

PPAR α Ligands Reduce PCB-Induced Endothelial Activation: Possible Interactions in Inflammation and Atherosclerosis

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Abstract Exposure to polychlorinated biphenyls (PCBs) can activate inflammatory responses in vascular endothelial cells. Activation of peroxisome proliferator-activated receptors (PPARs) by nutrients or synthetic agonists has been shown to block pro-inflammatory responses both in vitro and in vivo. Here we demonstrate that activation of PPAR α by synthetic agonists can reduce 3,3',4,4'-tetrachlorobiphenyl (PCB77)-induced endothelial cell activation. Primary vascular endothelial cells were pre-treated with the PPAR α ligands fenofibrate or WY14643 followed by exposure to PCB77. PPAR α activation protected endothelial cells against PCB77-induced expression of the pro-inflammatory proteins vascular cell adhesion molecule-1 (VCAM-1), cyclooxygenase-2 (COX-2), and PCB77-induced expression and activity of the aryl hydrocarbon receptor (AHR) responsive cytochrome P450 1A1 (CYP1A1). Furthermore, basal AHR expression was downregulated by fenofibrate and WY14643. We also investigated the possible interactions between PCBs, and

basal PPAR activity and protein expression. Treatment with PCB77 significantly reduced basal mRNA expression of PPAR α and the PPAR responsive gene CYP4A1, as well as PPAR α protein expression. Also, PCB77 exposure caused a significant decrease in basal PPAR-dependent reporter gene expression in MCF-7 cells. Overall, these findings suggest that PPAR α agonists can reduce PCB77 induction of endothelial cell activation by inhibition of the AHR pathway, and that coplanar PCB induced pro-inflammatory effects could be mediated, in part, by inhibition of PPAR α expression and function.

Keywords Atherosclerosis · Vascular endothelial cells · PPAR α · PPAR γ · PCB · AHR · CYP1A1 · VCAM-1 · COX-2 · Inflammation

Introduction

There is substantial evidence from epidemiological studies that cardiovascular diseases are linked to environmental exposure to halogenated aromatic hydrocarbons (HAHs), such as dioxins and polychlorinated biphenyls (PCBs). For example, there was a significant increase in mortality from cardiovascular diseases among Swedish capacitor manufacturing workers exposed to PCBs for at least 5 years [1], and elevated plasma lipids and atherosclerotic plaques have been observed among dioxin exposed workers from the former Czechoslovakia [2]. A higher incidence of cardiovascular diseases, specifically chronic ischemic heart disease and chronic rheumatic heart disease, was detected in studies on the dioxin-exposed population of Seveso, Italy, after an industrial accident in 1976 [3, 4]. Furthermore, a recent study reported increased hospitalization rates for coronary heart disease in populations residing near areas contaminated with persistent organic pollutants [5].

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Atherosclerotic lesions are thought to be initiated by vascular endothelial cell activation [6]. Since the endothelium is in immediate contact with the blood, endothelial cells are particularly susceptible to the effect of pro-inflammatory agents, such as cytokines and environmental contaminants present in the bloodstream [7]. We have previously reported that coplanar PCBs such as PCB77 can activate the aryl hydrocarbon receptor (AHR), the pro-inflammatory transcription factor nuclear factor kappa B (NF- κ B), increase oxidative stress, and induce expression of adhesion molecules and cytokines such as vascular cell adhesion molecule-1 (VCAM-1) and interleukin-6 (IL-6) in vascular endothelial cells [8, 9]. Increased expression of adhesion molecules and cytokines play a critical role in endothelial cell activation and vascular disease [10, 11]. Furthermore, cell culture studies using U937 macrophages have shown that exposure to dioxin, can increase the expression of matrix degrading metalloproteinases, cyclooxygenase-2 (COX-2), and interleukin 1 β , and promote foam cell formation [12]. The expression of adhesion molecules and cytokines further enhances the inflammation by recruiting monocytes, and facilitating their binding to and migration through the endothelium. This loss of the endothelial barrier function can increase the accumulation of foam cells in the subendothelial space, subsequently leading to the formation of fatty streaks and advanced atherosclerosis [7, 13]. Studies with both in vivo and in vitro models have shown that many of the pro-inflammatory effects induced by coplanar PCBs are dependent on AHR function [14, 15]. In contrast to dioxin or dioxin-like compound-induced toxicity through abnormal or prolonged AHR activation [16, 17], it has been proposed that intervention using nutritional or pharmaceutical AHR inhibitors could potentially reduce toxicity caused by xenobiotic AHR ligands [18].

