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Mechanistic relationships between hepatic genotoxicity and carcinogenicity in male B6C3F1 mice treated with polycyclic aromatic hydrocarbon mixtures

Tracie D. Phillips,

College of Veterinary Medicine and Biomedical Sciences, Texas A&M University, College Station, TX 77843

Molly Richardson,

School of Rural Public Health, Texas A&M University Health Science Center, College Station, TX 77843

Yi-Shing Lisa Cheng,

Baylor College of Dentistry, Texas A&M University Health Science Center, Dallas, TX 75246

Lingyu He,

School of Rural Public Health, Texas A&M University Health Science Center, College Station, TX 77843

Thomas J. McDonald,

School of Rural Public Health, Texas A&M University Health Science Center, College Station, TX 77843

Leslie H. Cizmas, Stephen H. Safe,

College of Veterinary Medicine and Biomedical Sciences, Texas A&M University, College Station, TX 77843

Kirby C. Donnelly#,

School of Rural Public Health, Texas A&M University Health Science Center, College Station, TX 77843

Fen Wang,

Institute of Biosciences and Technology, Texas A&M University Health Science Center, Houston, TX 77030

Bhagavatula Moorthy, and

Department of Pediatrics, Baylor College of Medicine, Houston, TX 77030

Guo-Dong Zhou*

Institute of Biosciences and Technology, Texas A&M University Health Science Center, Houston, TX 77030

^{*}Corresponding author at: Institute of Biosciences and Technology, Texas A&M University Health Science Center, Houston, TX 77030, USA. Tel: 1-713-677-7433; Fax: 1-713 677 7784.gzhou@ibt.tamhsc.edu (G.D. Zhou).

#Deceased

School of Rural Public Health, Texas A&M University Health Science Center, College Station, TX 77843

Abstract

The genotoxicity of a complex mixture [neutral fraction (NF)] from a wood preserving waste and reconstituted mixture (RM) mimicking the NF with seven major polycyclic aromatic hydrocarbons (PAHs) and benzo(a)pyrene (BaP) was investigated by determining DNA adducts and tumor incidence in male B6C3F1 mice exposed to 3 different doses of the chemical mixtures. The peak values of DNA adducts were observed after 24 h and the highest levels of PAH-DNA adducts were exhibited in mice administered NF+BaP, and the highest tumor incidence and mortality were also observed in this group. DNA adduct levels after 1, 7, or 21 d were significantly correlated with animal mortality and incidence of total tumors including liver, lung, and forestomach. However, only hepatic DNA adducts after 7 d significantly correlated with liver tumor incidence. Most proteins involved in DNA repair including ATM, pATR, Chk1, pChk1, DNA PKcs, XRCC1, FANCD2, Ku80, Mre11 and Brca2 were significantly lower in liver tumor tissue compared to non-tumor tissue. Expression of proteins involved in apoptosis and cell cycle regulation were also significantly different in tumor vs non-tumor tissues and it is possible that PAH-induced changes in these gene products are important for tumor development and growth.

Keywords

Polycyclic aromatic hydrocarbon mixtures; DNA adducts; Tumor incidence; Reverse Phase Protein Array; ³²P-postlabeling assay; Protein expression

Introduction

Cancer is a major public health problem in United States and worldwide and it is estimated that 1,665,540 new cancer cases will be diagnosed and 585,720 deaths from cancer will occur in the United States in 2014 (Siegel et al. 2014). Epidemiological and experimental data show that some cancers are caused by environmental factors, including exposure to complex mixtures such as carcinogenic PAHs from cigarette smoke, air and water pollution. Several PAHs, including BaP, have been listed by the U.S. Environmental Protection Agency as probable human carcinogens (Agency 2003).

PAHs are precarcinogens that are converted into carcinogens by drug metabolizing enzymes and the resulting reactive metabolites alkylate DNA or induce DNA damage that play essential roles in the initiation phase of carcinogenesis (Hecht 2003; Wogan et al. 2004). For example, DNA adducts can cause miscoding events in critical genes (Hecht 2012). Many DNA adducts serve as biomarkers of carcinogenic PAH exposure and cancer risk (Phillips and Venitt 2012; Poirier 1997; Urban et al. 2012). Significant positive linear correlations between levels of PAH-DNA adducts and cancer incidence have been reported in both epidemiological studies (Agudo et al. 2012; Tang et al. 2001; Veglia et al. 2008) and animal experiments (Culp et al. 1998). However, negative correlations between PAH-DNA adducts and tumor incidence have also been observed (Saieva et al. 2011; Siddens et al. 2012) and

the mechanic relationships between carcinogen-DNA adducts and tumor formation are still not fully understood.

