Evaluation of Polycyclic Aromatic Hydrocarbons in the Activation of Early Growth Response-1 and Peroxisome Proliferator Activated Receptors

Jeong-Ho Kim,* Kiyoshi Yamaguchi,* Seong-Ho Lee,* Patricia K. Tithof,* Gary S. Sayler,† Joo-Heon Yoon,‡ and Seung Joon Baek*^{,1}

*Department of Pathobiology, and †Department of Microbiology, University of Tennessee, Knoxville, Tennessee 37996, and ‡Department of Otorhinolaryngology, Yonsei University College of Medicine, Seoul, South Korea

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Polycyclic aromatic hydrocarbons (PAHs) are ubiquitous environmental and food contaminants with known or suspected carcinogenic properties. In this study, we have evaluated whether PAHs activate the early growth response (EGR-1) gene and bind to peroxisome proliferator-activated receptor alpha (PPAR α) and delta (PPAR β/δ) in cell culture systems. Luciferase reporter systems were employed and several PAHs were evaluated for their ability to activate EGR-1 and PPARs. Some PAHs enhanced EGR-1 expression and activated PPAR α and PPAR β . Among them, benz(a)anthracene was found to act as a relatively potent activator of PPAR α and PPAR β/δ , and to significantly enhance EGR-1 transcription. These in vitro assays were confirmed by Western blot analysis, using cell lysates of tissue samples from mouse trapped at a highly contaminated Superfund site in the Chattanooga Creek floodplain in Chattanooga, Tennessee. We have found that a PPAR target gene, glycogen synthase kinase-3ß (GSK-3β), was down-regulated and EGR-1 was up-regulated in the mouse samples of Chattanooga Creek. In addition, select PAHs repressed GSK-3B and induced CYP4A in FaO rat hepatoma cells. In conclusion, PAHs activate PPARa and PPAR β/δ , and up-regulate EGR-1 expression in vitro as well as in vivo. These data may provide a diversity of PAH activity in several biological pathways.

Key Words: PAHs; PPARα; PPARδ; EGR-1; GSK-3β.

Polycyclic aromatic hydrocarbons (PAHs) are toxic and ubiquitous environmental pollutants. They are generally formed and emitted into the environment as a result of incomplete combustion of fossil fuels, wood, and other organic materials and from industrial processes. Humans and animals are exposed to PAHs from air, water, dietary, and occupational sources, and also from cigarette smoke. Their very long biological half-lives in combination with biological effects at very low concentrations have caused health concerns about carcinogenicity (Rubin, 2001), teratogenicity (Couture *et al.*, 1990; Incardona *et al.*, 2004), and cardiovascular disease (Gustavsson *et al.*, 1996, 2001; Thirman *et al.*, 1994). Although many reports focus on elucidating the molecular mechanism of PAHs in several diseases, pathways other than aryl hydrocarbon receptor (AhR) activation have received little attention.

Many chronic diseases, including cancer and cardiovascular diseases have been linked to heredity and/or the environment, which can either enhance or inhibit the disease process. One such molecular link between disease and the environment is early growth response-1 (EGR-1, also known as NGFI-A, Zif268, Krox24, and Tis8). EGR-1 is the prototypical member of a family of zinc finger transcription factors that includes at least three other members, EGR-2, -3, and -4. EGR-1 is especially induced by a range of physiological and environmental stimuli including growth factors, cytokines, ultraviolet light, ionizing radiation, and mechanical injury (Gashler and Sukhatme, 1995; Khachigian and Collins, 1998). EGR-1 appears to be critically involved in several diseases including angiogenesis and tumor formation. Alteration in expression could contribute to the deleterious effects of PAH exposure. Recently, Martinez et al. reported that halogenated aromatic hydrocarbon, 2.3.7.8-tetrachlorodibenzo-p-dioxin (TCDD), induces EGR-1 expression in human lung carcinoma cells (Martinez et al., 2004). However, expression of EGR-1 in nonneoplastic cells underlies diverse pathophysiological responses such as survival responses to damaging irradiation (Huang et al., 1999), development of vascular occlusion in arteriosclerosis (Silverman and Collins, 1999), and formation of severe pulmonary emphysema (Zhang et al., 2000). In addition to these diverse actions, expression of EGR-1 is commonly down-regulated in tumor cells in contrast with their normal tissue counterparts (Calogero et al., 2004; Hao et al., 2002; Huang et al., 1997; Levin et al., 1995; Shozu et al., 2004). However, EGR-1 is expressed at a higher level and promotes cell growth in prostate cancer when compared with normal tissues (Eid et al., 1998; Thigpen et al., 1996). Thus,