The peroxisome proliferator-activated receptors (PPARs) are ligand-activated transcription factors that form part of the nuclear receptor superfamily [19]. There are three genes that belong to the PPAR family: PPAR α , PPAR γ , and PPAR β/δ . Upon activation, PPARs control gene expression by forming a heterodimer with the retinoid \times receptor that can recognize specific DNA sequences known as PPAR response elements (PPREs) [20]. PPAR α expression is localized in tissues with a high metabolic rate (e.g., liver, kidney, and skeletal muscle), and in various cell types in the vascular wall such as smooth muscle cells, endothelial cells, and monocytes/macrophages [21]. Nutritional and pharmaceutical PPAR α agonists have been shown to be protective against atherosclerosis by down-regulating underlying pro-inflammatory signaling pathways [22, 23]. PPAR α has been shown to negatively interfere with NF- κ B, and activator protein-1 (AP-1) signaling pathways [24], and can therefore prevent the

induction of inflammatory genes such as adhesion molecules and cytokines controlled by these transcription factors. Indeed, clinical and experimental evidence suggests that PPAR α activation decreases the incidence of cardiovascular diseases [21, 23].

The objective of the current study was to determine interactions between the PPAR α and AHR pathways in the vascular endothelium. We hypothesized that PPAR α agonists can block PCB77 activation of pro-inflammatory responses in endothelial cells and that PCBs could alter basal PPAR α function. Our results suggest that PPAR α can interfere with PCB77 activation of the AHR pathway, and that PCBs can reduce function and expression of PPAR α . Since PPAR α plays a critical role in inflammation, the negative PCB effects on PPAR α expression and function may increase susceptibility to other kinds of pro-inflammatory stimuli in exposed individuals.

Materials and Methods

Cell Culture and Experimental Media

Endothelial cells were isolated from porcine pulmonary arteries, as described earlier [25]. Arteries obtained during routine slaughter were donated from the College of Agriculture, University of Kentucky. Endothelial cells from passages 5–10 were used in the current experiments. MCF-7 cells, stably transfected with a luciferase gene driven by a triple repeat of the PPAR response element (PPRE), were utilized for selected experiments. Cells were subcultured in medium 199 (endothelial cells) and DMEM (MCF-7 cells) containing 10% (v/v) fetal bovine serum (FBS, HyClone Laboratories, Logan, UT, USA) using standard techniques.

The experimental media contained 1% (v/v) FBS. PCB 77 (0.001–3.4 μ M) and PPAR α agonists (10–20 μ M) were added from stock solutions prepared in dimethyl sulfoxide (DMSO; Sigma, St. Louis, MO, USA). The PPAR α agonists' fenofibrate and WY14643 were purchased from Sigma (St. Louis, MO, USA) and Cayman Chemical (Ann Arbor, MI, USA), respectively. These two different agonists were used to verify that the observed effects were PPAR α specific. PCB77 was kindly provided by Dr. Larry W. Robertson, University of Iowa. PCB77 was used at a concentration, which was previously shown to promote endothelial cell activation, and is based on previously reported serum concentrations occurring after acute exposure [8, 9, 26]. All treatment groups contained an equal amount of DMSO. The final DMSO concentration in the media never exceeded 0.05% (v/v) in all treatment groups. Cells were treated with PPAR α agonists and with PCBs for 6–18 h.

CYP1A1 Activity: Ethoxyresorufin-*o*-Deethylase (EROD)

Cellular cytochrome P450 1A1 (CYP1A1) activity or ethoxyresorufin-*o*-deethylase (EROD) activity was measured in 48 well plates (Costar, Corning Incorporated, NY, USA) using 7-ethoxyresorufin (Sigma, St. Louis, MO, USA) as a CYP1A1 substrate, as described previously [14, 27]. Briefly, cells were rinsed with phosphate buffered serum (PBS; 37°C) followed by the addition of 7-ethoxyresorufin (2 μ M final concentration). The reaction was measured using a Cytofluor 4000 (PE Biosystems, Foster City, CA, USA) containing excitation and emission filters for 530 and 590 nm, respectively.