Gene/protein expression involved in DNA repair, apoptosis, and cell cycle regulation play very important roles in chemical carcinogenesis (Henkler et al. 2012; Suzuki et al. 2012) and treatment with BaP alters expression of genes involved in xenobiotic metabolism, DNA repair, apoptosis, and cell cycle regulation (Hamouchene et al. 2011). Initial recognition of DNA adducts is a key event regulating DNA damage repair (Sugasawa 2011) and this includes nucleotide excision repair (NER). BaP-DNA adducts in liver and esophagus were significantly increased in DNA repair deficient mice compared to wild-type mice (John et al. 2012). It was reported that apoptosis may also be associated with decreased BaP adduct levels (Zhou et al. 2010) and failure to induce apoptosis in human bronchial epithelial cells exposed to cigarette smoke resulted in enhanced neoplastic transformation (Du et al. 2012).

PAH mixtures are known human carcinogens and the potency of several individual compounds has been defined in animal studies, however, minimal information is available on potential interactions of complex mixtures of PAHs. Mixture interactions may alter the absorption, distribution, metabolism or excretion of chemicals and some chemicals in PAH mixtures may either inhibit or promote chemical carcinogenesis. Therefore, these interactions may effectively increase or decrease the potential toxicity or genotoxicity of the mixture (Tarantini et al. 2011).

The current study investigated the genotoxicity and carcinogenicity of a single PAH---BaP, BaP plus a complex PAH mixture (NF) isolated from a wood preserving waste, and a reconstituted mixture (RM) containing BaP and six other major carcinogenic PAHs. The relationships between genotoxicity and carcinogenicity were studied and expression of proteins involved in DNA repair, apoptosis, and cell cycle regulation was also investigated in tumor and normal liver tissues.

Materials and Methods

Chemicals

Benz(a)anthracene (BaA), BaP, benzo(b)fluoranthene (BbF), benzo(k)fluoranthene (BkF), chrysene, and dibenz(a,h)anthracene (DA)) were obtained from Sigma-Aldrich (St. Louis, MO). Indeno(1,2,3-c,d)pyrene (IP) was purchased from Absolute Standards (Hamden, CT). Materials for DNA extraction (Gupta 1984; Randerath et al. 1981; Reddy and Randerath 1986; Zhou et al. 1999), and ³²postlabeling analysis (Phillips and Arlt 2007; Randerath et al. 1999; Reddy and Randerath 1986; Zhou et al. 2004) have been reported previously.

Extraction and PAH fractionation from complex chemical mixture

Wood preserving waste (WPW), a non-aqueous phase liquid was initially collected from a contaminated aquifer beneath a wood treatment plant. A liquid-liquid separation technique was used to separate this material into acid, base, and neutral fractions following the USEPA 3650B method ((EPA) 1996). A 30 ml volume of WPW oil, 60 ml of methylene chloride (CH₂Cl₂), and 60 ml sodium hydroxide (NaOH), pH 12, were added to a 2,000 ml separating funnel and shaken for 2 min. Once phases were separated, the upper aqueous

layer (acid fraction) was removed, and the lower organic layer (base/NF) was extracted two more times. One volume of base/NF, 3 volumes of $1N\ H_2SO_4$ and $1.5\ volumes$ of CH_2Cl_2 were shaken in a new separating funnel for 2 min. After the phases were separated, the organic phase was removed and aqueous phase was extracted one more time with CH_2Cl_2 (same volume as first extraction). The two organic layers were combined to form the NF. The NF contains major carcinogenic PAHs (Figure 1), including BaA, BaP, BbF, chrysene, BkF, DA, and IP.

Preparation of a reconstituted mixture

Based on the composition of PAHs in the NF, seven major carcinogenic PAHs, which are defined as class B2 carcinogens by USEPA, were selected for a RM. The concentrations of each PAH in the RM were simulated to those in the NF. The final concentrations for animal treatment are listed in Table 1.

Chemical analysis of the complex mixture

The analysis of the PAHs in the sample extracts was accomplished using a modified USEPA Method 8270C (GC/MS-SIM) as reported previously (Cizmas et al. 2003). The quantification of the analytes was based on authentic standards and internal standards. The identification of the analytes was based on retention times and mass spectra (primary and confirmation ions).

Animals

B6C3F1 male mice (21 days old, 7–11 g) were obtained from Harlan Sprague Dawley (Houston, TX). Animals were housed in the Program for Animal Resources of the Institute of Biosciences and Technology, Texas A&M University System Health Science Center. All procedures were conducted with the approval of the Institutional Animal Care and Use Committee of the Texas A&M University System Health Science Center (Protocol Number: 06001-R1).

