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***In vitro* metabolism of benzo[a]pyrene and dibenzo[def,p]chrysene in rodent and human hepatic microsomes**

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Abstract

Polycyclic aromatic hydrocarbons (PAHs) are ubiquitous and often carcinogenic contaminants released into the environment during natural and anthropogenic combustion processes. Benzo[a]pyrene (B[a]P) is the prototypical carcinogenic PAH, and dibenzo[def,p]chrysene (DBC) is a less prevalent, but highly potent transplacental carcinogenic PAH. Both are metabolically activated by isoforms of the cytochrome P450 enzyme superfamily to form reactive carcinogenic and cytotoxic metabolites. Metabolism of B[a]P and DBC was studied in hepatic microsomes of male Sprague-Dawley rats, naïve and pregnant female B6129SF1/J mice, and female humans, corresponding to available pharmacokinetic data. Michaelis–Menten saturation kinetic parameters including maximum rates of metabolism (V_{MAX} , nmol/min/mg microsomal protein), affinity constants (K_M , μM), and rates of intrinsic clearance (CL_{INT} , ml/min/kg body weight) were calculated from substrate depletion data. CL_{INT} was also estimated from substrate depletion data using the alternative *in vitro* half-life method. V_{MAX} and CL_{INT} were higher for B[a]P than DBC, regardless of species. Clearance for both B[a]P and DBC was highest in naïve female mice and lowest in female humans. Clearance rates of B[a]P and DBC in male rat were more similar to female human than to female mice. Clearance of DBC in liver microsomes from pregnant mice was reduced compared to naïve mice, consistent with reduced active P450 protein levels and elevated tissue concentrations and residence times for DBC observed in previous *in vivo* pharmacokinetic studies. These findings suggest that rats are a more appropriate model organism for human PAH metabolism, and that pregnancy's effects on metabolism should be further explored.

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Conflict of interest

The authors declare that there are no conflicts of interest.

Transparency document

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Keywords

Benzo[a]pyrene; Dibenzo[def,p]chrysene; Michaelis–Menten; V_{MAX} ; K_M ; Intrinsic clearance

1. Introduction

Polycyclic aromatic hydrocarbons (PAHs) are xenobiotics observed ubiquitously as complex mixtures throughout the environment. The release of PAHs into the environment results from natural and anthropogenic combustion of organic materials, including forest fires, fossil fuel powered transport, industrial manufacturing, smoking cigarettes and grilling meat (USEPA, 1988). PAHs are highly hydrophobic and non-volatile (Cerniglia, 1992) due to their structure of two or more fused benzene rings (Sims and Overcash, 1983), and are therefore persistent pollutants in the atmosphere, soil and water. As the number of fused benzene rings in the PAH increases, water solubility and volatility decrease (Wilson and Jones, 1993).

Many high molecular weight PAHs have been shown to be carcinogenic in humans and animals, including the prototypic carcinogenic PAH, benzo[a]pyrene (B[a]P), and the lesser studied but highly potent dibenzo[def,p]chrysene (DBC), also known as dibenzo[a,l]pyrene. Due to their adverse effects on human health, persistence in the environment, and tendency to be transformed into more reactive and dangerous metabolites, B[a]P and DBC have been classified by the International Agency for Research on Cancer (IARC) as Group 1, carcinogenic to humans, and Group 2A, probably carcinogenic to humans, respectively (WHO, 2010, 2012).

Major routes of PAH exposure occur via ingestion of contaminated food and water or inhalation of particulate matter in the ambient air (Miller and Ramos, 2001). Though natural forms of combustion such as forest fires, volcanic eruptions, and lightning contribute to human PAH exposure, anthropogenic sources including industrial manufacturing, petrol powered transportation, tobacco and cooking processes are principally responsible for the majority of human exposure (Guo et al., 2011). Once released into the atmosphere, PAHs immediately bind to particulate matter, which may then be inhaled or ingested to reach the metabolically active lungs and liver, and the rest of the body.