Measurement of VCAM-1, COX-2, CYP1A1, AHR, β -actin, and PPAR α protein expression

Total cellular protein was extracted, as described previously [8]. Cell monolayers were scraped and washed in cold PBS, pelleted, and incubated in 110 μ l of lysis buffer containing 20 mM Tris, 150 mM NaCl, 1 mM EDTA, 0.5 mM dithiothreitol, 0.5 mM phenylmethanesulfonyl fluoride (PMSF), 1 μ g/ml leupeptin, 1 μ g/ml pepstatin, 0.1% nonidet P-40, and 0.5% Triton X-100. Protein extracts were electrophoresed on 12% SDS-polyacrylamide gels transferred to nitrocellulose membranes. Proteins were probed with commercial rabbit and goat antibodies for VCAM-1, COX-2, CYP1A1, AHR (goat polyclonal IgG isotype, Santa Cruz Biotechnology, Santa Cruz, CA, USA), PPAR α (rabbit polyclonal IgG isotype, Cayman Chemical, Ann Arbor, MI, USA), and β -actin (rabbit polyclonal IgG, Sigma, St. Louis, MO, USA). β -actin was used as a loading control for normalizing expression of proteins of interest. Antibodies were diluted 1:1,000 or 1:3,000 in blocking buffer. Blots were developed with an Image Station 2000R (KODAK Molecular Imaging Systems, New Haven, CT, USA) using ECL (GE Healthcare, Piscataway, NJ, USA) for chemiluminescent detection.

PPAR α and CYP4A1 mRNA Expression Real Time Polymerase Chain Reaction (PCR)

RNA was extracted using TRIzol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's directions. Specific primers for porcine CYP4A1, PPAR α , and β -actin mRNAs were designed using Primer Express 3.0 software (Applied Biosystems, Foster City, CA, USA). The following primers were employed; CYP4A1 forward: 5' GGA ATC ATC CTC TCG CTC TTCA 3', reverse: 5'

GCA GAA CCC GGT GCA AAC 3', PPAR α forward: 5' TCG CCA TGC TGT CCT CTG T 3', reverse: 5' AAG GGT TTC CTC AGG CTC TTG 3', β -actin forward: 5' TCA TCA CCA TCG GCA ACG 3', reverse: 5' TTC CTG ATG TCC ACG TCG 3'. Real time PCR was conducted with the 7300 Real Time PCR System (Applied Biosystems) and using Sybr Green (Applied Biosystems) to measure gene expression according to manufacturer instructions. The CYP4A1 and PPAR α expression data obtained for individual samples were normalized to the corresponding β -actin expression.

PPAR Reporter Gene Studies

Since primary endothelial cells are difficult to transfect with high efficiency, a well-established model, previously used to study PPAR function, was used to study PPAR-dependent reporter gene expression [28]. The human breast cancer epithelial cell line MCF-7 was transfected with pGL3 vector (Promega, Madison, WI, USA) containing a triple repeat of PPRE in the promoter region driving luciferase gene expression and a HSV-TK-driven renilla. Transfected cells were selected using G418. Cells were plated in 48 well plates and exposed to PCBs 6 h prior to cell lysis. Cell lysis and reporter gene assay was performed using the dual reporter assay kit (Promega, Madison, WI, USA). Values were expressed as a luciferase/renilla ratio.

Statistical Analysis

All experiments were performed in triplicate. Comparisons between treatments were made by one-way or two-way ANOVA. Post-hoc comparisons of the means were made by Tukey tests. Statistical probability of $P < 0.05$ was considered significant. All statistical analyses were performed with Sigmastat (Systat Software, San Jose, CA, USA).

Results

PPAR α Agonists Downregulate the PCB77-Induced Expression of the Pro-inflammatory Proteins COX-2 and VCAM-1

Previous studies have demonstrated that exposure to coplanar PCBs and halogenated aromatic hydrocarbons (HAHs) can lead to increased expression of pro-inflammatory genes associated with endothelial activation and atherosclerosis [8, 12]. To determine if PPAR α can block PCB77-induced expression of the pro-inflammatory

proteins COX-2 and VCAM-1, porcine endothelial cells were pretreated with fenofibrate (FF) followed by exposure to PCB77, and total cellular proteins were extracted, followed by immunoblots. PCB77 significantly induced COX-2 protein expression, but pretreatment with FF significantly blocked this effect (Fig. 1). To determine if PPAR α activation can interfere with PCB77 induction of VCAM-1, endothelial cells were pretreated with FF or WY14643 followed by exposure to PCB77. PCB77 significantly induced VCAM-1 protein expression, which was partially blocked by pretreatment with FF or WY14643 (Fig. 2a and b). Basal COX-2 and VCAM-1 protein expression was not altered by treatment with either PPAR ligand.