Animal experiments

Mice were housed in cages in a temperature and humidity controlled facility with a daily photoperiod of 12 h light and 12 h dark. Animals were acclimatized for a week to the new environment, and then the mice were randomly allocated to ten experimental groups with 35 mice for each group. Mice had free access to water and fresh diet was provided in clean food bowls daily. The animals were subjected to a single intraperitioneal (i.p.) injection with BaP or a mixture of seven PAHs. Table 1 shows doses of PAHs applied in animal experiments. These chemicals, except for BaP, and their concentrations were similar to the residues extracted from the dense non-aqueous phase liquid collected at a wood preserving waste (WPW) Superfund Site (Phillips 2006). These chemicals were dissolved in DMSO:Corn oil (50:50). The vehicle volume for ip injection was about 7 μl per g body weight. Five animals were sacrificed by CO₂ asphyxiation and cervical dislocation at 1, 7, and 21 d after the PAH treatment. Five mice served as a control and therefore, were injected with the vehicle. Twenty to thirty mice for each group were maintained in the same environment for 280 d to observe tumor formation. Liver tissues were collected and were frozen in liquid nitrogen

immediately and then stored at -8° C until extractions of DNA and protein. Part of each tissue was fixed in paraformaldehyde (4%) for histological examination.

Analyses of DNA adducts

DNA extraction and ^{32}P -postlabeling analyses were performed as in previous experiments (Phillips and Arlt 2007; Zhou et al. 2005). DNA was isolated by solvent extraction combined with enzymatic digestion of protein and RNA (Gupta 1984; Randerath et al. 1981) and stored at -80°C until analysis. The nuclease P1-enhanced bisphosphate version of the ^{32}P -postlabeling method (Reddy and Randerath 1986) was used for analysis, with modifications of the chromatographic conditions (Mabon et al. 1996). Briefly, DNA (10 µg) was enzymatically degraded to normal (Np) and modified (Xp) deoxyribonucleoside 3'-monophosphates with micrococcal nuclease and spleen phosphodiesterase at pH 6.0 and 37°C for 3.5 h. After treatment of the mixture with nuclease P1 to convert normal nucleotides to nucleosides, modified nucleotides (Xp) were converted to 5'- ^{32}P -labeled deoxyribonucleoside 3',5'-bisphosphates (pXp) by incubation with carrier-free [γ - ^{32}P]ATP and polynucleotide kinase.

Radioactively labeled digests were applied to modified polyethyleneimine (PEI)-cellulose thin layers and chromatographed overnight (15–16 h) with 2.3 M sodium phosphate, pH 5.75 (D1), to purify bulky adducts. Labeled PAH-adducts retained in the lower (L, 2.8×1.0 cm) part of the D1 chromatogram were, after brief autoradiography on Cronex 4 X-ray film, each contact-transferred to individual acceptor sheets and resolved by two-dimensional thin-layer chromatography (TLC). The bulky DNA adducts were separated with 3.82 M lithium formate, 6.75 M urea, pH 3.35 and 0.72 M sodium phosphate, 0.45 M Tris–HCl, 7.65 M urea, pH 8.2 in the first (D3) and second (D4) dimensions, respectively. 32 P-labeled DNA adducts were visualized by screen-enhanced autoradiography at -80°C using Kodak XAR-5 film or with the aid of an InstantImager (Packard Instruments) (Zhou et al. 1999). Appropriate blank count rates were automatically subtracted by the instrument from sample values. The extent of covalent DNA modification was calculated from corrected sample count rates, the amount of DNA assayed (expressed as pmol DNA monomer units or DNA-P), and the specific activity of $[\gamma - ^{32}P]$ ATP according to the following formula:

$$Adducts/10^{9}normal\ nucleotides = \frac{DNA\ modification(s)[cpm]}{DNA-P[pmol] \times Spec.act._{ATP}[cpm/pmol]}$$

Quantitative data represented minimum estimates because 100% recovery presumably was not achieved.

Reverse phase protein array (RPPA) assay

The assay of RPPA (Byers et al. 2012; Malinowsky et al. 2013) was performed in the Functional Proteomics Reverse Phase Protein Array Core Facility at MD Anderson Cancer Center, Houston, TX. Briefly, tissue was lysed with RPPA lysis buffer and diluted in five 2-fold serial dilutions in dilution buffer (lysis buffer containing 1% SDS). Serial diluted lysates were arrayed on nitrocellulose-coated FAST slides (Whatman, Inc, Piscataway, NJ) by an Aushon 2470 Arrayer (Aushon BioSystems, Billerica, MA). Each slide was probed

with a validated primary antibody plus a biotin-conjugated secondary antibody. The signal was amplified using a Dako Cytomation—catalyzed system (Dako North America, Inc. Varpinteria, CA) and visualized by DAB colorimetric reaction. The slides were scanned, analyzed, and quantified using the customized-software Microvigene (VigeneTech Inc. North Billerica, MA) to generate spot intensity (He et al. 2012; Tibes et al. 2006).

Histological examination

At 280 days of age surviving mice were euthanized by CO₂ asphyxiation, followed by cervical dislocation, and a number of tissues (liver, lung, and forestomach) examined first by gross necropsy and then fixed in 10% formalin. Fixed tissues were routinely processed to paraffin blocks, and hematoxylin and eosin-stained sections were analyzed by a board-certified histopathologist as previously described (Shorey et al. 2012).

