

A PPAR-independent pathway to PUFA-induced COX-2 expression

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| 39 | Keywords: Arachidonic acid; Protein kinase C; Peroxisome-proliferator-activated receptor; |
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| 40 | Cyclooxygenase 2; (Bovine endometrium). |
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| 44 | |
| 45 | Summary |
| 46 | Polyunsaturated fatty acids (PUFAs) induce COX-2 in bovine endometrial stromal cells |
| 47 | through activation of peroxisome-proliferator-activated receptor alpha (PPAR $lpha$). We have |
| 48 | investigated alternative (PPAR-independent) pathways to COX-2 induction using a reporter |
| 49 | construct driven by a COX-2 gene promoter sequence lacking a PPAR response element. This |
| 50 | construct was induced by PUFAs, but not by PPAR agonists. PPAR-independent reporter gene |
| 51 | expression occurred 6 h after PPAR-dependent induction of the endogenous COX-2 gene. In |
| 52 | contrast to PPAR-dependent COX-2 induction, which is not affected by NF-κB inhibitors, the |
| 53 | PPAR-independent pathway was blocked by the NF-κB inhibitor MG132 or following deletion |
| 54 | of NF-κB sites in the COX-2 promoter. The PPAR-independent effect of PUFA was mimicked |
| 55 | by the PKC activators 4 β -PMA and prostaglandin $F_{2\alpha}$, but was not blocked by the PKC |
| 56 | inhibitor RO318425. The results demonstrate a pathway to the induction of COX-2 by PUFAs |
| 57 | requiring NF-κB but not PPAR or PKC. |

INTRODUCTION

Various hormones interact with G-protein-linked cell surface receptors to activate phospholipases and raise intracellular levels of polyunsaturated fatty acids (PUFAs). A good example is the effect of oxytocin exerted via oxytocin receptors on the epithelium of the uterine endometrium, which leads to activation of phospholipases C and A, releasing arachidonic acid and other PUFAs esterified in phospholipids (Flint et al., 1986). PUFAs produced in this way may then act as second messengers, both through activation of protein kinases and via other metabolic products (Khan et al., 1995; Maloberti et al., 2005).

PUFAs may also be secreted by the cells in which they are generated, to act on neighbouring cells as paracrine messengers (Cooke et al., 1991; Ronco et al., 2002). This has been suggested to occur in the endometrium in relation to both activation of cyclic AMP phosphodiesterase (Cheng et al., 2007) and induction of cyclooxygenase 2 (COX-2; also known as prostaglandin endoperoxide synthase 2; Sheldrick et al., 2007). Evidence for an effect on COX-2 includes the observations that PUFAs raise COX-2 levels in rat uterine stromal cells (Prigent-Tessier et al., 1996) and in bovine endometrial epithelial cells (Parent et al., 2003) through increased *COX-2* gene expression and increased levels of COX-2 transcripts (Asselin et al., 1997; Meade et al., 1999). Furthermore the oxytocin receptor is expressed in the bovine endometrial epithelium at a time when it is not expressed in the stroma (Robinson et al., 1999). Arachidonic acid has therefore been proposed to act as a paracrine signal released during the oestrous cycle by epithelial cells to induce COX-2 in the stroma (Sheldrick et al., 2007). The experiments described here were carried out to investigate the pathways by which PUFAs induce *COX-2* gene expression in endometrial stromal cells.

Previous experiments on the mechanism by which arachidonic acid raises COX-2 levels suggest that peroxisome-proliferator-activated receptors (PPARs) mediate PUFA-induced COX-2 expression in the bovine endometrium (MacLaren et al., 2006; Sheldrick et al., 2007). The isoform PPARα appears to be involved in bovine endometrial stromal cells. PPARα is activated directly by a range of PUFAs, including arachidonic acid. However, PPARα function is dependent on phosphorylation by protein kinase C (PKC; Blanquart et al., 2004), which is also activated by PUFAs (Shinomura et al., 1991; Khan et al., 1995). Furthermore, in some cells PKC induces COX-2 through a second pathway, involving NF-κB (Paik et al., 2000; Wu