PPAR α Agonists Block PCB77 Induction of CYP1A1 and Reduce Basal AHR Protein Expression

The majority of the pro-inflammatory effects of coplanar HAHs and PCBs are associated with activation of the AHR pathway [15, 29]. To determine if PPAR α can alter AHR pathway activation, protein expression and activity of the AHR responsive microsomal enzyme cytochrome P450 1A1 (CYP1A1) was measured in endothelial cells pretreated with PPAR α ligands followed by PCB77 exposure. CYP1A1 is a monooxygenase enzyme that is highly inducible by AHR ligands such as coplanar PCBs [27, 30]. Endothelial cell pretreatment with FF resulted in

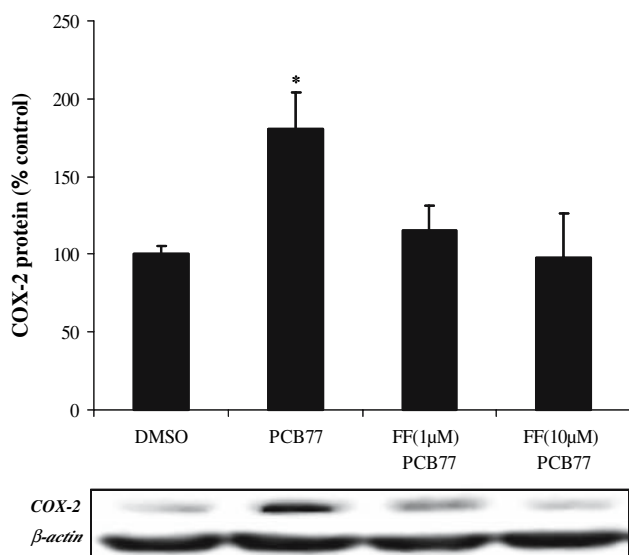


Fig. 1 Expression of COX-2 protein after pretreatment with the PPAR α ligand fenofibrate (FF; 1 or 10 μ M) followed by exposure to PCB77 (3.4 μ M). Cell cultures were pretreated with FF for 8 h followed by exposure to vehicle (DMSO), or PCB77 for 18 h followed by immunoblots. COX-2 densitometry values were normalized to β -actin signal. Blots represent one of the three replicates. Bars represent % of control values \pm SEM ($n = 3$); *Statistically significant difference ($P < 0.05$) relative to the control group