Statistical analysis

The relationships between values of PAH-DNA adducts at 1, 7, and 21 d and the mortality and tumor incidence were analyzed by linear regression analysis (Zar 2009). Statistical analysis of the data from individual mice was performed by using the unpaired and paired Student's *t*-tests (Zar 2009).

Results

Profiles and levels of hepatic PAH-DNA adducts

For acute experiments, animals were terminated at 1, 7, and 21 d after PAH treatment. Profiles and levels of hepatic DNA adduct formation were determined. Representative profiles of hepatic DNA adducts in mice treated with BaP, BaP+NF, or RM are displayed in Figure 2. Typical patterns of PAH-DNA adducts are exhibited in all three groups with only slight differences. Total levels of PAH-DNA adducts for all chemical groups at three time points, and three different doses are summarized in Table 2. PAH-DNA adducts formed quickly within one day after PAH treatment. High levels of adducts were observed after 24 h and they gradually decreased after 7 and 21 d. The NF+BaP groups showed higher levels of hepatic DNA adducts compared to the BaP or RM groups. Adduct levels at high doses were 81.6 ± 23.4 , 276.5 ± 19.0 , and 94.6 ± 14.9 in 10^9 normal nucleotides, respectively, for the BaP, NF+BaP, and RM groups. Analysis of all nine treatment groups (three doses and three time points) and comparison of DNA adduct levels between the NF+BaP and RM treatment by paired-sample t-test (Zar 2009) showed that DNA adduct levels were significantly higher among the mice dosed with NF+BaP than those dosed with RM (P=0.05). DNA adducts were also detected in mice 280 d after treatment with PAHs, however, hepatic PAH-DNA adducts were not observed at this time point (data not shown).

Histological observations

Acute histopathological changes were observed in the high dose RM group (Figure 3A, panel b). PAHs caused liver damage 7 d after treatment compared to the control group (Figure 3A, panel a). These results suggest that the high dose of carcinogenic PAHs not only induced genotoxicity, but also caused toxicity-based histological alterations.

Forty weeks after i.p treatment with PAHs, tumor formation was observed in the liver, lung, and forestomach of B6C3F1 mice. The tumor incidence was greater in liver than in lung or forestomach (Table 3). Upon dissection, gross liver tumor formation was observed (Figure 3B, panel c) and histological analysis indicated that most liver tumors were hepatomas (Figure 3B, panel e). High hepatic tumor incidence was observed in all three chemical groups.

The relationships between DNA adducts and mortality

Mortality was measured by the number of dead animals observed prior to the end (280 d) of the study. The groups that were challenged with NF+BaP had higher mortality than the groups dosed with either BaP or the RM. Interestingly, levels of DNA adducts at 1, 7 and 21 d were significantly correlated with animal mortality (Figure 4). The higher the levels of hepatic DNA adducts displayed at early time points, the higher was the mortality. Hepatic DNA adduct values after 1 d exhibited the most significant correlation with mouse mortality (P<0.001) compared to 7 (P<0.002) and 21 d (P<0.02). These results suggested that PAHs induced acute toxicity including genotoxicity could be associated with increased cancer related mortality. PAH-DNA adducts could be also significant biomarkers for such mortality.

The relationships between DNA adducts and tumor incidence

Tumor incidence for all groups was determined based on histological observation. Linear regression between PAH-DNA adducts and tumor incidence was determined and hepatic DNA adduct levels after 7 d significantly correlated with liver tumor incidence with P-values being <0.02 (Figure 5A). Interestingly, hepatic DNA adducts may be not only biomarkers of liver tumors, but also biomarkers for tumors formed in other organs. Figure 5B shows that levels of hepatic DNA adducts after 1, 7, and 21 d were significantly correlated with tumor incidence in multiple organs (liver, lung and forestomach). The P-values are <0.01, <0.005, and <0.05 for 1 d, 7, d and 21 d, respectively.

Comparisons of protein expression involved in DNA repair, apoptosis, and cell cycle regulation between tumor and normal tissues

A panel of 4 normal and 4 tumor tissues was profiled by RPPA to identify differences in expression of protein involved in chemical carcinogenesis such as DNA repair, apoptosis, and cell cycle regulation. Protein lysate was collected from hepatic tumor samples and normal liver tissues and it was evident that most DNA repair proteins including ATM, pATR, Chk1, pChk1, DNA PKcs, XRCC1, FANCD2, Ku80, Mre11 and Brca2 were expressed at significantly lower levels in tumor vs normal tissues (Table 4). Expression of proteins involved in apoptosis and cell cycle regulation were also significantly different in tumor vs non-tumor tissues. In tumor tissues, proteins PARP1, caspase3, caspase7, XIAP, p53, p-Rb (Ser807/811), and Cyclin B1 were downregulated. In contrast, Bcl-xL, Bax, pBad (Ser112), and P16/INK4A were upregulated.