¹ To whom correspondence should be addressed at Department of Pathobiology, College of Veterinary Medicine, University of Tennessee, 2407 River Drive, Knoxville, TN 37996. Fax: (865) 974-5616. E-mail: sbaek2@utk.edu.

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EGR-1 has multiple functions in tumorigenesis, and the exact biological function of EGR-1 may be dependent on cell context as well as tissue types (Baek *et al.*, 2004). Nonetheless, the EGR-1's role in vascular disease has been firmly established by the fact that EGR-1 controls the expression of several genes implicated in the pathogenesis of atherosclerosis and restenosis (Breslow, 1996; Harja *et al.*, 2004; McCaffrey *et al.*, 2000; Silverman and Collins, 1999).

Peroxisome proliferator activated receptors (PPARs) are another molecular link between chronic disease and the environment. PPARs are members of the nuclear receptor superfamily and exist as three subtypes designated α , β (or δ), and γ . Among those, PPAR α activation is responsible for the pleiotropic effects of peroxisome proliferator such as enzyme induction, peroxisome proliferation, liver enlargement, and tumors (Klaunig et al., 2003). PPARa also plays a critical role in regulation of cellular uptake and β -oxidation of fatty acids (Berger and Moller, 2002; Marx et al., 2004). In contrast, PPAR δ (also known as PPAR β) is widely expressed with relatively higher levels in brain, colon, and skin. Although there have been extensive studies on PPAR α , much less is known about the function of PPARS. Nonetheless, recent studies suggest that PPARS plays a role in colon cancer (Gupta et al., 2004; He et al., 1999; Wang et al., 2004), and preadipocyte proliferation (Hansen et al., 2001).

We hypothesized that transcription factors, EGR-1 and PPARs may link environmental toxic compounds to human diseases. The aim of this study is to determine whether PAHs affect EGR-1 and PPARs activity and to identify the different activity of PAHs in reporter system. In this study, fifteen PAHs, which are commonly found in the environment, were examined as potential activators of PPAR α or PPAR β/δ , and inducer of EGR-1 gene expression in A549 human lung adenocarcinoma cells and HCT-116 human colorectal adeno carcinoma cells. The luciferase reporter genes were used to measure the activity of PPARs and transactivation of the EGR-1 promoter. We have demonstrated that some PAHs may activate PPAR α and PPAR β/δ , and transactivate EGR-1 promoter activity. Among those, benz(a)anthracene (BaA) induces EGR-1 and PPAR activation in culture systems. Furthermore feral mice (Peramyscus gossypinus) trapped along the floodplain of a Superfund site with high levels of PAH contamination demonstrated significant up-regulation of EGR-1 and down-regulation of GSK-3β, a PPAR target gene. These data provide evidence for diverse effects of PAHs that may be important in diseases linked to environmental pollution such as cardiovascular disease and carcinogenesis.

MATERIALS AND METHODS

Cell culture and chemicals. A549 human lung adenocarcinoma cell lines and HCT-116 human colorectal carcinoma cell lines were obtained from the American Type Culture Collection (ATCC) (Manassas, VA). A549 cells were cultured in RPMI-1640 medium supplemented with 10% FBS and 10 μ g/ml gentamycin. HCT-116 cells were cultured in Modified McCoy 5A medium supplemented with 10% FBS and 10 μ g/ml gentamycin. FaO rat hepatoma cell lines were generously provided by Dr. Seong-Jin Kim (National Cancer Institute, Bethesda, MD) and were cultured in Dulbecco's modified Eagle's medium supplemented with 10% FBS and 10 μ g/ml gentamycin. All the PAHs were purchased from Sigma-Aldrich (St. Louis, MO) and completely dissolved in DMSO.