| 91 | and Wiltbank, 2001; Yan et al., 2002). As a result arachidonic acid may potentially induce |
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| 92 | COX-2 directly via PPARα and indirectly through PKC/NF-κB. |
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| 94 | To distinguish between PPAR-dependent and -independent pathways we have used a reporter |
| 95 | construct under the control of a naturally-occurring sequence derived from the bovine COX-2 |
| 96 | promoter. This construct, which includes most of the important transcription regulators present |
| 97 | in the COX-2 promoter but lacks an identifiable peroxisome-proliferator response element |
| 98 | (PPRE), allows the study of pathways to COX-2 induction in which PPARs are not involved. |
| 99 | Furthermore we have investigated the role of NF-kB in the function of the promoter by |
| 100 | sequential removal of the NF-κB sites from this construct. |
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| 102 | MATERIALS AND METHODS |
| 103 | |
| 104 | Materials |
| 105 | All PUFAs; synthetic PPAR agonists; the NSAIDs, indomethacin and NS398; prostaglandin |
| 106 | A_1 (PGA ₁); the phorbol ester 4 β -PMA and its inactive analogue 4 α -phorbol 12,13-didecanoate |
| 107 | $(4\alpha\text{-PDD})$; the protein kinase C inhibitor, RO 318425, and the NF-κB inhibitor, MG132 were |
| 108 | obtained from either Sigma (Poole, UK) or Calbiochem (Nottingham, UK). Conjugated |
| 109 | linoleic acid (Sigma catalogue number O 5507) was a mixture of cis- and trans-9,11- and - |
| 110 | 10,12-octadecadienoic acids. The PUFAs, indomethacin and prostaglandins were dissolved in |
| 111 | ethanol; all other compounds were added to culture medium in DMSO. Vehicle controls were |
| 112 | used as appropriate. PUFAs were stored in ethanol under N_2 at -20 $^{\circ}$ C in darkness. |
| 113 | |
| 114 | Cell culture |
| 115 | Bovine endometrial stromal cells isolated on day 16 of the oestrous cycle (Flint et al., 2002) |
| 116 | were maintained as described previously (Sheldrick et al., 2007). To test effects of PUFAs on |
| 117 | COX-2 promoter function, the fatty acids were added to culture medium for up to 48 h after |
| 118 | transfection. |
| 119 | |
| 120 | COX-2 reporter plasmids |
| 121 | A bovine COX-2 promoter fragment was cloned by PCR on the basis of the genomic sequence |
| 122 | (GenBank accession no. AF031699; Liu et al., 1999). The PCR product was sequenced to |
| 123 | confirm its identity, and inserted into the vector pCAT promoter 3 (Promega, Southampton, |

124 UK). The resulting construct was designated pCATCOXprom. This promoter sequence did not contain a PPRE (which is further upstream of the transcription start site; Meade et al., 1999). 125 126 NF-κB mutants (Lee et al., 2001) were prepared from pCATCOXprom by site directed 127 mutagenesis using the QuikChange multi site-directed mutagenesis kit (Stratagene, Cambridge, 128 UK) and checked by sequencing. The NF-κB consensus sequence GGGAATTCCC at -472 129 (numbering from the transcription start site) and the related sequence GGAACGTCCC at -454 130 were both mutated to GGCCTTCCC in separate plasmids (designated Del I and Del II 131 respectively). A third plasmid (Del I/II) contained both mutated sequences. All deletions were 132 confirmed by sequencing. pCAT basic and pCAT control vectors (Promega) were used as 133 controls.

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Transient transfections

Stromal cells at 50-60% confluence were washed and trypsinized before resuspension in Dulbecco Modified Eagles Medium (DMEM; Sigma, Poole, UK). They were then electroporated at 300 volts, 1650 μ F with 5 μ g plasmid DNA in a volume of 0.8 ml DMEM, at a concentration of 1-3 x 10⁶ cells per ml (conditions which had been optimized in preliminary experiments). Electroporated cells were suspended in DMEM containing 1% antibioticantimycotic and 10 % fetal bovine serum previously stripped with dextran-coated charcoal (5% w/v; DMEMSS) and transferred to 24-well plates (1.0-1.5 x 10⁵ cells per well). Test compounds were then added to the medium as required, and culture continued for 36 or 48 h. For time course experiments compounds were added at various times before the end of culture; thus in the case of cells treated for 3 h, compounds were added 3 h before culture was terminated at 48 h after medium change. At the end of culture cells were washed once in 1 ml ice-cold phosphate-buffered saline containing 0.2 mM sodium orthovanadate and lysed for 20 min on ice with reporter lysis buffer (100-200 μ l; Promega) for assay of chloramphenicol acetyl transferase and protein.

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Chloramphenicol acetyl transferase assay

- 152 Chloramphenicol acetyl transferase (CAT) was measured in cell lysates by a 2 phase method.
- 153 Pre-cleared cell lysates (30 µl) were incubated at 37° C with 10 µl 5 mM chloramphenicol in
- 154 250 mM Tris containing 5 mM EDTA, pH 7.8 in 96-well plates. After 5 min, 10 μl 75 μM HCl
- 155 containing 1.5 nmol [³H]acetylcoenzyme A (Amersham, Little Chalfont, UK) was added to
- each tube. The plates were incubated at 37° C for 3 h. The reaction mixture (48 µl) was

transferred to scintillation vials containing 0.5 ml 7 M urea and 5 ml 0.8% w/v toluene-PPO was added to each vial. The vials were capped and shaken for 15 sec.

[3H]Acetylchloramphenicol extracted into the toluene out of the aqueous phase was measured by scintillation counting after 20 min.