Discussion

Environmental PAHs are formed by the incomplete burning of hydrocarbons such as coal, gas, oil, and wood. They are generated from smoking, automobile exhaust, industrial activities, forest fires, and even grilled meats and are always found as mixtures in the environment and in biological tissues (Guo et al. 2011). In our current studies, we found there were seven major carcinogenetic PAHs in a NF extracted from a WPW dense nonaqueous phase liquid. This complex PAH mixture was carcinogenic in vivo and similar results were previously reported for this mixture by Culp and coworkers (Culp et al. 1998). RM was prepared based on the chemical compositions and relative PAH concentrations in the NF (Table 1). However, the biological effects caused by these two mixtures were significantly different. Levels of DNA adducts, mortality, and tumor incidence in the NF groups were much higher compared to those in RM groups (Table 2). These results suggest that some PAHs or other unknown compounds that were present in the NF, but not in the RM, played an important role in the formation of DNA adducts and tumors induced by the NF. It was reported that PAHs in binary mixtures modulate the efficiency of BaP to form DNA adducts in human cells (Tarantini et al. 2011) and chromium (VI) exposure can greatly enhance the mutagenicity and cytotoxicity of PAHs by inhibiting the nucleotide excision repair (Hu et al. 2004).

Environmental factors including tobacco smoke, diet, and exposure to environmental and occupational carcinogens are believed to be responsible for many cancers (J.J. 2008), and for humans PAHs are among most abundant chemical carcinogens in terms of levels and exposure. Most cancers are still not curable and therefore, cancer prevention is an important anticancer strategy, particularly in high risk populations such as smokers and others exposed to environmental and occupational carcinogens. Therefore, biomarkers of tumor formation are critical for the early detection of disease-related changes and the prediction of cancer risk (Kyrtopoulos 2006).

The ³²P-postlabeling assay was first developed in 1982 (Gupta et al. 1982) and was modified with nuclease P1-enhanced bisphosphate in 1986 (Reddy and Randerath 1986). Because of its high sensitivity and requirement of less than 10 µg DNA, the ³²P-postlabeling assay (Phillips and Arlt 2007) has been applied in a number of human and animal studies of detection for DNA adducts. However, it is difficult for the ³²P-postlabeling assay to identify the structures of most carcinogen-DNA adducts. BPDE-dG (Jeffrey et al. 1977; Randerath et al. 1998) is one of structure identified carcinogen-DNA adducts (Fig. 3). Our results showed that the BPDE-dG adduct has very similar trends with total DNA adducts. In future studies, high resolution mass spectrometry (Klaene et al. 2012) or a combination of ³²P-postlabeling assay and mass spectrometry should be considered in detection and structure identification of DNA adducts (Farmer and Singh 2008).

It was reported that complex PAH mixtures have been associated with increased cancer mortality (Warshawsky et al. 1996). In the current study we observed that levels of PAH-DNA adducts detected at 1, 7 and 21 d after PAH treatment were significantly correlated with animal mortality. Peak values of DNA adducts detected after 1 d may be a suitable

predictor for general toxicity that is associated with the chronic adverse effects, including mortality.

The correlation between levels of PAH-DNA adducts and tumor incidence was also observed in this investigation. For example, hepatic PAH-DNA adducts detected after 7 d significantly correlated with liver tumor incidence at 280 d. Adduct levels after 1 and 21 d showed the trends of linear regressions but the correlations were not significant (0.05<P<0.10). Interestingly, hepatic PAH-DNA adducts were significantly correlated with total tumor incidence in multiple organs (liver, lung, and forestomach) at 280 d, with P values being <0.01, <0.005, and <0.05 for 1, 7 and 21 d, respectively. Adduct levels after 7 d showed the best correlation with both liver tumor incidence and tumor incidence of multiple organs. One possible mechanism for these correlations could be that DNA repair systems that play important role after initial DNA damage from PAH exposure may be compromised. PAH-DNA adducts rapidly activate nucleotide excision repair (Henkler et al. 2012). Adducted nucleotides that were not repaired or removed within 7 days could lead to mutations. Accumulation of mutations is an important step for the tumor development (Vogelstein and Kinzler 2004).

