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ABSTRACT

Previously, we demonstrated the sensitivity of RPTEC/TERT1 cells, an immortalized human renal proximal tubule epithelial cell line, to two common environmental carcinogens, cadmium (Cd) and benzo[a]pyrene (B[a]P). Here, we measured BPDE-DNA adducts using a competitive ELISA method after cells were exposed to 0.01, 0.1, and 1 µM B[a]P to determine if these cells, which appear metabolically competent, produce BPDE metabolites that react with DNA. BPDE-DNA adducts were most significantly elevated at 1 μ M B[a]P after 18 and 24 h with 36.34 ± 9.14 (*n* = 3) and 59.75 ± 17.03 (*n* = 3) adducts/10⁸ nucleotides respectively. For mixture studies, cells were exposed to a non-cytotoxic concentration of Cd, 1 μ M, for 24 h and subsequently exposed to concentrations of B[a]P for 24 h. Under these conditions, adducts detected at $1 \mu M B[a]P$ after 24 h were significantly reduced, 17.28 ± 1.30 (n=3) adducts/10⁸ nucleotides, in comparison to the same concentration at previous time points without Cd pre-treatment. We explored the NRF2 antioxidant pathway and total glutathione levels in cells as possible mechanisms reducing adduct formation under coexposure. Results showed a significant increase in the expression of NRF2-responsive genes, GCLC, HMOX1, NQO1, after 1 μ M Cd \times 1 μ M B[a]P co-exposure. Additionally, total glutathione levels were significantly increased in cells exposed to 1 μ M Cd alone and 1 μ M $Cd \times 1 \mu M B[a]P$. Together, these results suggest that Cd may antagonize the formation of BPDE-DNA adducts in the RPTEC/TERT1 cell line under these conditions. We hypothesize that this occurs through priming of the antioxidant response pathway resulting in an increased capacity to detoxify BPDE prior to BPDE-DNA adduct formation.

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1. Introduction

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Over 90% of kidney cancers originate in the renal proximal tubule epithelial cells and are classified as renal cell carcinoma (RCC). However, only about 2% of kidney cancer cases can be attributed to a genetic predisposition [1,2]. The remaining cases occur in otherwise healthy

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individuals with no prior familial history [3]. Substantial evidence of environmental risk factors contributing to the development of RCC suggests that further scrutiny of human mutagens and carcinogens on the cellular and molecular level is warranted [4,5].

Exposure to polycyclic aromatic hydrocarbons (PAHs) has been associated with an elevated risk of many cancers including skin, lung, bladder, liver, and stomach [6]. PAHs are formed as byproducts of incomplete combustion and are ubiquitous in the environment. Major routes of exposure include inhalation and ingestion, which can result from cigarette smoking, consumption of grilled or contaminated foods, and atmospheric pollution associated with the burning of fossil fuels [7]. Increased consumption of chargrilled meats has been shown to directly correlate with elevated PAH exposure and risk of RCC [8,9]. Additional factors associated with the development of RCC include obesity, smoking, and hypertension [5,10,11].

The ability of cells to repair bulky PAH-DNA adducts may be altered by the presence of environmental contaminants such as cadmium (Cd). Cd has been shown to substitute for zinc ion co-factors in many DNA repair proteins and enzymes specifically responsible for recognizing and repairing DNA adducts [12]. Cd, a heavy metal and known nephrotoxicant, is present in the environment in food, cigarettes, and contaminated water runoff. Human exposure to Cd occurs primarily through inhalation of fine particulates (i.e. tobacco smoke) and consumption of foods such as rice, cereal, and mollusks [13]. Cd accumulates in the liver, kidneys, and bone and is suspected to promote cancers in these organs as well as in the lungs. Cd lacks strong mutagenic properties but may act as a co-carcinogen in the body by inhibiting DNA damage repair processes and increasing oxidative stress in cells [14,15].