Plasmids. EGR-1 promoter (-1260 to +35) linked to the luciferase gene (pEGR1260-Luc) was described previously (Baek et al., 2003). The plasmids used for studying PPAR α and PPAR β/δ activators were a reporter gene containing four copies of a Gal4 binding site (MH100×4-TK-Luc) and chimeric receptors (pCMX-Gal-mPPARa-LBD for PPARa and pCMX-GalmPPAR δ -LBD for PPAR δ). In this system, when a compound binds to the ligand binding domain (LBD) from PPARa or PPARS of the chimeric receptor (pCMX-Gal4-mPPARa-LBD or pCMX-Gal-mPPARô-LBD), then the DNA binding domain of the yeast Gal4 (denoted as Gal) binds to co-transfected Gal4 binding site and initiates transcription of the firefly luciferase (Luc). A reporter plasmid containing three copies of the PPAR response element (PPRE×3-TK-Luc) and a mouse PPARa cDNA (pCDNA3-mPPARa) were previously described (Nixon et al., 2003). This system directly measures activation of PPARa via transcriptional activation of the luciferase reporter gene as a result of PPARa binding to the PPAR response element (PPRE). All the PPAR and PPAR reporter plasmids were generously provided by Dr. Ronald M. Evans (Howard Hughes Medical Institute, CA).

Transient transfections and luciferase reporter assays. Cells $(1 \times 10^5$ cells/well) were cultured in twelve-well plates in culture medium containing 10% FBS. After growth for 16 h, the internal control, 0.05 µg pRL-null (Promega, WI) and 0.5 µg of the other plasmids were transfected using LipofectAMINE (Life Technologies, Inc.) according to the manufacturer's protocol. After 24 h, the media were changed and the cells were treated with the various PAHs dissolved in DMSO. The final concentration of DMSO did not exceed 0.1% (v/v) in any of the samples. Treatments with PAHs were performed under serum-free conditions. After 24 h treatment, the cells were washed with PBS and harvested in 1X luciferase lysis buffer. The luciferase activity was measured by a dual luciferase assay kit (Promega, WI), and normalized to the internal control, pRL-null (renilla luciferase) activity.

Animal studies. Feral mice (*Peramyscus gossypinus*) were trapped along the floodplain of the Chattanooga Superfund site and in a control site two miles upstream from the contaminated section of creek. All animal procedures were in compliance with the National Institute of Health guidelines on animal use and were approved by the University of Tennessee Institutional Animal Care and Use Committee. Traps were set each evening and checked each morning. Trapped animals were transported to the laboratory and anesthetized with CO₂. After the chest cavities were opened, the mice were exsanguinated by cardiac puncture and the distal aorta, heart and lungs perfused with phosphate buffered saline (PBS) to remove clotted blood. Sections of heart, lung, and colon were removed and snap-frozen in liquid nitrogen for Western blot analysis.

Western blot analysis. The level of protein expression was evaluated by Western blot analysis. FaO cells were grown to 60–80% confluency in 6 cm plates, followed by 24 h treatment of selected PAHs in the absence of serum. Total cell lysates were isolated using RIPA buffer (1× phosphate-buffered saline, 1% Nonidet P-40, 0.5% sodium deoxycholate, and 0.1% SDS) containing proteinase inhibitor, and the soluble protein concentrations were determined by BCA protein assay kit (Pierce, Rockford, IL). For feral mouse tissues, frozen samples were lysed in ice-cold RIPA buffer as described above. All lysate proteins were separated by SDS-PAGE and transferred for 1 h onto nitrocellulose membrane (Osmonics Inc., MN). The blots were blocked for 1 h with 5% skim milk in Tris-buffered saline and Tween 0.05% and probed with GSK-3 β (Cell Signaling, MA), EGR-1 (Santa Cruz Biotech., CA), CYP4A (Affinity Bioreagent, Golden, CO), or Actin (Santa Cruz Biotech., CA) antibody at 4°C overnight. After washing with Tris-buffered saline and Tween 0.05%, the blots were treated with horseradish peroxidase-conjugated secondary antibody for 1 h and washed several times. The signals were detected by the enhanced chemiluminescence system (Amersham Biosciences, Arlington Height, IL). The signal intensities were measured by NIH Image program (Scion Corp., MD).