Data obtained from RPPA indicated that some proteins involved in DNA repair such as ATM, pATR, Chk1, pChk1, Xrcc1, Ku80, Mre11 and apoptosis (PARP1, caspase3, caspase7, XIAP) were significantly decreased in liver tumors compared to normal liver tissues of B6C3F1 mice (Table 4). These results suggest that genetic alterations, especially genes related to DNA repair and apoptosis pathways, may play important roles in PAHmediated carcinogenesis. It was reported (Shi et al. 2004) that women with reduced DNA repair capacity were at an increased risk for developing breast cancer. Attenuation of DNA repair and apoptosis could increase the formation of carcinogen-DNA adducts (John et al. 2012) followed by DNA mutations and tumor formation. Our results indicate that proteins involved in cell cycle regulation showed both upregulation and downregulation. Similar results were reported by Hamouchene and coworkers (Hamouchene et al. 2011) who showed that BaP exposure increased gene expression of GDF15, JUN, RGC32, and EGR1, but deceased HDAC4 and Scaper. Thus, PAH-induced alterations in DNA repair and apoptotic pathways may facilitate persistent DNA adduct formation and changes in expression of these proteins or their corresponding mRNAs and may serve as biomarkers of exposure and risk. Future development and validation of these biomarkers at early time point after exposure is warranted.

DNA adducts and their levels are important for tumor formation in rats after exposure to 2-nitrofluorene (Cui et al. 1995) and our results also confirmed that DNA adducts could be useful biomarkers for tumor risk in mice after exposure to complex PAH mixtures. However, there are many factors involved in chemical carcinogenesis between formation of DNA adducts and tumor formation. Figure 6 shows a number of proteins/genes involved in DNA repair, apoptosis, and cell cycle regulation that may play important roles in chemical carcinogenesis. Dynamic observations of these proteins/genes after exposure to chemical carcinogens may help predict future tumor risk, and facilitate understanding the mechanisms of chemical carcinogenesis and will be investigated in future studies.

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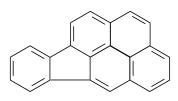
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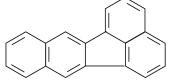
Chrysene

Benz(a)anthracene

Benzo(a)pyrene

Dibenz(a,h)anthracene





Indeno(1,2,3-cd)pyrene

Benzo(b)fluoranthrene

Benzo(k)fluoranthrene

Fig 1.

Chemical structures of the seven PAHs categorized as class B2 carcinogens according to the USEPA. These were present in the neutral fraction (NF) extracted from a wood preserving waste and were included in the reconstituted mixture (RM).

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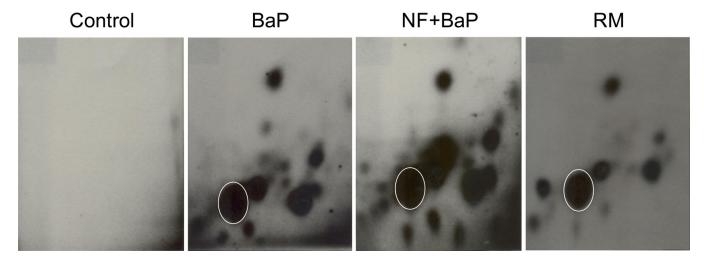
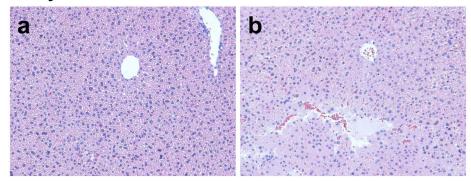


Fig 2. Typical representative profiles of ³²P-postlabeled PAH-DNA adducts detected in the livers of B6C3F1 mice. Animals were terminated at 1 d after PAH treatments. The circled spot is the benzo(a)pyrene-7,8-dihyrodiol-9,10-epoxide deoxy-guanosine (BPDE-dG) adduct.

A. Acute Toxicity



B. Tumor Formation



Normal Liver (10x)

Liver Hepatoma (10x)

Fig 3. Acute histological changes (A) in the liver of mice treated with reconstituted mixture (RM). Panel a: control, mouse treated with corn oil/DMSO (50:50). Panel b: mouse treated with the high dose of the reconstituted mixture (RM). Seven days after treatment, focal hemorrhage and tissue necrosis were observed. (B) PAH-induced hepatic tumor formation. Panel c: gross tumor formation observed in the high dose NF+BaP group during dissection. Panel d: liver tissue from control group treated with corn oil/DMSO (50:50). Panel e: liver hepatoma observed in the high dose NF+BaP group. All samples were obtained at 280 d after PAH treatment. NF, neutral fraction.