Because individuals are rarely exposed to singlechemical agents or carcinogens in the environment, it is important to study these compounds as humans might encounter them on a daily basis. Exposure to chemical mixtures can result in toxicological outcomes that substantially differ from the expected effects of each compound alone. Toxicants in mixtures may act through similar or distinctly different mechanisms of action. Chemicals and compounds can act antagonistically, additively, synergistically, or one chemical may potentiate the effects of another [16]. Ultimately, these interactions may substantially alter toxicity to different and possibly unexpected degrees. For example, in vitro studies have shown that exposure to binary combinations of PAHs including benzo[a]pyrene (B[a]P) and benzo[b]fluoranthene (B[b]F) results in a significant increase in the formation of DNA adducts than exposure to B[a]P alone. However, exposure to both B[a]P and benzo[k]fluoranthene (B[k]F), a similarly structured PAH, results in a significant reduction in the formation of DNA adducts than exposure to B[a]P alone [17–19]. The opposing results that occur even after exposure to compounds of the same toxicant class emphasize the need for studies investigating effects elicited after exposure to mixtures of toxicants from similar and different classes.

In order to study the mechanisms of mixture exposure, which may promote RCC, we utilized an immortalized human renal cell line, RPTEC/TERT1. The RPTEC/TERT1 cell

line was derived from the renal proximal tubule epithelial cells (RPTEC) of a normal, healthy male donor. These cells were immortalized with the catalytic subunit of the human telomerase reverse transcriptase enzyme (TERT1) [20]. Previously, we determined that RPTEC/TERT1 cells exhibit sensitivity and compound-specific responses to B[a]P and Cd treatment [21]. Our results were consistent with canonical biological responses to both environmental toxicants and demonstrate metabolic competency of the RPTEC/TERT1 cell line. To test our hypothesis that Cd may alter formation of adducts after B[a]P exposure, we have explored concentration-dependent formation of BPDE-DNA adducts through cellular bioactivation of B[a]P. We examined the persistence of those adducts under conditions of pre-treatment with Cd. We intended to determine the effects of Cd on the persistence of BPDE-DNA adducts as a function of time, co-exposure, and oxidative stress.

We hypothesize that exposure to a binary combination of the environmental carcinogens, Cd, a heavy metal, and B[a]P, a representative PAH, acts to alter DNA adduct formation in comparison to levels found after B[a]P exposure alone. As Cd is known to inhibit the recognition and/or repair of PAH-DNA adducts, it is plausible to find persistence of adducts under conditions of co-exposure [22]. Alternatively, co-exposure may result in an antagonistic response leading to the formation of fewer DNA adducts through increased detoxification or inhibition of bioactivation. However, our previous work in the RPTEC/TERT1 cell line suggests that the inhibition of bioactivation is unlikely [21]. The interaction of chronic, low level exposure to both Cd and PAHs over a lifetime may provide support for environmental contributions to the development of RCC in healthy individuals.

2. Materials and methods

2.1. Reagents

2.1.1. Chemicals

All chemicals were purchased from Sigma–Aldrich (St. Louis, MO) unless noted otherwise. Cadmium chloride (CdCl₂, 202908) was dissolved in fresh complete medium and delivered at 0.1% of the final culture volume to yield the appropriate target concentrations. Benzo[a]pyrene (B[a]P, B1760) was dissolved in dimethyl sulfoxide (DMSO, D8418) and delivered at 0.05% of the final culture volume to yield the appropriate target concentrations. B[a]P preparations and exposures were carried out under low light conditions.

2.1.2. DNA isolation reagents

Enzymes used for DNA isolation including RNaseT1, mRNAse A, and proteinase K were purchased from Sigma–Aldrich. Tris-buffered saturated phenol, phe-nol:chloroform:isoamyl (25:24:1), and 5 PRIME Phase Lock Gel, light, 15 mL tubes for DNA isolation were purchased from Fisher Scientific (Pittsburg, PA).

2.1.3. BPDE-DNA adduct ELISA reagents

Greiner Bio-One microplates (high-binding, white) were purchased from Fisher Scientific. I-Block casein-based blocking solution and CPD-Star Substrate with Emerald-II Enhancer were purchased from Life TechnologiesTM (Grand Island, NY). Polyclonal BPDE-DNA antiserum was kindly provided by Dr. Regina Santella. Biotin-labeled goat anti-rabbit secondary antibody (Cat #111-065-045) was purchased from Jackson ImmunoResearch (West Grove, PA). Streptavidin-alkaline phosphatase conjugate (Cat #21324) was a product of Pierce and purchased from Fisher Scientific. Standard BPDE-DNA adducts were prepared from highly purified calf thymus DNA (Sigma, St. Louis, MO) and benzo[*a*]pyrene-*r*-7,*t*-8-dihydrodiol*t*-9,10-epoxide(),(anti) from MRIGLOBAL Chemical Carcinogen Repository (Kansas City, MO) according to the procedures described by Jennette et al. [23].