Statistical analysis. For luciferase activities of transient transfection experiments, data were expressed as mean \pm SD for at least three independent repeats. For quantitative analyses, analysis of variance (ANOVA) with Tukey's multiple comparison test or *t*-test was used to compare mean values. SAS for Windows (9.1) (SAS Institute Inc., Cary, NC) statistical analysis software was used. A *p*-value of less than 0.05 was considered significant.

RESULTS

We have selected fifteen PAHs, containing two, three, four, or five aromatic rings. These PAHs were found to be in the highest concentrations in the Superfund site in Chattanooga, Tennessee (Elgayyar *et al.*, manuscript in preparation). Among these, benz(a)anthracene (BaA), benzo(a)pyrene (BaP), benzo(b)fluoranthene (BbF), benzo(k)fluoranthene (BkF), and chrysene (CHR) are considered to be carcinogenic, whereas acenaphthene (ANY), anthracene (ANT), naphthalene (NAP), pyrene (PYR), phenanthrene (PA), and triphenylene (TP) are not considered to be carcinogenic according to WHO (Table 1). A limited number of recent studies have shown that fluoranthene (FA) is an experimental carcinogen (Hecht *et al.*, 1995).

Effects of PAHs on EGR-1 Activation

Several PAHs were examined for the induction of *EGR-1* gene in A549 human lung carcinoma cell lines, which have been shown to induce EGR-1 expression (Martinez *et al.*, 2004). The pEGR1260-Luc construct was transiently transfected into A549 cells and the luciferase activity was measured in the presence of PAHs. The pEGR1260-Luc construct (Fig. 1A) contains 1.2 kb of the EGR-1 promoter (Baek *et al.*, 2005). All the PAHs tested here showed a significant increase in luciferase activity except BaP and TP (Fig. 1B). PA, BaA, and PA demonstrated higher luciferase activity, and PA also increased EGR-1 activity in 1 μ M concentration (Fig. 1C). As a positive control, sulindac sulfide (30 μ M) was used in the promoter assay (Baek *et al.*, 2005), showing a 4-fold induction of luciferase activity compared to vehicle treated sample.

Effects of PAHs on PPARa activation

In addition to cyctochrome p450 based biomarkers, peroxisome proliferators have been used for a pollution biomarker in aquatic organism (Cajaraville *et al.*, 2003). Among those, PPAR α has received much attention from many researchers since PPAR α expression is tightly regulated by various hormones. In addition, PPAR α plays a pivotal role in tumorigenesis and atherogenesis (Klaunig *et al.*, 2003; Marx *et al.*, 2004). Therefore, we have examined whether PAH compounds

TABLE 1 Characteristics of Polycyclic Aromatic Hydrocarbons Used in This Study

Compound	Structure	Molecular weight	Genotoxicity	Carcino- genicity	AhR activity
Acenaphthene		154	(?)	(?)	
Acenaphthylene	5	152	(?)		
Anthracene	ŬĈ	178	_	_	_
Benz(a)anthracene	œŶ	228	+	+	++
Benzo(a)pyrene		252	+	+	++
Benzo(b)fluoranthene		252	+	+	+++
Benzo(k)fluoranthene	\mathcal{D}	252	+	+	+++
Biphenylene		152			
Chrysene		228	+	+	+++
Fluoranthene	\otimes	202	+	(+)	+
1-Methylanthracene	CH3	192			
Naphthalene		128	_	(?)	
Phenanthrene	0J	178	(?)	(?)	
Pyrene	%	202	(?)	(?)	+
Triphenylene	Û	228	+	(-)	

Note. Genotoxicity and carcinigenecity according to WHO, 1998: +, positive; -, negative; ?, questionable; parentheses, result derived from small database (WHO, 1998). AhR-mediated activity (Machala *et al.*, 2001) value for benzo(*a*)pyrene (and PAHs with similar effects) was arbitrarily set as ++; values +++ and + indicate PAHs with more or less significant effects than benzo(*a*)pyrene; value of - indicates no effect.

activate PPAR α . The luciferase reporter construct and expression vector (Fig. 2A) were co-transfected into HCT-116 cells that have been shown a high transfection efficiency (Baek *et al.*, 2001). The cells were treated with 10 μ M of various PAHs for 24 h and luciferase activity was examined. The