An alternate, lesser investigated, method of exposure is the exchange between mother and fetus during gestation. The placenta is a vital gestational organ that permits the exchange of blood, nutrients, and oxygen between mother and fetus (Miller and Ramos, 2001). It is also capable of transferring maternal PAHs, resulting in significant decreases in infant birth weight, length, and head and chest circumference (Hunter et al., 2010). Neonates may also be exposed during lactation, as breast milk consists of approximately 7–10% fat (Mandel et al., 2005), providing a hydrophobic environment in which PAHs may stably exist. In mice, we have shown DBC to be a potent transplacental carcinogen affecting multiple organs (Castro et al., 2008; Shorey et al., 2012; Yu et al., 2006).

The liver is an important location of metabolic activation due to its production of many xenobiotic metabolizing enzymes involved in PAH bioactivation and detoxification (Miller

and Ramos, 2001). PAHs are procarcinogenic molecules, requiring enzymatic processing to form reactive metabolites capable of exerting carcinogenic effects. The cytochrome P450 superfamily is largely responsible for PAH metabolism and is highly inducible in the liver (Nebert et al., 2004).

The principal CYP 450 isoforms responsible for activation of most carcinogenic PAHs are CYP 1A1 and 1B1 (Shimada and Fujii-Kuriyama, 2004), which work in conjunction with epoxide hydrolases, peroxidases and other enzymes to transform the B[a]P and DBC parent compounds to a variety of intermediates including epoxides, phenols, quinones and diol epoxides (Fig. 1). Examination of these metabolites found the epoxides and diol epoxides to be extremely reactive with DNA (Conney, 1982), readily forming adducts that resulted in mutation in growth control genes such as p53 (Pfeifer et al., 2002).

The purpose of this study was to determine the *in vitro* Michaelis–Menten kinetic parameters involved in the initial step in the metabolism of B[a]P and DBC across multiple species using liver microsomes obtained from male Sprague-Dawley rats, naïve and pregnant female B6129SFI/J mice, and female human. These species were selected due to the volume of previous literature on B[a]P in the rat, our ongoing research into the mode of action and pregnancy-dependent pharmacokinetics of the transplacental carcinogenicity of DBC in the mouse, and the ultimate goal of predicting the disposition of PAHs in humans (Castro et al., 2008; Crowell et al., 2011; Shorey et al., 2012; Yu et al., 2006). Kinetic parameters obtained from these studies are critical for revising preliminary physiologically based pharmacokinetic (PBPK) models for each PAH (Crowell et al., 2011) to facilitate comparisons between target tissue doses from high-dose toxicity studies with pregnant and non-pregnant animals to humans under realistic exposure conditions.

2. Materials and methods

2.1. Reagents and chemicals

DBC, DBC-11,12-diol, DBC-11,12,13,14-teraoles, B[a]P-7,8-diol, and B[a]P-7,8,9,10-tetraols were synthesized according to previously reported methods (Krzeminski et al., 1994; Luch et al., 1998; Sharma et al., 2004). B[a]P, sodium sulfate, sulfuric acid, acetone, methanol, ethyl acetate, phosphate buffered saline, magnesium chloride, tris acetate, potassium chloride, potassium pyrophosphate, ethylenediaminetetraacetic acid (EDTA) and nicotinamide adenine dinucleotide phosphate (NADPH) were purchased from Sigma (St. Louis, MO). All solvents were of high-performance liquid chromatography (HPLC) grade.

Highly concentrated stocks of B[a]P and DBC were diluted with acetone to create working solutions with concentrations ranging from 0.01 to 2.1 mM (B[a]P) and 0.004–0.9 mM (DBC) for use as analytical and internal standards, as well as substrates in metabolism assays. Solutions were stored at –80 °C in Teflon capped amber vials wrapped with parafilm to prevent degradation or evaporation. Theoretical stock concentrations were confirmed by HPLC with fluorescence detection (FLD).

2.2. Animals

Female B6129SF1/J mice, male 129S/SvImj mice, and male Sprague-Dawley rats (Jackson Laboratory, Bar Harbor, ME, USA) were housed separately in suspended plastic cages with chipped bedding, in rooms maintained at 21 ± 2 °C and $50 \pm 10\%$ relative humidity with a 12-h light/dark cycle. Animals were given a minimum acclimation period of seven days before breeding or harvesting tissues. Lab diet certified rodent chow and water were provided *ad libitum*. For metabolism studies with liver microsomes from pregnant animals, naïve female B6129SF1/J mice were bred with 129S/SvImj males after the one-week acclimation period. The appearance of a vaginal plug was designated gestational day (GD) 0, as in previous studies (Castro et al., 2008; Yu et al., 2006). Livers from pregnant mice were isolated on GD17 corresponding to the exposure day used in prior pharmacokinetic and carcinogenicity studies with DBC (Castro et al., 2008; Crowell et al., 2011; Shorey et al., 2012; Yu et al., 2006). The animal facility is accredited by the American Association for Accreditation of Laboratory Animal Care (AAALAC). All animal protocols were approved by the Institutional Animal Care and Use Committee at Pacific Northwest National Laboratory and studies were performed in accordance with the National Institutes of Health guidelines for the care and use of laboratory animals (NIH, 2011).