2.2. Cell culture

RPTEC/TERT1 cells and culture medium were purchased from Evercyte Laboratories (Vienna, Austria), and grown according to Evercyte's instructions. Cells were cultured at 37 °C in a humidified atmosphere containing 5% CO₂. RPTEC/TERT1 cells were passaged approximately once or twice per week and subcultured at a 1:2 or 1:3 ratio. Cell culture vessels were purchased from Fisher Scientific and CellTreat[®] Scientific Products (Shirley, MA) and were tissue culture treated to promote adherent cell growth.

2.3. Cell exposure

Cd was dissolved in fresh complete medium and delivered at 0.1% of final volume to give appropriate dose ranges. B[a]P was dissolved in DMSO and delivered at 0.05% final volume to give appropriate concentration ranges. B[a]P exposures were conducted under low light conditions. Regardless of exposure format, final volume percentage of each chemical was maintained. For co-exposure experiments, 1 μ M Cd was used to pre-treat cells for 24 h before B[a]P exposure. The Cd pre-treatment concentration was determined based on previous characterization of the cell line's responses to various Cd concentrations. One micromolar Cd was the highest concentration tested that showed no significant cytotoxicity at 24-h, 48-h, or 1-week post-exposure while demonstrating significantly increased cellular responses at the level of the gene and protein [21].

For DNA isolation, RPTEC/TERT1 cells were treated at confluence in T75 cm² tissue culture treated flasks. After exposure time points, cells were washed twice with cold $1 \times$ PBS, collected by centrifugation at 4°C, and stored at -80 °C until DNA was isolated.

2.4. Gene expression

RPTEC/TERT1 cells were grown to confluence in 60 mm dishes and exposed to Cd or B[a]P as described above. Cells were exposed in triplicate for each concentration and time point examined. Total RNA was isolated from cells after appropriate time points using the QIAshredder (QIAGEN, 79656, Valencia, CA) and RNeasy extraction kit (QIAGEN, 74136) following the manufacturer's instruction. RNA concentration and purity were assessed using a Thermo Scientific Nanodrop 2000c spectrophotometer.

Table 1

Primer-probe sets used for RPTEC/TERT1 gene expression, Applied Biosystems[®] TaqMan[®] gene expression assays.

Gene ID	Gene function	Gene location	Assay ID
GCLC	Antioxidant	6p12	Hs00155249_m1
HMOX1	Antioxidant	22q13.1	Hs01110250_m1
NQO1	Quinone reduction antioxidant	16q22.1	Hs00168547_m1
ACTB	Reference	7p22.1	Hs99999903_m1

RNA samples were diluted to $0.5 \,\mu g/\mu L$ in nuclease-free water.

Two microliters of each RNA sample were used for cDNA synthesis reactions to deliver 1 µg template in a 20 µL total reaction volume. cDNA was synthesized using iScript cDNA synthesis (BioRad, 170-8891, Hercules, CA) protocol as follows: 5 min at 25 °C, 30 min at 42 °C, and 5 min at 85 °C. RNA templates and cDNA were stored at -20 °C until use. Gene expression was determined using primer-probe sets from Applied Biosystems[®] TagMan[®] Gene Expression Assays. Actin, beta (ACTB) was used as a reference gene. Primers used are listed in Table 1. The thermal cycling protocol followed the manufacturer's instructions: 50 °C for 2 min and 95 °C for 10 min followed by 40 cycles of 95 °C for 15 s and 60 °C for 1 min. Reactions were conducted in 20 µL volumes with each sample being run in duplicate. All reactions were carried out using a BioRad C1000TM thermal cycler equipped with a CFX96TM Real-Time PCR Detection System.

2.5. DNA isolation

Genomic DNA was isolated with a standard phenol chloroform extraction. Briefly, cell pellets were thawed and incubated with $1 \times$ TE buffer, RNaseT1, mRNAse A, and SDS for 45 min at 37 °C. Pellets were incubated with proteinase K for 60 min at 60 °C and then overnight at 37 °C. Deproteinized DNA was extracted using 5 PRIME Phase Lock Gel light, 15 mL, tubes to increase yield from the aqueous phase. Precipitated DNA was spooled onto a glass pipette, transferred to 70% ethanol, and collected by centrifugation (18,000 rcf for 10 min). Ethanol was decanted and DNA was allowed to dry completely before reconstituting in sterile, DNA grade water. DNA concentration and purity were assessed using a Thermo Scientific Nanodrop 2000c spectrophotometer.