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Immunoblotting

163 Cells were lysed and extracted for immunoblotting as described previously (Sheldrick et al., 164 2007). For immunoblotting, boiled lysates (10 µg protein) were subjected to electrophoresis on 10% acrylamide gels (5% stacking gels) before electroblotting onto Protran BA83 (Schleicher 165 166 & Schuell, Anderman and Company Kingston-upon-Thames, Surrey, UK) membranes in 25 167 mM Tris pH 8.3 containing 148 mM glycine and 20% (v/v) methanol. For detection of COX-2 168 bands (at molecular mass = 72 kDa), membranes were probed with COX-2 antibody (C-20; SC 169 1745, Santa Cruz Biotechnology, obtained through Autogen Bioclear, Calne, UK; 1:250 dilution in a solution containing 1% (w/v) Marvel milk powder and 0.5% (w/v) Tween 20 in 170 171 phosphate buffered saline). Second antibody was donkey anti-goat IgG-horseradish peroxidase (SC 2020; Santa Cruz Biotechnology; 1:11 000 dilution in 3% (w/v) Marvel, 0.5% (w/v) 172 173 Tween 20 in phosphate buffered saline), and visualization was by enhanced 174 chemiluminescence (Amersham) using Kodak BioMax Light film. Colour markers (molecular 175 masses 29 000 – 205 000; Sigma) were used to identify molecular masses of proteins. Band 176 intensities were quantified using Kodak 1D digital image analysis software.

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Protein assay

Protein concentrations in lysates prepared for electrophoresis were measured before the addition of β-mercaptoethanol and bromophenol blue using BCA reagents (Perbio Science, Cramlington, UK; Bradford, 1976).

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Experimental design and analysis

Experimental treatments were carried out in 24-well plates, with each replicate treatment of cells in at least 3 wells. Within each experiment all cells were transfected in a single batch before transfer to 24-well plates, and as a result there were no differences between wells in transfection efficiency. Data were therefore expressed in terms of cell protein, and not corrected for transfection efficiency. To account for differences in band intensity between immunoblots, all blots included 2 control samples, and all experimental treatments were related

to the control bands. All bands were included for analysis. Statistical analysis of treatment effects was by analysis of variance using Genstat version 8 (Lawes Agricultural Trust; obtained through VSN International Ltd, Hemel Hempstead, UK). Where significant effects were detected (P<0.05) individual differences were tested using a post-hoc test (Bonferroni's multiple comparison test). Values are given as means \pm s.e.m of results of separate experiments. In the figures significant treatment effects (from controls or zero-time treatments) are indicated by different letters above bars.

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RESULTS

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Structure of the COX-2 promoter

The nucleotide sequence of the bovine COX-2 gene promoter is shown in Fig. 1. PPARs form heterodimers with the 9-cis retinoic acid receptor (RXR) to activate target genes by binding to specific response elements (PPREs). PPREs contain two direct repeats of PuGGTCA half-sites separated by one base (usually A) with an additional 4 bases (usually AACT) comprising the 5' part of binding site, giving the consensus sequence AACTAGGNCAAAGGTCA (the hexameric half-sites being italicised; Ijpenberg et al., 1997). The 5' half of this response element is occupied by a PPAR, the 3' half binding the RXR. The bovine COX-2 promoter region contains a group of potential half-binding sites (Fig. 1), but the bovine equivalent of the human COX-2 PPRE, which is 3.9 kb upstream of the transcription start site (Meade et al., 1999), was not present. A sequence (AAGAGGGGAAAAGTTT) resembling the PPRE of human lipoprotein lipase (AAGAGGGGAAAGGGCA; Ijpenberg et al., 1997) is located -109 bp upstream of the translation start site in the bovine sequence. This has a perfect 5' PPAR binding motif, but differs in the last 5 nucleotides from the consensus RXR binding motif. However, as shown below, none of these motifs conferred responsiveness to PPARα in stromal cells, possibly because there was insufficient homology with actual PPREs, because there are species differences between bovine and other consensus sequences, or because of the proximity of the site at -109 to the transcription start site.

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pCATCOXprom was not responsive to PPAR activation

As anticipated due to the absence of a PPRE, and in contrast to the effects of PPAR agonists on endogenous *COX-2* gene expression (Sheldrick et al., 2007), the *COX-2* promoter was not responsive to the PPARα agonists ciprofibrate, methylclofenapate, indomethacin, NS398 or prostaglandin A₁ or to the PPARγ agonist pioglitazone (Fig. 2). Nonetheless arachidonic acid

- 224 did induce CAT expression, indicating the presence of a PPAR-independent pathway for COX-
- 225 2. The response to PUFA varied between experiments in the range 2.0- to 16.2-fold (mean, 6.1-
- fold; N = 10). In previous experiments measuring endogenous COX-2 protein, ciprofibrate was
- as effective a PPAR agonist as WY14643, suggesting that the failure to induce CAT expression
- was unlikely to reflect poor bioavailability.