2.3. Microsome preparation

Pooled female human liver microsomes generated from 25 human females, 31–86 years of age, were obtained from Celsis *In Vitro* Technologies (Baltimore, MD, USA). Rodent hepatic microsomes were prepared according to the methods of Guengerich (Guengerich, 1994). Briefly, naïve and GD 17 female B6129SF1/J mice and male Sprague-Dawley rats were euthanized by CO₂ asphyxiation, exsanguinated, and liver tissue excised. Tissue was immediately placed in ice-cold 0.1 M PBS solution (pH 7.4). Livers were subsequently weighed and placed in 2 mL per gram tissue of homogenization buffer (1 M tris acetate, pH 7.4; 1 M KCl; 100 mM EDTA; d-H₂O), and homogenized by 6–8 passes of a drill press. Homogenate was centrifuged in a Beckman ultracentrifuge at 4 °C, $10,000 \times g$, for 30 min. Supernatant (S9 fraction) was decanted, and centrifuged for an additional 60 min at 4 °C and $105,000 \times g$. Supernatant (cytosolic fraction) was decanted and stored at –80 °C. Wash buffer (100 mM potassium pyrophosphate; 100 mM EDTA; d-H₂O) was added in excess to the ultracentrifuge tube to dislodge the pellets (microsomal fraction), which were then pooled and centrifuged a final time for 45 min at 4 °C and $105,000 \times g$. Supernatant was discarded and the pellets resuspended with 0.5 mL per gram original tissue of resuspension buffer (1 M tris acetate; 100 mM EDTA; d-H₂O). Resuspended pellets were homogenized with 5–6 passes of the drill press. Microsomes were aliquoted by 0.5 mL volumes and stored at –80 °C. Microsomal protein (MSP) concentration was determined by bicinchoninic acid spectrophotometric assay (BCA), using bovine serum albumin as a standard of known concentration; CO difference spectra assays were performed to determine the microsomal CYP concentration.

2.4. In vitro metabolism studies

Reactions were conducted in 500 µL total volumes, consisting of 0.1 M phosphate buffer (pH 7.4), MSP (0.2–1.0 mg/ml), MgCl₂ (3 mM), and excess NADPH (1.5 mM). Reactions

were initiated by the addition of the chemical of interest (DBC or B[a]P) dissolved in an acetone vehicle (total organic in reaction less than 1%, v/v) to a final reaction concentration of 0.05–35 μM (B[a]P) or 0.05–9.0 μM (DBC) and were incubated at 37 °C for 0–30 min. Control experiments were performed to verify that reaction kinetics were linear with regard to time and MSP content, and without key substrates to establish baseline chromatography (without PAH) and substrate stability over time (without NADPH). Reactions were quenched with the addition of 500 μL 0.9 M H_2SO_4 , followed by the addition of a 5 μL internal standard (B[a]P for DBC kinetic experiments; DBC for B[a]P kinetic experiments). Samples were then vortexed and placed on ice until extraction.

Liquid–liquid extractions were performed as follows: 250 mg Na_2SO_4 was added to each sample, followed by 500 μL ethyl acetate. Samples were vortexed, then centrifuged for 10 min at 4 °C and $1600 \times g$. The supernatant was drawn off using a glass Pasteur pipette to a new tube, and then the extraction repeated. The combined supernatants were blown to dryness under a gentle stream of nitrogen. Samples were reconstituted in 500 μL methanol and analyzed as described below.