2.6. BPDE-DNA adduct ELISA

BPDE-DNA adducts were measured by a competitive ELISA method [24–26]. Briefly, 96-well white microplates were coated by adding 50 pg BPDE-substituted DNA in PBS to each microwell. The DNA was sonicated and denatured in a boiling water bath for 5 min before coating. Plates were allowed to dry overnight and washed twelve times the next day with washing buffer ($1 \times PBS/0.05\%$ Tween 20). All subsequent wash steps were also performed twelve times. Plates were treated with I-Block ($200 \mu L/well$) for 90 min at $37 \,^{\circ}$ C to prevent non-specific binding. Standard curves and samples were prepared by mixing and incubating with the previously characterized polyclonal BPDE-DNA antisera at 1:3,000,000 in I-Block buffer [25]. A 5-point standard

curve was used, in triplicate, to give a range of 0.312-10 fM adducts/well. Unknown samples were assessed at $10 \mu g$ DNA per well in triplicate after sonication and denaturation. The plate was washed after incubation with primary antibody, and a biotin-labeled goat anti-rabbit secondary antibody (1:2500 in I-Block) was incubated with in each well for 1 h. After an additional wash, the plate was incubated for 1 h with streptavidin-alkaline phosphatase conjugate (1:40,000 in I-Block). After one more wash step, the CPD-Star Substrate with Emerald-II Enhancer was used to produce and amplify signal. Luminescence was read with a Tecan Infinite[®] 200 PRO multimode reader (Tecan, San Jose, CA).

Adducts were calculated for unknown samples based on percent inhibition of the standard curve and expressed as average number of adducts per 10⁸ nucleotides. Nonspecific background signal detected in vehicle control groups was subtracted.

2.7. Total glutathione assay

After determined exposure time points, cells were trypsinized, collected, and washed twice in $1 \times$ cold PBS. Total glutathione levels in cells were determined using OxiSelectTM Total Glutathione (GSSG/GSH) Assay Kit (Cell Biolabs, Inc., San Diego, CA) according to the manufacturer's instructions. Cell isolates were diluted at 1:100 for use within the linear range of the assay.

2.8. Statistical analysis

One- and two-way ANOVAs were performed using the GraphPad Prism analytical software, version 6.0 (San Diego, CA). Data total glutathione assays were analyzed using a one-way ANOVA and Dunnett's multiple comparison tests. Data for gene expression were analyzed using a two-way ANOVA and Tukey's *post hoc* test. An α of 0.05 was used as the criteria for determining significance.

General linear models were used to test for differences among treatments, treatment groups, and time points for the BPDE-DNA adduct ELISA. Where the initial GLM analysis of variance (GLM-ANOVA) indicated a significant difference, *post hoc* mean comparisons were conducted using a Tukey correction. Statistical testing was conducted using IBM SPSS Statistics version 19 software (Armonk, NY). An α of 0.05 was used as the criteria for determining significance.

3. Results

3.1. BPDE-DNA adducts are formed and detected after exposure to B[a]P but altered after co-exposure to B[a]P and Cd

After 18 h of exposure to B[a]P alone, BPDE-DNA adducts were detected in RPTEC/TERT1 DNA samples. Although there appeared to be a dose-dependent increase in adduct formation after 18 h, exposure to 1 μ M B[a]P was significantly increased over DMSO vehicle control or lower concentrations, 0.01 and 0.1 μ M B[a]P. After 24 h of exposure to B[a]P alone, adduct formation was most

Table 2

BPDE-DNA adducts formed after B[a]P and Cd exposure in RPTEC/TERT1 cells detected by ELISA.