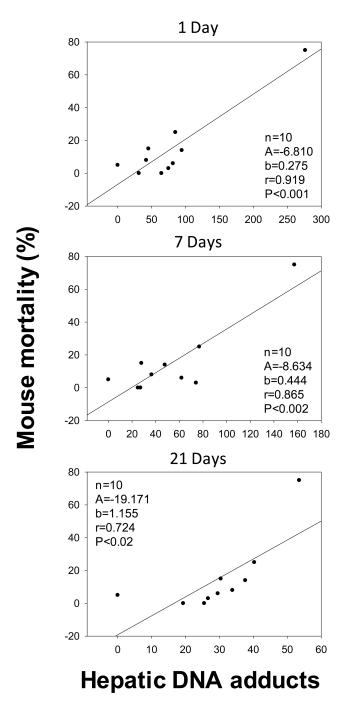
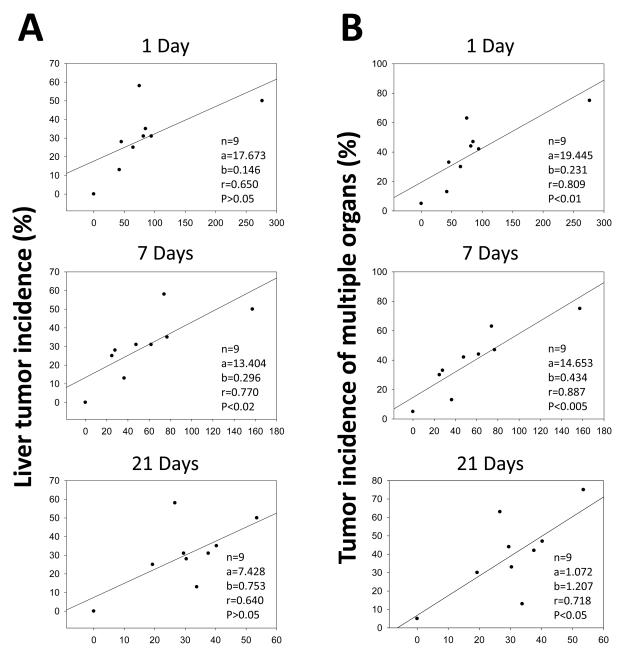


Fig 4. Correlations between hepatic DNA adduct levels (adducts in 10^9 normal nucleotides) at 1, 7, and 21 d, and mouse mortality prior to 280 d. Linear correlations were statistically significant for 1 d (P<0.001), 7 d (P<0.002) and 21 d (P<0.02).



Hepatic DNA adduct levels

Fig 5.

Correlation between hepatic DNA adduct levels (adducts in 10⁹ normal nucleotides) at 1, 7, and 21 d, and liver tumor incidence (A). Adduct levels detected at 7 d, but not at 1 or 21 d, were significantly correlated with liver tumor incidence. (B) Correlation between hepatic DNA adduct levels (adducts in 10⁹ normal nucleotides) at 1, 7, and 21 d, and tumor incidence in multiple organs (liver, lung, and forestomach). Levels of hepatic DNA adducts at 1, 7, and 21 d were all significantly associated with tumor incidence in multiple organs (liver, lung and forestomach).

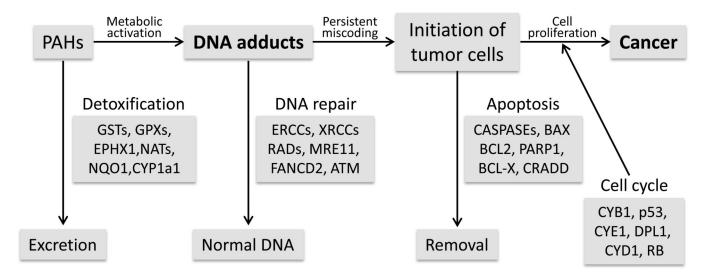


Fig 6.Possible mechanisms by which various proteins/genes involved in DNA repair, apoptosis, and cell cycle regulation play important roles in PAH-mediated carcinogenesis.

Table 1

Quantities of PAHs used in mouse treatments

ۇ ك			Ch	Chemical composition (µg/mouse)	osition (nou/gn	se)	
Group	dn	BaP	BaA	Chrysene	BbF	BkF	DA	Ш
Control		-	-	-	-	-	-	-
	Low	50	-	-	-	-	-	-
BaP	Middle	125	-	-	-	-	-	-
	High	375	-	-	-	-	-	-
	Low	50	0.85	58.0	0.34	0.21	0.02	0.11
NF+BaP	Middle	125	2.12	2.12	0.84	0.52	0.04	0.28
	High	375	5.3	5.3	2.1	1.3	0.1	<i>L</i> :0
	Low	50	0.62	<i>LL</i> '0	0.34	0.21	0.02	01.0
RM	Middle	125	1.56	1.92	0.84	0.52	0.04	0.24
	High	375	3.9	4.8	2.1	1.3	0.1	9.0

Benzo(a)pyrene (BaP), benz(a)anthracene (BaA), chrysene, benzo(b)fluoranthene (BbF), benzo(k)fluoranthene (BkF), dibenz(a,h)anthracene (DA), and indeno(1,2,3-c,d)pyrene (IP).