2.5. Sample analysis

DBC, B[a]P, and their hydroxylated metabolites were quantified by reverse-phase HPLC using an Agilent 1100 HPLC system equipped with a fluorescence detector (Santa Clara, CA, USA). Twenty μL of reconstituted sample were injected onto an Ascentis 25 cm \times 4.6 mm, 5 μm C_{18} column (Sigma–Aldrich, St. Louis, MO, USA). A water:acetonitrile gradient from 45:55 to 0:100 was employed from 0 to 10 min, and then held at 100% acetonitrile until 22 min, at a constant flow rate of 0.95 mL/min. Excitation and emission wavelengths were 245 and 430 for tetraol metabolites, 360 and 430 for diol metabolites, and 235 and 430 for B[a]P and DBC. Elution times were 20.9 and 18.1 min for DBC and B[a]P, 10.7 and 8.5 min for DBC-diol and B[a]P-diol, 5.4, 6.0, and 6.6 min for DBC tetraols, and 3.3, 3.9, and 4.6 min for B[a]P tetraols. The limits of reliable quantitation (LOQ) for B[a]P and DBC were 0.004 and 0.02 μM , respectively.

2.6. Data analysis

Initial enzyme velocity (V_s , nmol/mL/min) was obtained by linear regression of parent PAH concentration (μM) versus time (min) data from individual kinetic experiments; a representative time course appears in Fig. 2. Regressions included at least six data points and were only considered valid if R^2 was equal to or greater than 0.7. To determine kinetic parameters, nonlinear regressions of protein-normalized initial velocities versus initial reaction substrate concentrations ($[\text{S}]$, μM) were performed using R: A Language and Environment for Statistical Computing Version 2.13.1 (R Foundation for Statistical Computing, Vienna, Austria) according to the Michaelis–Menten model:

$$V_s = \frac{V_{MAX} [\text{S}]}{K_M + [\text{S}]} \quad (1)$$

where V_{MAX} (nmol/min/mg MSP) is the maximum enzyme velocity and K_M (μ M) is the Michaelis constant, defined as the initial reaction concentration required to reach half the V_{MAX} .

From V_{MAX} and K_M , intrinsic clearance rates (CL_{INT}) were predicted by applying the methods of Obach et al. (1997). CL_{INT} was scaled up from microsomal (mL/min/mg MSP) to tissue (mL/min/g liver) to organismal (mL/min/kg body weight) levels utilizing liver weights of 55, 41.8, 36.6 and 25.7 g per kg body weight for naïve and pregnant (GD 17) mice, rats and humans, respectively (Brown et al., 1997; Oflaherty et al., 1992), as described in Eq. (2).

$$CL_{INT} = \frac{V_{MAX}}{K_M} * \frac{mg \text{ MSP}}{1 \text{ g liver weight}} * \frac{g \text{ liver}}{1 \text{ kg body weight}} \quad (2)$$

Though the organism level CL_{INT} is more useful in development of physiologically based pharmacokinetic (PBPK) models, tissue level CL_{INT} involves fewer extrapolations from the microsomal level at which experiments were conducted, and was therefore also included for comparison. Alternatively, intrinsic clearances were also determined directly using the *in vitro* half-life method (Obach, 1999) for comparison to values determined from classic Michaelis–Menten regressions of substrate depletion data. Scaled *in vitro* metabolic parameters such as these can be used in physiologically based pharmacokinetic models alongside parameters describing physiology and biochemistry governing chemical disposition within an organism, as described previously (Crowell et al., 2013).

3. Results

PAH metabolism was observed regardless of sex or species of organism. This was demonstrated by decreasing parent compound and increasing metabolite peak area count over time (Fig. 3). Michaelis–Menten regressions appear alongside measured data points in Fig. 4.

In all species, V_{MAX} was higher for B[a]P than DBC, by a margin of 2.8 fold in humans, 6.5 fold in rats, and 6.7 fold in mice (Table 1 Table 1). Estimates of CL_{INT} at both the tissue and organism level, regardless of method of calculation, were also universally higher for B[a]P than DBC. For those calculated from Michaelis–Menten regressions, CL_{INT} for B[a]P exceeded that of DBC by a margin of 2.6 fold in male rats, 9.0 fold in female mice, and 3.3 fold in female humans; for those calculated by the *in vitro* half-life method, B[a]P CL_{INT} exceeded DBC CL_{INT} by a margin of 1.4 fold in rats, 10.6 fold in mice, and 13.3 fold in humans (Table 1). Michaelis constants did not have a similar chemical specific trend: K_M s for DBC were marginally higher than for B[a]P in the female mouse and female human, while the reverse was true for male rat (Table 1).