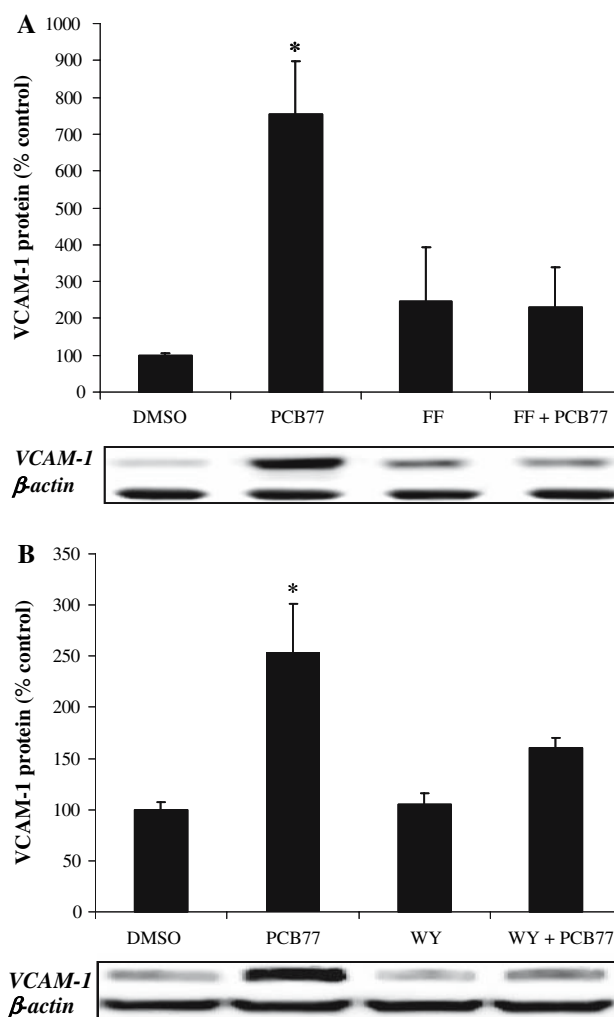


Fig. 2 (a and b): Expression of VCAM-1 protein after pretreatment with the PPAR α ligands fenofibrate (FF) or WY14643 (WY) at a final concentration of 10 and 20 μ M, respectively, followed by exposure to PCB77 (3.4 μ M). Cell cultures were pretreated with FF (a) or WY (b) for 8 h followed by exposure to vehicle (DMSO), or PCB77 for 18 h followed by immunoblots. VCAM-1 densitometry values were normalized to β -actin signal. Blots represent one of the three replicates. Bars represent % of control values \pm SEM ($n = 3$); *Statistically significant difference ($P < 0.05$) relative to the control group

a significant reduction in PCB77-induced CYP1A1 activity (Fig. 3a). Furthermore, treatment with WY14643 significantly reduced CYP1A1 protein induction by PCB77 (Fig. 3b). To determine if PPAR α activation affected AHR protein expression, endothelial cells were treated with FF prior to measurement of AHR expression by immunoblots. Treatment with FF resulted in a significant reduction of AHR protein expression (Fig. 4a), which was confirmed using the PPAR α ligand WY14643 (Fig. 4b). These results suggest that PPAR α activation disrupts the AHR pathway in endothelial cells by reducing AHR protein expression.

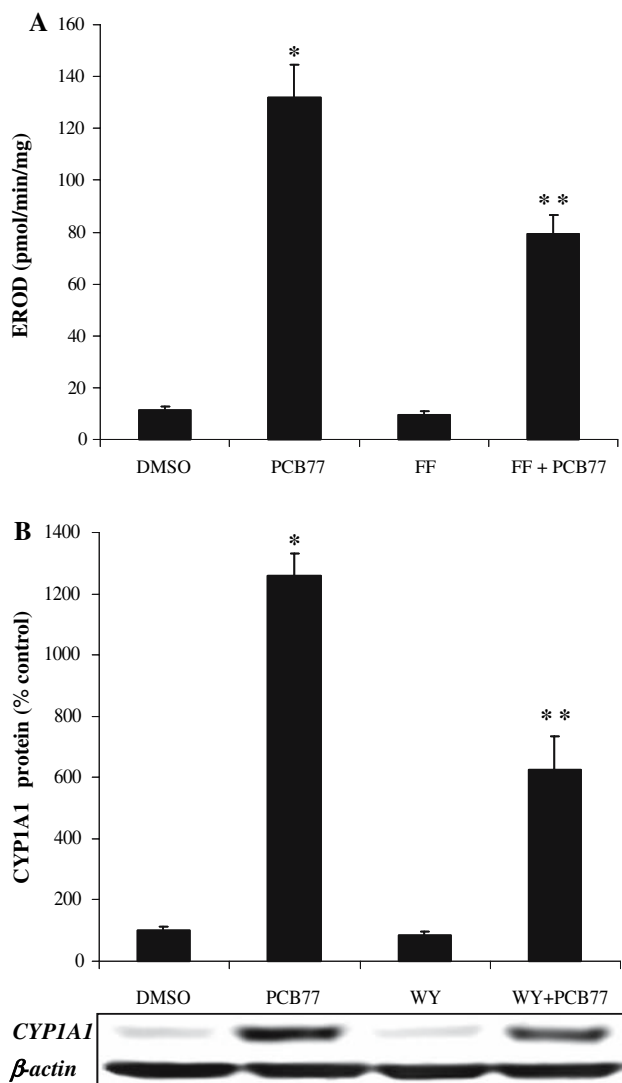


Fig. 3 (a and b): CYP1A1 activity (a) and protein expression (b) after pretreatment with the PPAR α ligands fenofibrate (FF) or WY 14643 (WY) at a final concentration of 10 and 20 μ M, respectively, followed by exposure to PCB77 (3.4 μ M). Cell cultures were pretreated with FF (a) or WY (b) for 8 h followed by exposure to vehicle (DMSO), or PCB77 for 18 h followed by EROD assay or immunoblots. CYP1A1 densitometry values were normalized to β -actin signal. Blots represent one of the three replicates. Bars represent CYP1A1 activity or protein expression (% of control values) \pm SEM ($n = 3$); * and ** represent statistically significant difference ($P < 0.05$) from vehicle and PCB treated cells, respectively