Exposure duration	Treatment	Average adducts/10 ⁸ nucleotides \pm SEM, $n = 3$
18 h B[a]P	DMSO 0.01 µM B[a]P 0.1 µM B[a]P 1 µM B[a]P	$\begin{array}{c} 0.0 \pm 0.15 \\ 0.58 \pm 0.37 \\ 9.92 \pm 2.66 \\ 36.34 \pm 9.14 \end{array}$
24 h B[a]P	DMSO 0.01 µM B[a]P 0.1 µM B[a]P 1 µM B[a]P	$\begin{array}{c} 0.0 \pm 0.21 \\ 0.0 \pm 0.82 \\ 5.72 \pm 1.43 \\ 59.75 \pm 17.03 \end{array}$
24 h 1 μM Cd × 24 h B[a]P	DMSO 0.01 µM B[a]P 0.1 µM B[a]P 1 µM B[a]P	0.0 ± 0.16 1.18 ± 0.14 7.88 ± 1.33 17.28 ± 1.30

significantly increased at $1 \mu M B[a]P$ in comparison to DMSO vehicle control, 0.01 and 0.1 $\mu M B[a]P$ at both 18 and 24h post-exposure. Fewer adducts were detected after 24h of exposure to 0.1 $\mu M B[a]P$ in comparison to the same concentration at 18h although the difference was not statistically significant (Fig. 1, Table 2).

In order to assess the ability of Cd to alter adduct formation and persistence, adducts were analyzed under conditions of Cd and B[a]P co-exposure. Cells were exposed to Cd alone for 18 and 24 h to verify the absence of adducts. There were no BPDE-DNA adducts found above background at either time point after Cd exposure (data not shown). For co-exposure, cells were exposed to a non-cytotoxic concentration of Cd, 1 µM, for 24 h. Cytotoxicity of each compound was based on previous work [21]. After 24 h, cells were exposed to DMSO vehicle control or appropriate concentrations of B[a]P for 24h. Adducts detected in groups exposed to lower concentrations of B[a]P remained relatively unchanged between treatment groups. However, cells exposed to 1 μ M Cd \times 1 μ M B[a]P demonstrated significantly reduced levels of adducts in comparison to 1 µM B[a]P alone at either time point (Fig. 1, Table 2).

3.2. Exposure to Cd increases expression of NRF2 responsive genes

Gene expression changes of the NRF2 responsive genes, glutamate-cysteine ligase, catalytic subunit (GCLC), heme oxygenase 1 (HMOX1), and NAD(P)H dehydrogenase, quinone 1 (NQO1), were examined after exposure to determine if Cd alone or Cd and B[a]P together appeared to induce an antioxidant response that may increase BPDE detoxification and reduce BPDE-DNA adduct formation under co-exposure conditions at $1 \mu M Cd \times 1 \mu M B[a]P$. While GCLC was detected, there was no change among treatment groups after a 24-h exposure to Cd (Fig. 2A). After 24 h of exposure to 0.1, 1, and 10 µM Cd, there was nearly a 3-fold increase in HMOX1 at 10 µM Cd in comparison to untreated cells and all other concentrations (Fig. 2B). Additionally, all concentrations of Cd showed approximately a 2-3-fold increase in NQO1 over that of untreated cells (Fig. 2C).

Twenty-four hours of B[a]P exposure did not increase gene expression of GCLC, HMOX1, or NQO1 (Fig. 3A–C). All 🚥 18 Hrs B[a]P 🔲 24 Hrs B[a]P 🗔 24 Hrs Cd x 24 Hrs B[a]P



Fig. 1. BPDE-DNA adducts are formed and detected in RPTEC/TERT1 cells after exposure to B[a]P but reduced under co-exposure to 1 μ M Cd and 1 μ M B[a]P. After 18 and 24 h of treatment with B[a]P a significant increase in the number of adducts was detected at 1 μ M B[a]P. Treatment of cells for 24 h with 1 μ M Cd before a 24 h exposure to concentrations of B[a]P showed a significant decrease in adducts detected at 1 μ M B[a]P in comparison to B[a]P alone. Bars represent average adducts/10⁸ nucleotides (*n*=3)±SEM. *Significant difference from 1 μ M B[a]P at 18 h, *p* < 0.01, *Significant difference from 1 μ M B[a]P at 24 h, *p* < 0.01.

genes were detected at basal levels by real time PCR. However, co-exposure significantly increased gene expression of all three genes at the highest concentration of $1 \mu M$ Cd $\times 1 \mu M B[a]P$ over vehicle control and other co-exposure groups. *GCLC* gene expression was increased by approximately 2-fold, *HMOX1* gene expression was increased by approximately 3-fold, and *NQO1* gene expression was increased by approximately 4-fold (Fig. 4A–C). This suggests that co-exposure, under these conditions, triggers a stronger transcriptional antioxidant response than Cd alone.