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Time course of effects of PUFAs on CAT expression

- Further evidence that the effect of arachidonic acid (Fig. 2) was not due to activation of a
- 232 PPAR was obtained from time course studies. At concentrations above 10 μM, arachidonic
- acid increased CAT expression between 12 and 24 h after addition (Fig. 3a). Similar responses
- were observed with n-6 PUFAs other than arachidonic acid (Fig. 3b), although these were
- delayed until 24 36 h. In contrast, induction of the endogenous *COX-2* gene (as indicated by
- 236 immunoblotting) peaked at 6 h after exposure to agonist (Fig. 3a; data of Sheldrick et al.,
- 237 2007). The lag before expression of CAT relative to induction of the endogenous COX-2
- 238 protein did not reflect the time taken by the cells to recover from electroporation, as the PPAR
- agonist indomethacin raised COX-2 levels within 6 h in cells that had been electroporated (Fig.
- 240 3c).

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Involvement of PKC and NF-κB in induction of CAT expression

- To further differentiate between PPAR-dependent and -independent effects, two experimental
- strategies were used to define the roles of PKC and NF-κB in pathways to COX-2 induction.
- 245 Firstly, cells transfected with pCATCOXprom were treated with (i) 4β-PMA (which
- specifically activates PKC; Nishizuka, 1984), (ii) prostaglandin F_{2α} (which activates PKC
- 247 though the FP receptor and phospholipase C; Abayasekara et al., 1993), (iii) the PKC inhibitor
- 248 RO318425, or (iv) the NF-κB inhibitor MG132 (Wu et al., 2004). Secondly, cells were
- transfected with constructs derived from pCATCOXprom but lacking NF-κB sites.

- Expression of pCATCOXprom was induced by 4β -PMA and prostaglandin $F_{2\alpha}$ (Fig. 4a). When
- 252 these compounds were added with arachidonic acid the effects of arachidonic acid and 4β-
- 253 PMA or arachidonic acid and prostaglandin $F_{2\alpha}$ were additive but not synergistic (i.e. there was
- 254 no statistically significant interaction). The inactive phorbol ester 4α -phorbol 12,13-
- 255 didecanoate (2 μM) was ineffective. Consistent with a role for NF-κB, the effects of
- arachidonic acid and 4β-PMA were both blocked by the NF-κB inhibitor, MG132 (Fig. 4b).
- MG132 also blocked induction of CAT expression by $PGF_{2\alpha}$ (data not shown) and reduced

basal CAT expression when added alone. In contrast to the effect of MG132, the specific PKC inhibitor RO318425 blocked the response to 4β -PMA (Fig. 4c) but not the response to arachidonic acid (Fig. 4d).

In experiments to test the effects of PUFAs on the COX-2 promoter following NF- κ B site deletion, arachidonic acid failed to induce CAT expression in any of the three NF- κ B mutant constructs (Fig. 4e). Similar results were obtained with 4 β -PMA and prostaglandin F_{2 α}, and with dihomogammalinolenic, linoleic and conjugated linolenic acids (data not shown). As expected on the basis of the effect of MG132 observed in Fig. 4b, removal of the NF- κ B sites from pCATCOXprom reduced basal CAT expression by the constructs to below 1% of the control level.

DISCUSSION

Previous experiments showed that one pathway to induction of COX-2 in uterine stromal cells treated with arachidonic acid involves PPARα (Sheldrick et al., 2007). However in previous experiments it was not possible to distinguish between an effect of arachidonic acid as a PPAR agonist and that involving activation of PKC and NF-κB. Inhibitors were used to block either PKC or NF-κB, but blocking PKC led to inhibition of the PPAR pathway (as expected based on the requirement for PPARs to be phosphorylated by PKC; Blanquart et al., 2004). Furthermore the PPARα antagonist MK886 was ineffective in uterine stromal cells (Sheldrick et al., 2007). As a result it was difficult to identify PPAR-independent pathways. This difficulty has been avoided in the present experiments by using a reporter construct which is unresponsive to PPARs. As a PPAR-responsive reporter construct was not available (the PPRE not being identified to date in the bovine gene), endogenous COX-2 expression acted as a positive control.