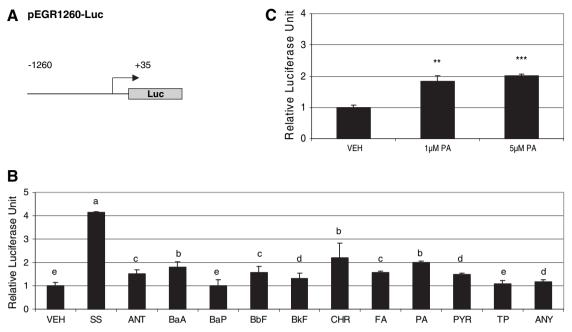


FIG. 1. PAH effects on EGR-1 expression. (A) Schematic diagram of pEGR1260-Luc construct. (B) A549 cells were exposed to PAHs after transfection with a pERG1260-Luc construct. Cells were treated with 5 μ M PA and 10 μ M of all other PAHs for 24 h. Sulindac sulfide was used as a positive control for EGR-1 activator (Baek *et al.*, 2004, 2005). Transfection efficiency for luciferase activity was normalized to the Renilla luciferase (pRL-null vector) activity. The *y*-axis shows relative luciferase unit (firefly luciferase unit/Renilla luciferase unit) by the fold increase in PAH treatment over vehicle treatment. The results are expressed as mean \pm SD of three independent transfections and different letters indicate significant difference (Tukey's multiple comparision test, *p* < 0.05). VEH, vehicle; SS, sulindac sulfide; ANT, anthracene; BaA, benz(a)anthracene; BaP, benzo(a)pyrene; BbF, benzo(b)fluoranthene; BkF, benzo(k)fluoranthene; CHR, chrysene; FA, fluoranthene; PA, phenanthrene; PYR, pyrene TP, triphenylene; ANY, acenaphthylene. (C) A549 cells were transfected and exposed to 1 μ M, 5 μ M of PA. The luciferase activity was measured, and relative luciferase unit was shown at *y*-axis. *t*-test, ***p* < 0.001, compared to vehicle.

luciferase activity to PAH treatment was compared to that of vehicle treatment. BaA was estimated as a strong activator among PAHs tested here. Other PAHs showed slight increases or little changes in luciferase activity (Fig. 2B). As a positive control, Wy14643 was used in this system and we found around 55-fold induction of luciferase activity, compared to vehicle treated samples (data not shown). Transfected cells were also exposed to BaA and luciferase activity was significantly increased in a dose dependent manner (Fig. 2C). It is important to note that the results shown in Figure 2 reflect solely activator activity with exogenous transfectants. Therefore, to examine the transactivation of PPARa, A549 cells were transiently cotransfected with a pPPRE×3-TK-Luc and pCDNA3-mPPARa expression vector (Fig. 3A), and then treated with vehicle, and 1 or 10 µM of BaA. As shown in Figure 3B, we observed a significant induction of luciferase activity in a dose dependent manner, indicating that BaA binds to PPARa and transactivates the gene containing PPRE sites in the promoter.

Effects of PAHs on PPAR β/δ activation

It has been suggested that PPAR β/δ may play an important role in tumorigenesis and atherogenesis (Gupta *et al.*, 2004; He *et al.*, 1999; Marx *et al.*, 2004; Tong *et al.*, 2000). PAH compounds were tested for PPAR β/δ activity in HCT-116 cells transiently transfected with pMH100×4-TK-Luc and pCMX- Gal4-mPPAR δ -LBD constructs (Fig. 4A). The PPAR β/δ activity of each PAH was compared with vehicle. Interestingly, treatment with BaA also induced strong luciferase activity among PAHs tested here. However, other PAHs also slightly increased luciferase activity, which is significant (Fig. 4B). BaA also increased luciferase activity in a dose dependent manner (Fig. 4C).

