglitazone. 3T3-L1 cells do

make 15d-PGJ₂, and COX-2 is among the proteins upregulated during differentiation induced by DM. However, despite increased expression of the enzyme, no differences in formation of 15d-PGJ₂ or the more abundant conventional PGs, PGE₂ and PGI₂ were observed during differentiation. Furthermore, addition of an NSAID or a selective COX-2 inhibitor, at doses shown to suppress PG formation, failed to alter induction of differentiation by either DM or ciglitazone. Consistent results (data not shown) were obtained in a second murine fibroblast cell line, 3T3-F442A. The failure of indomethacin to inhibit completely the production of 15d-PGJ₂ could represent the contribution of auto-oxidative products, as previously described (52).

Characterizing the dose-related effects of the NSAIDs was important as concentrations higher than those necessary to achieve maximal inhibition of PG formation drive adipocyte maturation by ligation of PPAR γ . As reported previously, exogenous 15d-PGJ₂ can also function as a PPAR γ agonist (41); however, the EC₅₀ for this response, 4.5 μ M, exceeds endogenous concentrations by orders of magnitude. Lower concentrations also failed to modulate the response to ciglitazone or DM, as might be expected if it bound to a PPAR γ coactivator protein (53).

The intrinsic reactivity of the highly electrophilic α,β -unsaturated ketone moiety of 15d-PGJ₂ somewhat complicates estimates of its availability in biologic systems. It readily cross-links *in vitro* with many biomolecules (23). The reactivity of the CP ring has been best characterized for another, the CP-PG, Δ^{12} -PGJ₂ (54, 55). This compound undergoes inactivating conjugation with glutathione and L-cysteine both *in vitro* and *in vivo* (54, 55). When [³H]- Δ^{12} -PGJ₂ is injected into the rat, only 10% of the infused material is recovered in the urine, whereas 90% is concentrated in the bile (55). Consistent with these observations, addition of albumin reduces the potency of exogenously added 15d-PGJ₂ to activate PPAR γ in transfected cells by roughly one order of magnitude (56).

These observations may contribute, in part, to the marked discrepancy between the concentrations of exogenous 15d-PGJ₂ necessary to activate PPAR γ -dependent differentiation and the endogenous concentrations actually detected. However, free intracellular levels remained orders of magnitude below those necessary to drive differentiation. Indeed, conjugation of 15d-PGJ₂ to glutathione only attains steady state after 2 hours *in vitro* (57), whereas PGs typically exert their actions within minutes in biologic milieu. Consistent with this discordance between *in vitro* observation and *in vivo* plausibility, glutathione depletion failed to elevate intracellular levels of 15d-PGJ₂. Furthermore, although the recovery of intracellular [³H]15d-PGJ₂ almost doubled after cleavage of potential chemical bonds with biomolecular thiols by treatment briefly with alkali (16), it remained less than 2 nM, far below that necessary to activate PPAR.

Estimation of PGs and their metabolites in urine has proven to be a sensitive approach to the detection of minor and transient alterations in their biosynthesis (58). The 15d-PGJ₂ that we detected does reflect endogenous biosynthesis of the compound. Administration of COX inhibitors depressed excretion of the PG. Furthermore, although PGD₂ can dehydrate spontaneously to 15d-PGJ₂ in urine *in vitro*, this was a most unlikely source of artifact. Rapid extraction of freshly voided urine into organic solvent, which would stabilize excreted PGJ₂, failed to permit detection of PGJ₂ in human urine. Unsurprisingly, 15d-PGJ₂ has eluded detection by less sensitive approaches using mass spectrometry in the past (27). There has been a recent brief report of its detection in human urine as a pentafluorobenzyl (PFB) derivative using GC-MS (25). However, these results are controversial. We found that a contaminating peak co-eluted with the PFB derivative of authentic 15d-PGJ₂ and resisted chromatographic separation (data not shown). The comparable effects of selective inhibitors of COX-2, traditional NSAIDs, and aspirin on urinary 15-PGJ₂, suggest that COX-2 was the dominant source of formation of this PG *in vitro* and *in vivo*.

Biosynthesis of 15d-PGJ₂ is not increased in a variety of settings in which PPAR γ activation by its endogenous ligand(s) has been implicated. We have previously reported that LPS administration to human volunteers results in transient flu-like systemic symptoms that are ameliorated by prior inhibition of COX-2 (39). Coincident with the symptoms, expression of both COX isoforms is detectable *ex vivo*, whereas biosynthesis of the conventional COX products, including TxA₂, is markedly augmented (39). Despite this evidence of marked and functionally relevant COX activation and the putative role of 15d-PGJ₂-dependent PPAR γ ligation in the propagation of inflammation (20, 59), no alteration in the small quantities of urinary 15d-PGJ₂ that we detected was observed. Similarly, despite the putative role of COX-2-derived 15d-PGJ₂ in the resolution of inflammation (60), no alteration in biosynthesis occurred as symptoms resolved. The efficacy of COX-2 inhibitors and NSAIDs in the treatment of arthritis (61, 62) attests to the relevance of COX products in the mediation of pain and inflammation in this disease. Despite the ready detection of PGE₂ in synovial fluid obtained from patients with arthritis who were not taking NSAIDs or related drugs, 15d-PGJ₂ was undetectable. Finally, PPAR γ activation by its endogenous ligand(s) has been implicated in both obesity and diabetes. We failed to detect any alteration in 15d-PGJ₂ biosynthesis in patients with diabetes, with or without concurrent obesity.

In summary, 15d-PGJ₂ is formed in small amounts by COX-2 in fibroblasts *in vitro* and is detectable as a minor product of COX-2 in human urine. It does not mediate PPAR γ -dependent adipocyte maturation. Its biosynthesis is also unaltered in diverse set-

tings of inflammation in humans in which its roles as a PPAR γ ligand and a modulator of NF- κ B signaling have been implicated.

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