The results reported here confirm the involvement of PPARs in COX-2 induction, as the synthetic PPAR agonists ciprofibrate, methylclofenapate, indomethacin and NS398, which induce the endogenous gene (Fig. 3c; Sheldrick et al., 2007), did not induce the PPRE-negative CAT construct. However the data revealed at least one PPAR-independent pathway for induction of *COX-2* promoter function by PUFAs, as the *COX-2* promoter construct used was induced by arachidonic acid and other unsaturated fatty acids (Fig. 3a and b). This was confirmed by the time course of CAT induction by PUFAs (occurring at 12 - 48 h; Fig. 3), which differed from that of the endogenous *COX-2* gene observed by immunoblotting (which

| 292 | peaked at 6 h). The delay in expression of CAT by PUFAs was not due to damage caused to |
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| 293 | the cells during electroporation, as expression of COX-2 from the endogenous gene was |
| 294 | induced in electroporated cells at 6 h by indomethacin, which has previously been shown to be |
| 295 | an effective PPAR agonist. |
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| 297 | One potential PPAR-independent route to COX-2 induction involves PKC and NF-κB (Scheme |
| 298 | 1). All three PKC activators tested here: PUFAs (Shinomura et al., 1991; Khan et al., 1995), |
| 299 | 4β-PMA (Nishizuka, 1984) and prostaglandin $F_{2\alpha}$ (Abayasekara et al., 1993) induced CAT |
| 300 | expression, and each of these effects was blocked by the NF-κB inhibitor, MG132. |
| 301 | Furthermore the role of NF-kB was confirmed by the deletion studies which showed that |
| 302 | induction of the COX-2 promoter by arachidonic acid, 4β -PMA or prostaglandin $F_{2\alpha}$ depended |
| 303 | on the presence of an NF-κB responsive element. These findings are consistent with pathways |
| 304 | c, d, e and f in Scheme 1. |
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| 306 | The effects of arachidonic acid and 4β-PMA on CAT expression were also prevented by the I- |
| 307 | κB kinase inhibitor parthenolide (4 μM ; data not shown). Parthenolide maintains I- κB in the |
| 308 | non-phosphorylated state, and as a result inhibits translocation of NF- κB to the nucleus (Bork |
| 309 | et al., 1997). Its action is therefore similar to that of MG132, which reduces the rate of I - κB |
| 310 | degradation. However, in contrast to the effect of MG132, parthenolide increased basal CAT |
| 311 | expression. Among the non-specific effects of sesquiterpene lactones such as parthenolide is |
| 312 | the inhibition of 5-lipoxygenase (Capasso, 1986; Sumner et al., 1992). As lipoxygenase |
| 313 | products down-regulate COX-2 (Ait-Said et al., 2003), and the lipoxygenase inhibitor |
| 314 | nordihydroguaiaretic acid increases COX-2 levels in stromal cells (E.L.R. Sheldrick, |
| 315 | unpublished work), an increase in CAT expression in response to parthenolide may reflect a |
| 316 | side effect of this compound. |
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| 318 | The NF-κB inhibitor MG132 decreased CAT expression dramatically, and deletion of NF-κB |
| 319 | sites from pCATCOXprom had the same effect. These effects reflect the importance of NF- κB |
| 320 | in regulating COX-2 promoter function. A reduction in basal COX-2 promoter function also |
| 321 | follows NF-κB site mutation in human endometrial stromal cells (Tamura et al., 2002) and |
| 322 | alveolar epithelial cancer cells (Huang et al., 2003). The reduced level of CAT in these |
| 323 | experiments suggests that a small increase in reporter expression in response to arachidonic |

acid or 4β -PMA would have been detectable, had there been an alternative activatory site. The

double mutant was less active than either of the single mutants, but was not induced significantly by PUFA.

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Use of the PKC inhibitor RO318425 showed that although the effect of 4 β -PMA (Scheme 1, pathway e, f) was dependent on PKC, the effect of arachidonic acid was not. This allowed the effect of arachidonic acid to be dissociated from PKC, revealing a PKC-independent pathway to NF- κ B activation by arachidonic acid (pathway g in Scheme 1). The existence of a PKC-independent pathway is consistent with the additive effects of arachidonic acid and PKC activators (Fig. 4a). Potential intermediates in this pathway include reactive oxygen species, which are involved in the activation of NF- κ B by PUFAs with a time course consistent with that observed here (24 – 48 h; Mazière et al., 1999; Bécuwe et al., 2003). The effect of arachidonic acid was blocked by inhibition of NF- κ B (Fig. 4b), ruling out an arachidonic acid-stimulated pathway to COX-2 induction independent of both PKC and NF- κ B.

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Certain isoforms of PKC are down-regulated following activation, leading to refractoriness (Standeart et al., 1993). In separate studies we have confirmed that this occurs in bovine endometrial stromal cells exposed to 4β-PMA for 24 h, as shown by a reduction in phosphorylation by PKC of cyclic AMP phosphodiesterase (Cheng et al., 2007). PKC would therefore be expected to be down-regulated following activation by arachidonic acid. On this basis, if the PPAR-independent induction of CAT expression by arachidonic acid required PKC activation, the effect might be expected to be lost after 24 h. The fact that it was not (as the effects of PUFAs were only observed at or after 24 h; Fig. 3) at first sight confirms the suggestion that this effect is not exerted via PKC. However this conclusion is difficult to reconcile with the action of 4β-PMA, which increased CAT expression at 48 h, the effect being blocked by RO318425 (Fig. 4c). One possible explanation is that the isoform of PKC responsible for CAT induction by 4β-PMA was resistant to down-regulation by phorbol ester. There is a precedent for this in that PKC isoforms differ in the speed and extent of their downregulation by phorbol esters in a cell-specific manner (Adams and Gullick, 1989; Standaert et al., 1993). There are therefore 3 reasons for concluding that PKC is not involved in the response to arachidonic acid: (i) the response is not blocked by RO318425; (ii) the responses to arachidonic acid and 4β -PMA or prostaglandin $F_{2\alpha}$ are additive, and (iii) the effect occurs at a time when PKC would be expected to be down-regulated. In the light of these findings we conclude that arachidonic acid activates COX-2 promoter function in the absence of a PPRE,