GSK-3β and EGR-1 Expression in the Presence of PAHs

CYP4A is induced by PPARa activation in liver tissue (Johnson *et al.*, 1996). GSK-3 β is repressed when FaO rat hepatoma cells are exposed to PPAR α and PPAR β/δ ligands (Vanden Heuvel et al., 2003). We have shown that some PAHs are able to bind the PPAR α and PPAR β/δ as an activator. To determine whether PAHs affects PPARa downstream in FaO cells, we performed Western blot analysis using cell lysates prepared from FaO cells treated with BaA, BaP, or PA. As shown in Figure 5A, all the PAHs tested increase CYP4A expression and suppressed GSK-3ß expression relative to vehicle-treated cells. BaA was the relatively strong suppressor of GSK-3ß expression, whereas BaP or PA was the weak suppressor. These data are consistent with previous data showing that BaA is the relatively strongest PPAR α and PPAR β/δ activator, whereas BaP and PA are relatively weak (Figs. 2 and 4). Next, we examined GSK-3β expression in vivo

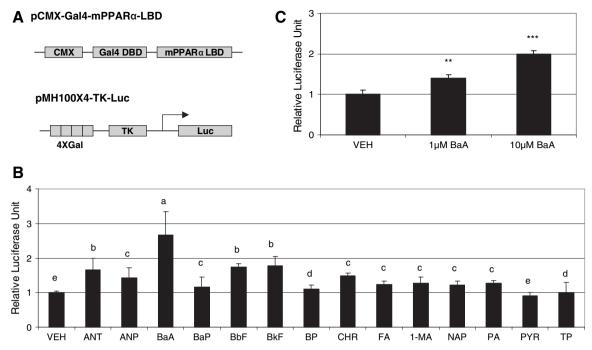


FIG. 2. PAH effects ozn PPAR α binding affinity. (A) Schematic diagram of pCMX-Gal4-mPPAR α -LBD and pMH100X4-TK-Luc vector. (B) HCT-116 cells were transfected with the vectors shown in Figure 2A, and the cells were treated with 10 μ M of various PAHs indicated for 24 h. Transfection efficiency for luciferase activity was normalized to Renilla luciferase (pRL-null vector) activity. The *y*-axis shows relative luciferase unit (firefly luciferase unit/Renilla luciferase unit) by the fold increase in PAH treatment over vehicle treatment. The results are expressed as mean ± SD of four independent transfections and different letters indicate significant difference (Tukey's multiple comparison test, *p* < 0.05). ANP, acenaphthene; BP, biphenylene; 1-MA, 1-methylanthracene; NAP, naphthalene. Other abbreviations are listed in Figure 1B. (C) The transfected cells were treated with the indicated concentration of BaA for 24 h and luciferase activity was measured. *t*-test, ***p* < 0.001 compared to vehicle.

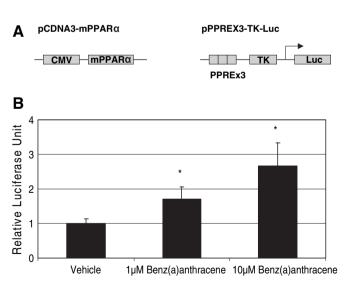


FIG. 3. PAH effects on PPAR α transactivation in the promoter containing PPRE sites in A549 cells. (A) Schematic diagram of pCDNA3-mPPAR α expression vector and pPPREX3-TK-Luc reporter vector. (B) A549 cells were transfected and the cells were treated with either vehicle or indicated BaA for 24 h. *p < 0.05 compared to vehicle. Luciferase activity was measured as described above.

using feral mice (Peramyscuc gossypinus) trapped along the floodplain of a highly contaminated area of the Chattanooga Creek and also along a noncontaminated area. We evaluated GSK-3β expression in colon, heart, and lung tissues of mice from both areas. As shown in Figure 5B, GSK-3B was downregulated in heart but not lung from mice trapped in the contaminated area of the Chattanooga Creek when compared to mice trapped in the control area. However, GSK-3β was not detected in colon tissue from both mice trapped from either site. We also measured EGR-1 expression and found that EGR-1 was induced in colon samples from mice trapped in the Superfund site when compared to control. However, no EGR-1 was detected in heart or lung samples from either site. EGR-1 induction in the in vivo study is consistent with in vitro data, showing that PAHs increase EGR-1 expression as assessed by a reporter system (Fig. 1). Taken together, these results suggest that some PAHs, particularly BaA activates PPARa and PPAR β/δ , and alters the PPAR target gene expression such as CYP4A and GSK-3β.