3.3. Cd and Cd \times B[a]P exposure increase total glutathione levels in RPTEC/TERT1 cells

Total glutathione was measured after Cd exposure for 24 h and after co-exposure with B[a]P. Total glutathione levels were approximately double in cells treated with 1 and 10 μ M Cd in comparison to untreated groups after 24 h. Cells treated with 0.1 μ M Cd exhibited a slight increase in total glutathione, but this increase was not statistically significant. Total glutathione was also significantly increased in cells pre-treated with 1 μ M Cd for 24 h followed by exposure to 0.01, 0.1, and 1 μ M B[a]P for 24 h (Fig. 5). This supports our hypothesis that Cd induces a biochemical antioxidant response, and co-exposure to Cd and B[a]P results in a substantial increase in reduced glutathione (GSH) levels possibly greater than those induced by Cd alone.

4. Discussion

The multifaceted effects that have environmental mixtures on the human body have been notoriously problematic to resolve. For over a decade, scientists have faced

exceedingly difficult challenges in chemical toxicological research when studying chemical mixtures designed to address gaps in our knowledge [27]. However, the importance of pursuing chemical mixture experiments continues to increase with the rise in diseases that have no clear genetic predisposition. Assessments by the American Cancer Society credit heritable mutations in the development of only 5% of all cancers [3]. Likewise, current evidence suggests that an overwhelming 90% of human disease burden, especially degenerative conditions, can be attributed to environmental factors such as exposure, lifestyle, and diet [28–31]. Humans encounter mixtures of chemical compounds daily and throughout their lives; however, relatively little research to date has aimed to address the differential effects that mixtures have on molecular and mechanistic endpoints in comparison to studies focused on individual chemicals and compounds. A recent review on PAH mixtures toxicology illustrates these complexities but also provides strong rationale for approaching these issues [32]. Biological processes and molecular factors that counteract the development of cancer (e.g. increased antioxidant or detoxification capacity) must be studied in the context of exposure to these mixtures. Environmental toxicants that interfere with efficient processing and accurate repair of DNA adducts may increase the mutagenicity of other toxicants by decreasing DNA repair capacity. In this context, such environmental toxicants function as cocarcinogens.

In an effort to characterize the effects of a simple binary mixture on renal proximal tubule cells, we have examined cellular responses of the RPTEC/TERT1 immortalized cell line to B[a]P and Cd, two distinctly different carcinogens. Our previous studies with this cell line have demonstrated its sensitivity to both B[a]P and Cd as well as compound-specific responses [21]. Here, we confirm





Fig. 2. RPTEC/TERT1 cells respond to 24h Cd exposure by upregulating *HMOX1* and *NQ01* but not *GCLC*. After 24h of treatment with Cd at various concentrations, RPTEC/TERT1 cells showed no change in (A) *GCLC* at any concentration. There was a significant increase in gene expression at the highest concentration, 10 μ M Cd, of (B) *HMOX1* and (C) *NQ01*. Bars represent mean fold expression (n=3) \pm SEM. All genes of interest were normalized to *ACTB*. 0 μ M, where denoted, was set as 1. *Significant difference from 0 μ M Cd, p < 0.01, #Significant difference from 10 μ M Cd, p < 0.01, and *Significant difference from 0 μ M Cd, p < 0.05.

Fig. 3. Twenty-four hours of B[a]P exposure does not induce changes in *GCLC*, *HMOX1*, or *NQO1*. None significantly differ. Bars represent mean fold expression $(n=3)\pm$ SEM. All genes of interest were normalized to *ACTB*. DMSO, where denoted, was set as 1.

Total Glutathione (µM)



1µM Cd x B[a]P Concentrations

Fig. 4. Co-exposure conditions with Cd and B[a]P result in upregulation of *GCLC*, *HMOX1*, and *NQO1* in RPTEC/TERT1 cells. Cells were exposed to 1 μ M Cd for 24 h followed by a 24 h exposure to B[a]P at different concentrations. One micromolar Cd was previously determined to be non-cytotoxic to RPTEC/TERT1 cells after 24 h. RPTEC/TERT1 cells demonstrated significant upregulation of (A) *GCLC*, (B) *HMOX1*, and (C) *NQO1* after exposure to 1 μ M Cd × 1 μ M B[a]P. Bars represent mean fold expression (n = 3) ± SEM. All genes of interest were normalized to *ACTB*. DMSO, where denoted, was set as 1. *Significant difference from DMSO, p < 0.01 and #Significant difference from 1 μ M Cd × 1 μ M B[a]P, p < 0.01.