| 358 | demonstrating the existence of a PPAR-independent route to COX-2 expression, and that NF- |
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| 359 | κB (but not PKC) is involved in this response. |
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| 364 | We thank Pat Fisher for isolating the endometrial stromal cells. This work was funded by the |
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| 514 | Figure Legends |
|-----|---|
| 515 | |
| 516 | Figure 1. (a) Fragment of bovine COX-2 gene promoter. Numbering as given in GenBank |
| 517 | accession no. AF031699 at left, from the transcription start site (TSS) at right. Throughout this |
| 518 | paper, numbering is from the transcription start site. Block arrows mark transcription and |
| 519 | translation start sites (indicated by TSS and coding sequence respectively). Thin arrows mark |
| 520 | PCR primers used to produce the insert in pCATCOXprom. Half sites of native PPREs (after |
| 521 | Ijpenberg et al., 1997) are underlined and highlighted. The almost perfect match to the native |
| 522 | PPRE present in the human lipoprotein lipase promoter at position -10993 of the bovine |
| 523 | COX-2 promoter sequence is highlighted (13 out of 17 nucleotides being identical). The NF-κB |
| 524 | sites I and II are in bold and underlined. (b) Mutations introduced in both NF-κB sites (Lee et |
| 525 | al., 2001). |
| 526 | |
| 527 | Figure 2. CAT expression in stromal cells transfected with pCATCOXprom and treated with |
| 528 | PPAR agonists. Cells transfected by electroporation were cultured for 36 h with or without the |
| 529 | following PPAR agonists which have previously been shown to raise endogenous COX-2 |
| 530 | levels in stromal cells (Sheldrick et al., 2007): ciprofibrate (50 μ M), methylclofenapate (50 |
| 531 | $\mu M),$ indomethacin (10 $\mu M),$ NS398 (10 $\mu M)$ and prostaglandin A_1 (3 $\mu M). The PPAR\gamma$ |
| 532 | agonist pioglitazone (50µM) has previously been shown to be ineffective. Arachidonic acid, |
| 533 | but none of the other PPAR α agonists, induced CAT expression. |
| 534 | |
| 535 | Figure 3. Time course of induction of CAT expression by arachidonic acid and other PUFAs. |
| 536 | (a) Stromal cells transfected with pCATCOXprom were cultured in the presence of arachidonic |
| 537 | acid (50 μM) for up to 48 h. Closed bars, CAT expression; open bars, endogenous COX-2 |
| 538 | determined by immunoblotting in non-transfected cells (data of Sheldrick et al., 2007). COX-2 |
| 539 | levels were raised at 3 – 12 h, whereas CAT expression was raised only after 12 h. Inset shows |
| 540 | representative immunoblotting signals for COX-2. (b) Specificity of CAT induction by PUFAs. |
| 541 | Stromal cells transfected with pCATCOXprom were cultured in the presence or absence of |
| 542 | dihomogammalinolenic acid (DGLA; 50 μM), conjugated linoleic acid (CLA; 50 μM) or |
| 543 | linoleic acid (LA; 2 $\mu M)$ for up to 48h. The concentration of linoleic acid tested, 2 $\mu M,$ was |
| 544 | optimal in dose response experiments (data not shown). (c) Electroporated cells remained |
| 545 | responsive to PPAR ligand. To test whether the failure of PPAR ligands to induce COX-2 |
| 546 | promoter function at 6 h (Figs. 2 and 3a) reflected damage to cells caused by electroporation, |