DISCUSSION

Humans and other living organisms are constantly exposed to a large number of potentially toxic environmental

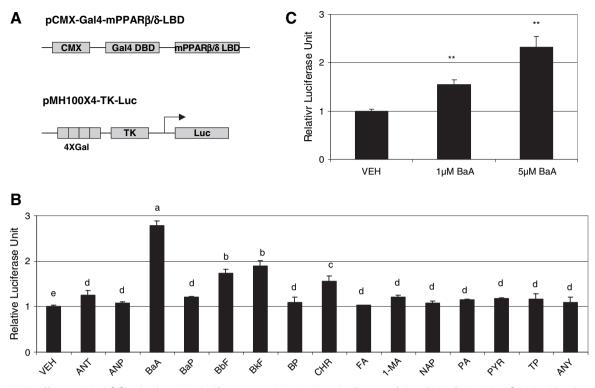


FIG. 4. PAHs effects on PPARβ/δ activation using luciferase assay. (A) A schematic diagram of the pCMX-Gal4-mPPARδ-LBD chimeric gene and the pMH100X4-TK-Luc reporter vector used in this study. (B) HCT-116 cells were transfected with vectors shown in A, and the cells were treated with 10 μ M of various PAHs for 24 h. Transfection efficiency for luciferase activity was normalized to the Renilla luciferase (pRL-null vector) activity. The *y*-axis shows relative luciferase unit (firefly luciferase unit/Renilla luciferase unit). The results are expressed as mean ± SD of three independent transfections and different letters indicate significant difference (Tukey's multiple comparision test, *p* < 0.05). The abbreviations are described in Figure 2B. (C) Transfected cells were treated with indicated concentrations of BaA for 24 h and luciferase activity was measured. *t*-test,. ***p* < 0.001, compared to vehicle treatment.

chemicals, including the ubiquitous polycyclic aromatic hydrocarbons (PAHs). A number of PAHs have been known to affect several chronic diseases, including cancer and cardiovascular disease, in experimental animals following oral, pulmonary, dermal, or subcutaneous administration. Most of the biological effects of PAHs are considered to be mediated via aryl hydrocarbon receptor (AhR)-dependent gene expression (Nebert et al., 2000). PAHs also cause oxidative DNA damage and DNA adduct formation, which is the most widely accepted mechanisms of PAH-induced tumorigenesis and cardiovascular disease (Lee and Blair, 2001; Toraason et al., 2001; Wogan et al., 2004). Compared to genotoxicity or AhR-mediated toxicities of PAHs, knowledge about the nongenotoxic effects of PAHs remains to be elucidated. The major goal of our investigation was to determine the cellular mechanisms that are induced by PAHs in AhR independent manner. EGR-1 and PPARs are the transcription factors that control many genes in the presence of stimuli, including chemical compounds. The ability of select PAHs to activate the EGR-1 gene and to bind to PPAR α and PPAR β/δ was examined in this study using several PAHs commonly found in a Superfund site in Chattanooga, Tennessee. We found that some PAHs activated EGR-1 gene and enhanced PPAR α and PPAR β/δ activity in vitro. The latter

resulted in the suppression of GSK-3 β expression, which may play an important role in tumorigenesis.

EGR-1 can induce expression of a set of vasculature genes, such as PDGF-A and B chain, bFGF, TGF- β , TNF- α , and intracellular adhesion molecule-1. Expression of the EGR-1 is elevated in prostate cancer and correlates with tumor progression. Thus, EGR-1 is the key mediator in orchestrating the functional characteristics of the vessel wall and tumorigenesis. However, EGR-1 can be related to anti-tumorigenesis and protumorigenesis, depending on cell and tissue types. While EGR-1 induces anti-tumorigenic proteins including p53, PTEN, and NAG-1 (Baek et al., 2005), EGR-1 is expressed at a higher level and promotes cell growth in prostate cancer (Eid et al., 1998; Thigpen et al., 1996). Thus, EGR-1 could play a role in both cell proliferation and growth arrest. In contrast to EGR-1 functions in tumorigenesis, experimental evidence is emerging to link EGR-1 to chronic vascular and inflammatory stress in vivo. The role of EGR-1 in atherosclerosis related to PAH exposure has not been examined. In this study, EGR-1 was not up-regulated in heart tissue from mice exposed naturally to environmental contaminants; however, vascular tissue was not examined. Further studies are warranted to evaluate the potential role of EGR-1 in atherosclerosis progression related to atherosclerosis and heart disease.