Maximum velocities of PAH depletion were lowest in female humans and highest in female mice, regardless of parent compound (Table 1, Fig. 5). For DBC, V_{MAX} ranged from 0.009 ± 0.005 nmol/min/mg MSP in the female human to 0.646 ± 0.055 nmol/min/mg MSP in the naïve female mouse; in B[a]P from 0.025 ± 0.007 nmol/min/mg MSP in the female human

and 4.35 ± 0.628 nmol/min/mg MSP in the naïve female mouse. For both compounds, V_{MAX} in rodents far exceeded that in humans (Table 1).

Michaelis constants for DBC metabolism were generally comparable between rodents, ranging from 6.63 μ M in female mice to 7.28 μ M in male rats, while the K_M for DBC metabolism in female humans was markedly lower (0.92 μ M). For B[a]P, K_M in humans was also the lowest by a large margin, at 0.73 μ M, but K_M in male rats exceeded that in female mice by 3.6 fold (Table 1, Fig. 5).

The effect of pregnancy on DBC metabolism kinetics was evaluated in female mice. Both V_{MAX} and K_M were reduced in the pregnant animal, by 35% and 51% respectively. Calculated organismal level CL_{INTS} were comparable between pregnant and naïve mice, despite a 30% higher tissue level CL_{INT} in the pregnant animal using the Michaelis–Menten regression method (Table 1). For CL_{INT} estimated *via* the *in vitro* half-life method, the tissue level CL_{INT} is 28% lower in the pregnant animal, and scaling up to organism level CL_{INT} results in a 45% lower clearance rate in the pregnant animal (Table 1).

Finally, estimates of CL_{INT} by the *in vitro* half-life method were generally within twofold of those calculated by the Michaelis–Menten method for both PAHs and all species. However, for both PAHs, tissue and organism level CL_{INTS} estimated by the Michaelis–Menten method were comparable for female humans and male rats, while female mice were markedly higher; this effect was greater for B[a]P than for DBC. A similar trend was apparent for CL_{INT} calculated by the *in vitro* half-life method for B[a]P metabolism, while CL_{INT} calculated using this method for DBC metabolism were dissimilar in all organisms (Table 1, Fig. 6).

4. Discussion

Elevated maximum enzyme velocities (V_{MAX}) and CL_{INTS} were observed for B[a]P compared to DBC (Table 1). Composed of six fused benzene rings, DBC is larger and bulkier than five ringed B[a]P. This additional benzene ring creates steric hindrance during biotransformation (Nelson and Cox, 2008), possibly explaining the slower rates of conversion of DBC substrate to metabolite product.

The fourfold higher K_M observed for B[a]P in the male rat relative to naïve mouse indicates that the CYP system of the mouse has a higher affinity for B[a]P relative to the rat, and when coupled with the higher observed mouse V_{MAX} , may explain the two order of magnitude disparity of B[a]P organism level CL_{INT} between mouse and rat (Fig. 6). The high error of the estimate of K_M here (as well as for other kinetic parameters) may also stem from the hydrophobic nature of B[a]P and DBC in an essentially aqueous experimental system.

While clearance values obtained from the *in vitro* half-life method were generally similar to those calculated from Michaelis–Menten analyses, there are limitations to this method that should be considered. Estimates of clearance are made using much less data than for Michaelis–Menten regressions, and therefore may not accurately reflect clearance values or their variability. The *in vitro* half-life method also required that reaction concentrations fall