547 cells were cultured with the PPAR ligand indomethacin with (Electroporated) and without (Control) electroporation. COX-2 levels were measured by immunoblotting after 6 h treatment. 548 549 Indomethacin increased COX-2 levels both before and after electroporation, showing that 550 expression from the endogenous COX-2 gene at 6 h was not prevented by electroporation. 551 552 Figure 4. Induction of CAT expression by arachidonic acid, 4β -PMA and prostaglandin $F_{2\alpha}$. 553 Stromal cells transfected with pCATCOXprom (or related NF-kB deletion mutants) were 554 cultured with arachidonic acid (50 μM) for 48 h with or without the PKC activators 4β-PMA (2 μM) and prostaglandin $F_{2\alpha}$ (3 μM), the PKC inhibitor RO318425 (500 μM), the NF- κB 555 556 inhibitor MG132 (30 μM) or the inactive phorbol ester 4α-phorbol 12,13-didecanoate (2 μM). 557 (a) CAT expression increased in response to arachidonic acid, 4β -PMA and prostaglandin $F_{2\alpha}$, and the effects of arachidonic acid and either 4 β -PMA or prostaglandin $F_{2\alpha}$ were additive. 4α -558 559 Phorbol 12,13-didecanoate (4α-PDD) had no effect. (b) The effects of both arachidonic acid 560 and 4β-PMA were prevented by MG132. Note that CAT expression was decreased by MG132 561 added alone. (c) The effect of 4β-PMA was prevented by the PKC inhibitor RO318425, but (d) the effect of arachidonic acid was not. (e) The response to arachidonic acid was blocked by 562 563 deletion of either of the NF-κB sites in the reporter plasmid pCATCOXprom. Del I, -472/-463 mutation; Del II, -454/-445 mutation; Del I/II, mutations at both -454/-445 and -472/-463. 564 565 Basal CAT expression in mutation Del I/II was lower than for the other mutant constructs, but 566 the effect of arachidonic acid was not statistically significant. 567 Scheme 1. Scheme illustrating possible pathways by which arachidonic acid (AA) induces 568 COX-2 expression (or, in reporter experiments, expression of chloramphenicol acetyl 569 570 transferase, CAT) in stromal cells. AA induces expression of the endogenous COX-2 gene 571 through activation of PPARa (a) (Sheldrick et al., 2007). Protein kinase C (PKC) activates 572 PPARα through phosphorylation (b) (Blanquart et al., 2004). These pathways are shown as 573 dotted lines because the COX-2 promoter construct used in the present experiments to express 574 CAT lacked a peroxisome-proliferator-activated receptor response element, and did not 575 respond to PPARα activation. PKC may be activated directly by AA (c), indirectly by 576 prostaglandin $F_{2\alpha}$ synthesised from AA via the FP receptor and phospholipase C (d) or by the 577 phorbol ester 4β-PMA (e). PKC induces COX-2 via NF-κB (f). The experiments reported here 578 suggest a third pathway by which AA induces COX-2 via an unspecified intermediate X acting

through NF-κB (g). We have not attempted to identify isoforms of PKC or NF-κB, or

| 580 | intermediates | between F | PKC and NF | -κB. After | Sheldrick et a | 1. (2007) | . which | provided | evidence |
|-----|---------------|-----------|------------|------------|----------------|-----------|---------|----------|----------|
| | | | | | | | | | |

for pathways a - f, but not for pathway g.