PCBs Decrease PPAR Transcriptional Function, Responsive Gene Expression, and Protein Expression

It has been previously shown that PPAR α expression and activity are attenuated during inflammation [31, 32]. To determine if PCB77 exposure can affect basal PPAR α function in endothelial cells, mRNA expression of PPAR α and the PPAR responsive gene CYP4A1 were measured after treating cells with increasing concentrations of

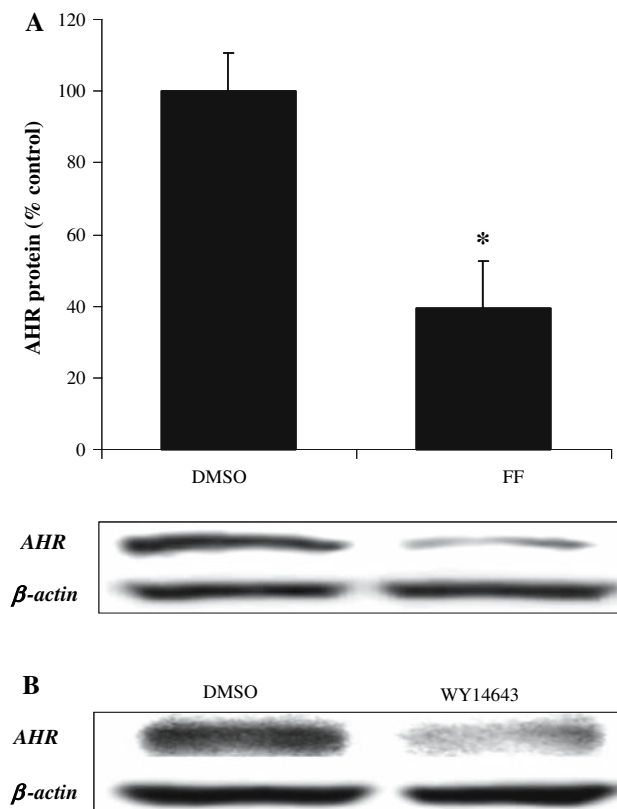


Fig. 4 Expression of AHR protein after treatment with the PPAR α ligands: fenofibrate (FF; 10 μ M) (a) or WY14643 (20 μ M) (b). Cell cultures were treated with FF or WY14643 for 18 h followed by immunoblots. AHR densitometry values were normalized to β -actin signal. Blots represent one of the three replicates. Bars represent % of control values \pm SEM ($n = 3$); *Statistically significant difference ($P < 0.05$) relative to the control group

PCB77. Treatment with PCB77 significantly reduced PPAR α and CYP4A1 mRNA expression at the lowest concentration used in these experiments (1 nM) (Fig. 5). We then measured the effects of PCB77 on PPAR α protein expression. Treatment with PCB77 led to a dose-dependent decrease in PPAR α protein expression (Fig. 6). To determine if PCB77 can affect PPAR transcriptional function, a PPAR responsive element driven reporter gene (luciferase) was used in MCF-7 cells. Treatment with PCB77 significantly reduced reporter gene expression (Fig. 7). These results suggest that PCB77 treatment reduced PPAR α expression and transcriptional activity.

Discussion

The results from this study demonstrate that PPAR α activation can protect endothelial cells from PCB-induced damage by decreasing expression of AHR protein and subsequent inhibition of the AHR pathway, and that