Treatment

Fig. 5. Cd exposure increases total glutathione in RPTEC/TERT1 cells. Total glutathione levels were significantly increased after 24 h of exposure to 1 and 10 μ M Cd. Total glutathione levels were also significantly increased after exposure to 1 μ M Cd for 24 h followed by a 24 h exposure to B[a]P concentrations. Bars represent mean total glutathione levels (n = 3) ± SEM. *Significant difference from 0 μ M Cd, p < 0.05.

the metabolism of B[a]P to metabolites which form DNA adducts under these conditions. We detected BPDE-DNA adducts at 18 and 24h post-exposure to B[a]P alone. At 24 h post-exposure, there were fewer adducts detected at intermediate concentrations, 0.01 and 0.1 µM B[a]P, than at 18 h post-exposure. While the numbers of adducts were not significantly reduced, the decrease at these concentrations suggests that there may be some removal or repair of the initial adducts. However, the limited sensitivity of the ELISA method at the lower concentrations of B[a]P tested in these experiments makes these suggestions speculative. At the highest concentration of $B[a]P(1 \mu M)$ tested alone, we found that significantly greater adduct levels remained at both 18- and 24-h time points. At this concentration of B[a]P, detoxification and repair mechanisms may have been unable to process the amount of B[a]P metabolites in the examined time period. There may a threshold effect, possibly short-term but in excess of 24 h, in which bioactivation exceeds detoxification and repair, which generates BPDE-DNA adducts at a rate greater than the rate at which DNA repair processes can remove them. This conclusion is warranted especially if B[a]P treatments alone in this experiment do not induce an antioxidant response, increase detoxification capacity, or increase DNA repair capacity. In contrast, when cells were pre-treated with 1 μ M Cd, BPDE-DNA adducts formed after 1 μ M B[a]P exposure were significantly reduced. This effect was not observed to this degree at other concentrations of B[a]P after Cd pre-treatment. It is possible that CYP-mediated biotransformation at the lower concentrations of B[a]P was occurring without exceeding detoxification and/or DNA repair capabilities. This would allow cellular mitigation of BPDE-DNA adducts without an increase in induction as supported, in part, by our previous work [21]. However, as mentioned previously, the sensitivity of the ELISA method at lower concentrations of B[a]P used in these experiments may not be adequate to distinguish statistically significant differences in BPDE-DNA adduct levels among the treatment and co-treatment groups with adequate precision. We suspect that B[a]P is metabolized at the lower concentrations, but more sensitive analytical methods are necessary to discriminate significant differences in adduct formation and persistence based on treatment regimen. We suggest that future experiments be designed to address such experimental possibilities and statistical power limitations.

The reduction in adducts under co-exposure at the highest concentration of B[a]P may be a function $Cd \times B[a]P$ priming the detoxification system through the NRF2 antioxidant pathway. These effects appear to result in increased levels of glutathione and increased inactivation or detoxification of BPDE prior to adduct formation. Future measurements of BPDE-DNA conjugates in conjunction with glutathione levels should be used to confirm this supposition. B[a]P exposure alone, under these conditions, does not appear to induce any such response. While it appears that Cd alone induces a significant antioxidant response resulting in increased levels of glutathione, Cd and B[a]P together at the highest concentrations tested induce an even more robust response. This response to both Cd and B[a]P in our experiments seems to mitigate DNA adduction formation through enhanced detoxification capacity. Our previous work does not support one alternative explanation that Cd inhibits the formation of BDPE through CYP feedback inhibition [21].