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Table 2
Total DNA adduct levels in liver of B6C3F1 male mice

Chemical	Group	DNA addu	ects in 10 ⁹ nucl	leotides*
Chemicai	Отопр	1 day	7 days	21 days
Control	CO:DMSO	ND**	ND**	ND**
	Low	64.6±6.5	25.0±2.2	19.3±1.9
BaP	Middle	74.9±10.8	74.1±34.8	26.7±3.4
	High	81.6±23.4	61.9±9.2	29.5±3.4
	Low	45.4±2.3	27.9±1.8	30.4±2.8
NF+BaP	Middle	85.1±14.5	76.9±6.5	40.3±5.9
	High	276.5±19.0	157.3±41.5	53.5±3.5
	Low	31.3±3.7	27.1±3.4	25.5±3.6
RM	Middle	42.1±11.2	36.5±4.5	33.8±5.2
	High	94.6±14.9	47.8±5.4	37.6±5.1

^{*} Mean±SEM, n=4;

^{**} Not detectable.

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Table 3

ale mice	Tumor incidence (%)	Liver Multiple organ
.1 ma	Tr	Live
dortality and tumor incidence in B6C3F1 male mice		Mortality (%)
ncide		u
and tumor i		Group
fortality a		Chemical

	(Tum	Tumor incidence (%)
hemical	Group	n	Mortality (%)	Liver	Multiple organs*
Control	CO:DMSO	20	5	0	5
	Low	20	0	25	30
ВаР	Middle	22	3	58	63
	High	22	9	31	44
	row	20	15	28	33
NF+BaP	Middle	20	25	35	47
	High	20	75	50	SL
	***MOT	26	0	-	-
RM	Middle	26	8	13	13
	High	28	14	31	42

* Multiple organs include liver, lung, and forestomach.

** The results of tumor incidence including liver tumor incidence and multiplicity tumor incidence are outliers and need to be further investigated in the future.

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Table4

Protein expression in liver tumor and normal liver of B6C3F1mice treated with carcinogenic PAHs

ramway	Protein	Tumor (T)	Normal (N)	T/N	${\it P}$ value
	АТМ	0.590 ± 0.046	0.761 ± 0.126	0.967	0.0432
	pATR	0.569 ± 0.037	0.748 ± 0.019	0.761	0.0001
	Chk1	0.788±0.028	0.961 ± 0.042	0.820	0.0005
	Chk2	1.640±0.128	1.493±0.115	1.098	0.1386
	pChK1	0.743±0.033	0.894 ± 0.017	0.831	0.0002
	DNA PKcs	0.564 ± 0.035	0.772 ± 0.015	0.731	<0.0001
DNA	ERCC1	4.413±0.347	3.450 ± 0.554	1.279	0.0257
repair	XRCC1	0.508 ± 0.032	0.640 ± 0.011	0.794	0.0002
	FANCD2	1.543±0.093	1.896±0.192	0.814	0.0162
	Ku80	0.585 ± 0.050	0.696±0.027	0.841	0.0081
	MSH2	2.510±0.218	2.274±0.313	1.104	0.2629
	Mre11	0.870 ± 0.040	1.194 ± 0.160	0.729	0.0077
	RAD50	1.465±0.152	1.362±0.166	1.076	0.3985
	Brca2	0.513 ± 0.036	0.680 ± 0.025	0.754	0.0003
	PARP1 (Cleaved)	0.905 ± 0.108	1.105 ± 0.072	0.819	0.0217
	Caspase3	0.363 ± 0.012	0.466 ± 0.015	0.779	<0.0001
	Caspase7 (Cleaved)	0.480±0.024	0.584±0.017	0.822	0.0004
Apoptosis	Bcl-xL	1.501 ± 0.143	1.075 ± 0.056	1.396	0.0014
	Bcl-2	1.301 ± 0.074	1.234 ± 0.136	1.054	0.4207
	Bax	1.485 ± 0.180	0.910 ± 0.015	1.632	0.0007
	pBad(Ser112)	0.988 ± 0.074	0.778 ± 0.031	1.270	0.0019
	XIAP	1.224±0.141	2.495 ± 0.059	0.491	0.0006
	P16/INK4A	1.314 ± 0.117	1.108 ± 0.099	1.186	0.0363
Cell cycle	p53	0.693 ± 0.036	0.884 ± 0.071	0.784	0.0030
regulation	Phopho-Rb (Ser807/811)	0.756±0.043	0.992±0.061	0.762	0.0007

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Pathway	Protein	Tumor (T)	Normal (N)	N/L	P value
	Rb	1.244±0.087	1.244 \pm 0.087 1.057 \pm 0.099 1.177	1.177	0.0293
	Cyclin B1	0.726±0.033	0.726±0.033	0.691	<0.0001
	Cyclin D1	0.947±0.068	0.947±0.068 0.885±0.110 1.070	1.070	0.3744
	Nrf2	1.013 ± 0.046	1.013±0.046 1.142±0.030 0.887	0.887	0.0034
Others	Egfr	0.405 ± 0.034	0.405 ± 0.034 0.455 ± 0.013 0.890	0.890	0.0329
	Cox2	0.527±0.029	0.609±0.050	0.865	0.0291

Normal tissues were obtained from control group, n=4.

Tumor tissues were from the BaP group (3 samples) and RM group (1 sample).

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