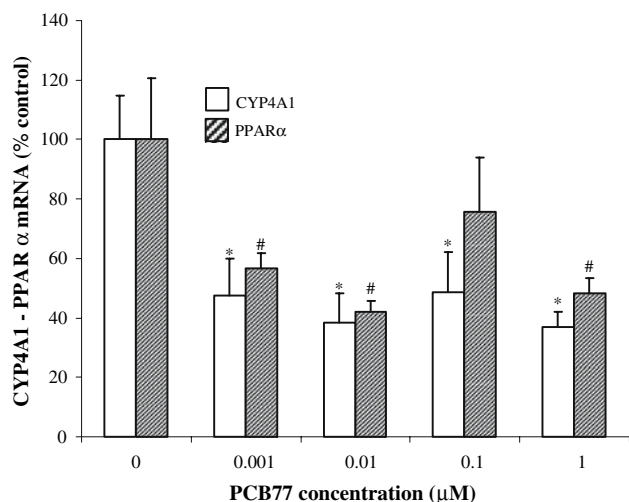


Fig. 5 Effects of PCB77 (0.001–1 μM) on mRNA expression of PPAR α and the PPAR α responsive gene CYP4A1. Cells were exposed to increasing concentrations of PCB77 for 6 h. mRNA expression was measured by real time PCR using SYBR Green for detection. PPAR α and CYP4A1 expression was normalized to β -actin signal. Bars represent % control values \pm SEM ($n = 3$); * and # represent statistically significant differences ($P < 0.05$) from the respective control groups

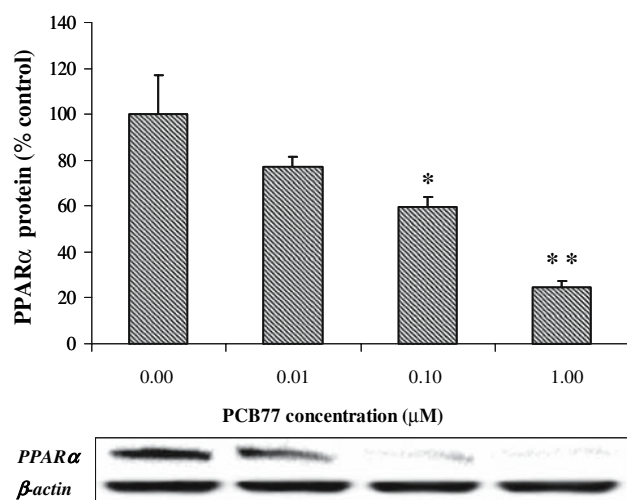


Fig. 6 Effects of PCB77 (0.01–1 μM) on PPAR α protein expression. Cells were exposed to increasing concentrations of PCB77 for 18 h followed by immunoblots. PPAR α densitometry values were normalized to β -actin signal. Blots represent one of the three replicates. Bars represent % of control values \pm SEM ($n = 3$ replicates per treatment); * and ** represent statistically significant differences ($P < 0.05$) relative to the control group, and different PCB77 concentrations

coplanar PCBs can negatively affect PPAR function, and decrease PPAR α mRNA and protein expression. It has been shown, in both cell culture and whole animal models that many of the pro-inflammatory and cardiotoxic effects of coplanar HAHs, including PCBs, are mediated by the AHR pathway [29], and the downstream activation of pro-inflammatory signaling cascades [15]. The results from the

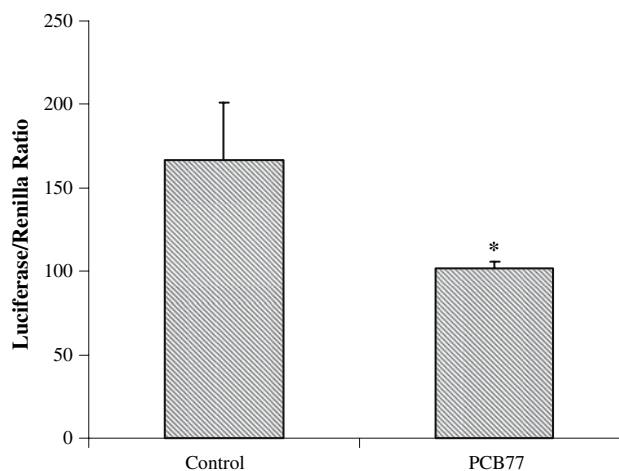


Fig. 7 Effects of PCB77 (3.4 μM) on PPAR transcriptional activity in MCF-7 cells stably transfected with a luciferase reporter gene driven by a triple repeat of PPRE. Cells were exposed to PCB77 for 6 h. Luciferase readings were normalized by readings obtained from measuring the CMV-driven renilla. Bars represent means \pm SEM ($n = 3$). *Statistically significant difference ($P < 0.05$) relative to the control group

experiments described above suggest that activation of PPAR α blocked PCB77 induction of the responsive pro-inflammatory proteins COX-2 and VCAM-1. Both of these proteins play an important role in endothelial cell activation and the pathogenesis of atherosclerosis. Induced COX-2 activity leads to increased production of prostaglandins, and reactive oxygen species [33, 34], while VCAM-1 is a cell surface receptor recognized by immune cells (monocytes and T lymphocytes) that promotes cell migration to the site of vascular injury [7]. These two proteins have been previously shown to be induced by HAHs [9, 12], and their upregulation is dependent on the presence of a functional AHR [9, 35].