The binding of B[a]P to the aryl hydrocarbon receptor increases the expression of xenobiotic response element (XRE) genes and their encoded enzymes, which are responsible for metabolizing B[a]P to reactive intermediates [33,34]. These reactive intermediates, along with reactive oxygen species (ROS) from heavy metals, can increase the transcription of antioxidant response element (ARE) genes through NRF2 binding [35,36]. Activation of this gene battery may be responsible for metabolite detoxification. We suspect that this process, under our experimental coexposure conditions, reduced level of adducts detected at 1 µM B[a]P following 1 µM Cd pre-treatment. We found the expression of NRF2-targeted genes, GCLC, NQO1, HMOX1, to be significantly increased under experimental conditions coinciding with the most significant reduction in BPDE-DNA adducts under co-exposure. Our results are similar to *in vivo* studies which have discovered that priming the NRF2 system decreases the levels of adducts formed after B[a]P exposure. Nrf2 knockout mice develop more tumors than wild-type mice when treated with B[a]P alone. When mice are given a Nrf2 activator with B[a]P, wild-type mice develop half as many tumors. However, tumor reduction is not seen in Nrf2 knockout mice given a Nrf2 activator with B[a]P exposure [35]. In other studies including transformed kidney cell lines from humans and rats, Cd has been shown to induce the NRF2 pathway through an oxidative stress mechanism [37-39]. Future studies in the RPTEC/TERT1 cell line should consider the application of a NRF2 inhibitor to mimic in vivo Nrf2 knockout conditions and further verify the responses seen after B[a]P exposure. Several NRF2 activating agents, both natural and synthetic, have been examined as chemoprotectives for chronic disease and overall cell health. Flavonoids, for example, are naturally occurring antioxidants found in cruciferous vegetables, apples, and onions. They have been shown to increase NRF2 mediated expression of *NQO1* and *GST*. Additionally, naturally occurring phytochemicals such as chalcones and coumarins have been shown to act similarly by inducing NRF2 expression of *NQO1* and *GST* to act as anti-inflammatories and antioxidants [40]. Similarly, bardoxolone methyl, a synthetic NRF2 activator derived from natural antioxidants, has been successful in increasing kidney function and halting the progression of renal injury in Phase 2 clinical trials in patients with chronic kidney disease [41].

Our goals were to measure responses in RPTEC/TERT1 cells to defined, non-cytotoxic mixtures of two distinctly different toxicants. While our results suggest an antagonistic, or perhaps a hormetic effect, on the endpoint examined, DNA adduct formation and persistence, further studies are necessary to explore these results and determine effects on other downstream biomarkers. Of particular interest is the mutagenicity of BPDE-DNA adducts under such conditions of co-exposure especially at lower, environmentally relevant concentrations. We hypothesize that increased detoxification capacity is responsible for the reduced levels of BPDE-DNA adducts which may protect cells from these premutagenic lesions. This could be interpreted as a hormetic effect [42]. It is also possible that, while detoxification capacity is increased, subsequent DNA repair is inhibited by the presence of Cd. Cd inhibition of DNA repair may promote repair mistakes or error-prone translesion synthesis of the remaining adducts leading to a relative increase in mutagenicity under these conditions (for a review, see [43]).

Apparent NRF2 activation and the increased total glutathione levels found in Cd and co-exposure groups are evidence of cellular oxidative stress. Cd and Cd compounds are Group 1 carcinogens and are known to cause cancer in humans [13]. DNA damage caused indirectly by Cd, such as oxidative insult and repression of DNA damage repair, must be considered in mutational investigations. Quantifying the levels of PAH-DNA adducts in human studies can serve as a biomarker for exposure as well as provide information on an individual's DNA repair capacity and mutagenic risk [24]. However, it would be ideal to also measure mutation frequency to confirm mutagenic potential as a function of adduct formation under controlled conditions with *in vitro* and *in vivo* models to better represent and understand mechanisms by which mixtures impact humans.

We have conducted these studies in the RPTEC/TERT1 human immortalized cell line because they were derived from a normal, healthy individual, have proven to be metabolically competent, and exhibit canonical responses similar to human kidney cells when exposed to the selected environmental toxicants. However, we acknowledge the difficulties in carrying out robust, controlled experimentation on chemical mixtures. Due to the complicated nature of mixtures toxicology, it remains challenging to extrapolate the results obtained in this or any *in vitro* or *in vivo* model to actual human risk. Nevertheless, *in vitro* models permit higher throughput screening of many compounds and mixtures. As they improve to better model the tissue of interest, *in vitro* models will prove extremely valuable in mixtures toxicology. Advancements in high throughput screening have allowed scientists to begin to elucidate increasingly complex mechanisms and interactions inherent to mixtures toxicology. As research continues on both environmental and genetic components of disease, the causes of conditions such as cancer are becoming more recognized as environmentally mediated. Thus, it can be hypothesized that the majority of genetic changes resulting in cancer are acquired over a lifetime through one's interactions with the environment. Therefore, it is critical to better understand the molecular processes that contribute to the initiation of cancer.

Transparency document

The Transparency document associated with this article can be found in the online version.

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