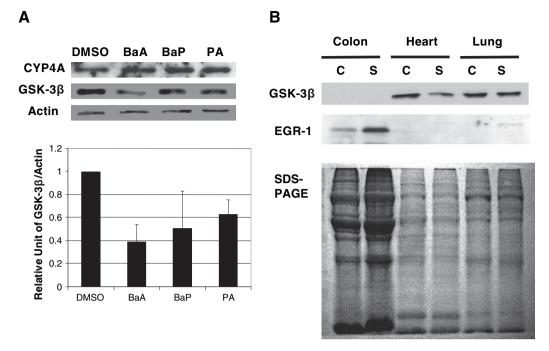


FIG. 5. Regulation of GSK-3 β , CYP4A, and EGR-1 by PAHs. (A) FaO rat hepatoma cells were treated with 10 μ M of various PAHs for 24 h (DMSO, vehicle control; BaA, benz(a)anthracene; BaP, benzo(a)pyrene; PA, phenanthrene). Western blotting was performed using a GSK-3 β antibody (Cell Signaling, Beverly, MA) and CYP4A antibody (Affinity Bioreagent, Golden, CO). Equal loading (30 μ g) was confirmed by determining actin immunoreactivity. The relative GSK-3 β expression levels normalized by actin from three independent experiments are shown on the bottom. (B) Mouse tissue samples from colon, heart, and lung were isolated from mouse tissue collected from the control site (C) or from the Superfund site (S) in Chattanooga, Tennessee. Cell lysates (100 μ g) were subjected to Western blot analysis for GSK-3 β and EGR-1 protein expression. Equal loading was confirmed by Bio-Safe Coomassie staining (Bio-Rad, CA) of total proteins shown in bottom. The blot shown is a representative of two independent experiments.

PPAR α is a key contributor in the processes of peroxisome proliferation, hypertrophy, cell proliferation, and hepatocarcinogenesis in vivo. Its over-expression is observed in advanced prostate cancer (Collett et al., 2000), and activation of PPARa promotes cell proliferation in breast cancer cells (Klaunig et al., 2003; Suchanek et al., 2002). In addition, activation of PPAR α has been demonstrated to modulate many aspects of lipoprotein metabolism and inflammation in vitro, as well as in animal and human studies (Israelian-Konaraki and Reaven, 2004). Thus, activation of PPAR α may play a role in disease such as tumorigenesis and atherogenesis. On the other hand, the activation of PPAR β/δ plays an anti-apoptotic role in keratinocytes via transcriptional control of the AKT signaling pathway (Di-Poi et al., 2002). Genetic disruption of PPARβ/δ also decreases the tumorigenicity of human colon cancer cells transplanted into mice (Park et al., 2001). Our results support the contention that some PAHs with known carcinogenic activity are relatively strong PPAR α and PPAR β/δ activators as assessed by reporter system. These results support that some PAHs may induce chronic disease through PPAR activation mechanism other than AhR activation. These results also suggest that minute differences in PAH structure result in the activation of two different PPARs.

In this report, we have shown that GSK-3 β is suppressed in heart tissue from mice trapped in a highly contaminated

Superfund site as well as in the FaO cells. GSK-3 β is known to be a negative regulator of cardiac hypertrophy (Hardt and Sadoshima, 2002), and we have recently reported that AKT/GSK-3 β plays an important role in apoptosis (Yamaguchi *et al.*, 2004). Therefore, the suppression of GSK-3 β by environmental contaminants such as PAHs may be important in the processes by which environmental pollution accelerates cardiac disorders or tumorigenesis such as cardiomyopathy or cancer; however, further studies may be required to elucidate the exact molecular mechanism.

In conclusion, our data suggest that some PAHs, particulary BaA, are able to activate EGR-1 promoter and act as an activator of PPAR α and PPAR β/δ *in vitro*. BaA can activate target genes of PPAR α and PPAR β/δ , thereby repressing the GSK-3 β expression *in vitro* and *in vivo* and inducing the CYP4A expression *in vitro*. The repression of GSK-3 β and activation of EGR-1 by some PAHs may provide a novel approach to elucidating the various effects of PAHs on human chronic disease.

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