582

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Figure 1
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9
     (a)
                                                                 PCR F primer
      241 ttttttgggg ggtggggag gggggtggca tttcaccttc agacacccct cccgactaca -1350
10
      301 agacagaatc aaacaaatgc aaaagatgat gaaattgaaa attgtcaaaa tcacccttcc -1290
11
      361 ttcatatcca ttccatccga attcctcatc aaattacctt ccaaaatgag cctagaaaac -1230
12
      421 tggccctaat aaacccagac aactgaaaag tgagtgtaat gttgtacttt aatcctttgt -1170
13
      481 cacaactcat tatcctagag aagtaggcag aaaacatgca cacaaaccaa tctttaaaaa -1110
14
      541 tetaetteet teaattggta teteetgtga aaagetggeg accaaaacag tgaaaateae -1050
15
      601 tgcagcagag aaagaaatgc cataaagcag atcttttgga cggtgagcat ccttaaacat -990
16
      661 cccggtcacc tccgcttcac ttgttcatgg gaggagaggt tacagcggtg acccctcaat -930
17
      721 ctcagattga ctatttacag gataagtgct cagggtcagt tttaggggga ggaatttacc -870
18
      781 tttatttcct ctctccaaga agcaacgtcg gggtggtggc atgaagaaca gtatttttta
19
      841 tgttgaaaac aacttagcta gaaaaagaac cttacaggta cgtatacact gactactgct
20
      901 attacagttc attcggtcaa gtaaaaaaat aaaaaaatta aagtattaat aaaaaatatt
21
22
      961 aaagtattaa taaaaaatta aaaattaatg gatgtgcgta catctacacg aggttaagag
     23
     1081 cttggcccga ttctccccag tctctccaga ctgtgacttc ctcggccccc aaaagccgag
                                                                              -510
24
25
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27
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29
30
31
32
33
                                                     NF<sub>K</sub>B I
     1141 ggaccagacg tagggggacg ggggtggggg cgcgcgaggg aattccctgc gctccggaac
                                                                              -450
     1201 gtcccagccc cctgatcaat tccgggagag gaagcaagcc aggaggccgc cctccccacc
                                                                              -390
                                                                              -330
     1261 cacgccctca cccagaaaat ctacctcgga ccccctctcg gaagcctcgg tcccaaccgc
     1321 gacagaagga aaggaatctg gcgcgaagcc ggcgccagcg gcggagaaac cggccgaggg
     1381 gcgctggaag gtctgaactg gaagggggcg atcctctttg cacccaccgt aggaagccaa
                                                                              -210
     1441 gcggcgactc cccaagtttc cgattttcta gtttgcgtag ctaaaaaaaa aagaaagaaa
                                                                              -150
     1501 tectacecca accegggtet tgegcaattg tttaagtaga aagaggggga aaagttttgg
     1561 aagggggga ggaaaggcgg aaagaaacag tcatgccgtc acgtgggcta tttcacgcat
                                                                               -30
                                         TSS
34
     1621 aaaaggaag\mathbf{g} teeteteegt tagetteea\mathbf{g} \mathbf{t}\mathbf{t}gteaaagg actegeagtg aacgteagag
                                                                               +31
35
36
37
                     PCR R primer
                                                                               +91
     1681 gacgcccgg aactccgcac cgccctcctc cggccccgca gctccgatcc gcgcacctcc
38
                                                  coding sequence
39
     1741 acgcctccgc cggccccgcg ccccgcccgc agccgagatg ctcgcccggg ccctgctgct
40
41
42
43
44
     (b)
45
                                5'-NF-κB
                                                       3'-NF-κB
46
                     cgcgcgagggaatt<u>ccc</u>tgc gctccggaacgtcccag
47
                                  48
                     cgcgcga.ggccttccctgc gctccgg.ccgtcccag
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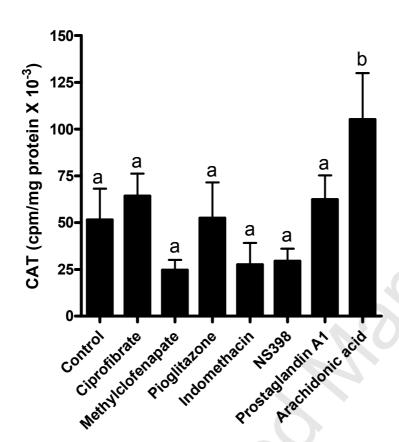


Figure 3

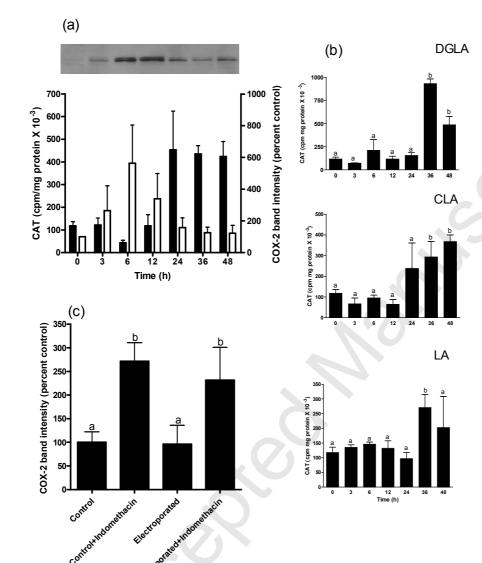
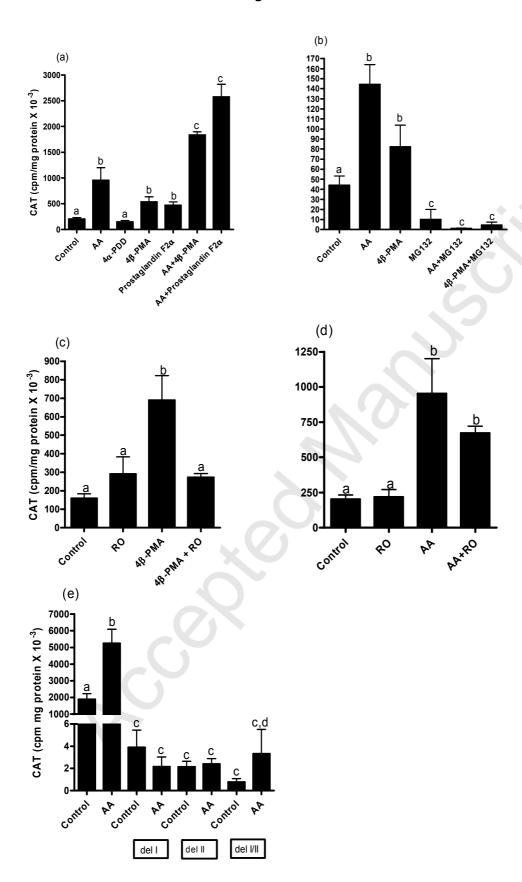


Figure 4



Scheme 1