The results from this study suggest that PPAR α activation can significantly reduce PCB77 effects by reducing AHR protein expression and the induction of AHR-regulated gene expression. It has been previously shown that PPAR α activation can block pro-inflammatory signaling pathways by various mechanisms, which include inhibition of transcription factors AP-1 and NF- κ B. This inhibition occurs through various mechanisms: direct interaction with these proteins [36], inducing expression of the NF- κ B inhibitory protein I κ B α [37], reducing basal expression of the pro-inflammatory transcription factors NF- κ B and CAAT enhancer binding protein β (C/EBP- β) [38], and the IL-6 receptor complex IL-6R/gp80 and gp130 [39]. Our results are supportive of previous studies showing that PPAR α activation can alter induction of AHR responsive genes, and reduce AHR expression in rat liver and HepG2 cells [40]. Specifically, we demonstrated in endothelial cells that PPAR α activation could significantly reduce

basal AHR expression, and CYP1A1 induction by PCB77. A recent study using rat liver epithelial cells showed that decreased AHR expression by siRNA targeted gene silencing can reduce CYP1A1 induction by the AHR ligands benzo-a-pyrene and benzo-b-fluoranthene [41]. However, studies using Caco-2 cells, a colorectal carcinoma cell line, suggest that PPAR α activation can potentiate AHR expression and CYP1A1 induction [42]. Overall, these data suggest that the outcome of PPAR α and AHR pathway interactions are tissue and cell type specific.

In the current study, we focused on PPAR α . Both PPARs α and γ are expressed in the vasculature, including the endothelium, and there is evidence that treatment with 2,3,7,8-tetrachlorodibenzo-*p*-dioxin (TCDD) can suppress PPAR γ expression [43, 44]. Our data suggest that PCBs can decrease both PPARs α and γ basal protein expression [45]. Various mechanisms triggered by coplanar PCB exposure could be associated with downregulating expression and function of PPARs α and γ . Previous studies have demonstrated that during adipogenesis, TCDD can cause suppression of PPAR γ expression through activation of MEK/ERK dependent mechanisms [43], and that these effects could be blocked by AHR inhibitors [46].

Another possible mechanism, by which PCBs decrease basal PPAR α expression and function, is by promoting oxidative stress and the activation of pro-inflammatory signaling cascades. PPAR α has been shown to be reduced by age-dependent and diet-induced oxidative stress, as well as exposure to pro-inflammatory cytokines [47–49]. Others and we have shown previously that exposure to coplanar PCBs such as PCB77 and PCB126 can activate NF- κ B, induce pro-inflammatory cytokine production and oxidative stress [8, 9, 15]. Finally, PPAR α downregulation by PCB77 could be due to AHR interference with transcription factors that regulate PPAR α expression. It has been shown that the PPAR α gene promoter contains multiple binding sites for Sp1 [50, 51], and AHR activation can affect Sp1 function to either promote or repress gene expression [52, 53].

The inhibitory effects of PCB exposure on the basal expression and function of the anti-inflammatory PPAR α could increase susceptibility to other types of pro-inflammatory stimuli. Various PPAR knockout models have shown that genetic elimination of PPAR α or γ expression is associated with increased susceptibility to pro-inflammatory agents [54, 55]. Dioxins and coplanar PCBs have been shown to increase expression of genes associated with systemic inflammation and atherosclerosis [9, 12, 15]. These effects, coupled with a decrease in expression of protective PPARs, may increase the potential for PCB toxicity in the vascular endothelium.

Both cell culture and whole animal studies have previously shown that treatment with PPAR α ligands can

significantly reduce inflammation and atherosclerosis [24, 56]. The data presented in the current study show that PCBs can decrease basal expression and function of PPAR α . Furthermore, our data demonstrate that PPAR α agonists can inhibit PCB-mediated endothelial cell activation by downregulating AHR expression and function. These observations present a novel mechanism, by which endothelial cell activation and inflammation induced by chlorinated AHR ligands can be attenuated by PPAR agonists.

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