well below K_M , information which may not be well established for a given compound. Relatedly, the clearance values cannot be used to determine V_{MAX} and K_M , and therefore are only reflective of clearance in the first-order range of substrate concentrations (i.e., clearance of low exposures). The *in vitro* half-life method is certainly useful as an initial, rapid, and low cost method of investigation of metabolism kinetics. Additionally, in situations where substrate solubility may hinder accurate Michaelis–Menten analyses, as may be the case for B[a]P in male rat, the availability of an alternate method is invaluable. In the pregnant mouse, V_{MAX} and K_M of DBC depletion were reduced compared to the non-pregnant mouse (Table 1, Fig. 5). It has been shown that pregnant rats in mid to late gestation (GD13 and GD19) have significantly decreased hepatic expression of CYP1A1 (He et al., 2005)—one of the primary isoforms involved in PAH metabolism (Shimada and Fujii-Kuriyama, 2004). While comprehensive information on enzyme expression and activity during pregnancy in the mouse is not available, we have recently used a novel functional proteomic approach, activity based protein profiling, to assess the relative abundances of active enzyme protein levels in pregnant and naïve mice, and found a 2–10 fold reduction in over 30 active P450 enzyme isoforms in the livers of pregnant mice (Crowell et al., 2013). These findings support the observed reduction in overall microsomal V_{MAX} levels but say little about PAH-specific K_M s. If the depletion of parent PAH were dependent on a single enzyme, we would not expect to see a change in enzyme affinity with a purported reduction in enzyme expression or abundance. However, because multiple CYP isoforms contribute to PAH metabolism, the observed reduction in overall apparent K_M may be attributable to an uneven reduction in several isoforms that contribute to DBC metabolism.

Regardless, overall microsomal apparent K_M s estimated from studies with higher molecular weight PAHs are subject to greater error than those determined for lower molecular weight PAHs. This is due to a corresponding decreased water solubility, even with closely related PAHs like B[a]P and DBC, that hinders the ability to conduct studies over a sufficiently broad concentration range to adequately characterize the shift between first-order and zero-order reaction rates. This dose-dependent shift in overall tissue kinetics is further complicated by potential changes in isoform ratios as a result of pregnancy. As a result, overall tissue and organismal level CL_{INT} values determined using the *in vitro* half-life method that does not depend upon accurate estimates of K_M are more likely relevant for comparing across species and pregnancy status as long as exposures are well below those that cause metabolic saturation.

When using rodent data to estimate human risk it is vital to understand interspecies variability in both expression and catalytic activity of relevant CYP isoforms. It was previously believed that CYP1A1 was predominantly responsible for PAH detoxification and activation (Conney, 1982). Studies showed that PAH exposure induced increased expression of CYP1A1; however, CYP1B1 has been shown to transform PAH parent compounds into reactive and carcinogenic metabolites at a higher rate than CYP1A1 (Murray et al., 2001). Both isoforms are predominantly expressed in organs other than the liver (Shimada, 2006) and are highly inducible by PAHs (Murray et al., 2001; Uno et al., 2004). Strong regulatory conservation of CYP1A isoforms has been observed between male

rat and humans (Mugford and Kedderis, 1998), corroborating our finding that human metabolism kinetics were more similar to male rat than to female mouse. Clearance values for B[a]P and DBC in female mice were generally far in excess of those in male rat or female human (Table 1, Fig. 6), supporting the previous reports that rats may provide a more accurate model for human risk assessment.

In order to establish more accurate human dosimetry, *in vitro* studies using human tissues must be continued. As the CYP iso-forms involved in the metabolism of B[a]P and DBC are highly inducible, future studies should focus on how pre-exposure to PAHs affects kinetic parameters. PAHs occur environmentally as complex mixtures of varying composition, making accurate risk assessment and exposure estimates in humans difficult to evaluate (Wenzl et al., 2006). To produce more accurate human models, studies should be performed applying PAHs as simple as well as environmentally relevant mixtures. Though the liver is considered an important target of PAH parent compounds due to its production of many of the enzymes involved in bioactivation and detoxification (Miller and Ramos, 2001), lung and other tissues are also metabolically active target sites and should be investigated. To better quantify rates of production and depletion of carcinogenic metabolites, studies of downstream metabolites as the initial substrate should be performed. Focus should be placed on the hydroxylated diol and tetraol metabolites as the carcinogenic epoxide metabolites are too unstable for *in vitro* study.

The findings reported here provide an important basis for further development of quantitative models of PAH disposition, such as those previously described (Crowell et al., 2011, 2013), and facilitate more robust extrapolations and estimations of human risk.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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HIGHLIGHTS

- PAH clearance was highest in naïve female mice and lowest in female humans.
- PAH clearance rates in male rat were more similar to female human than female mice.
- Metabolism of B[a]P was generally more rapid than DBC, regardless of organism.
- DBC metabolism kinetics were reduced in the pregnant mouse relative to naïve mice.

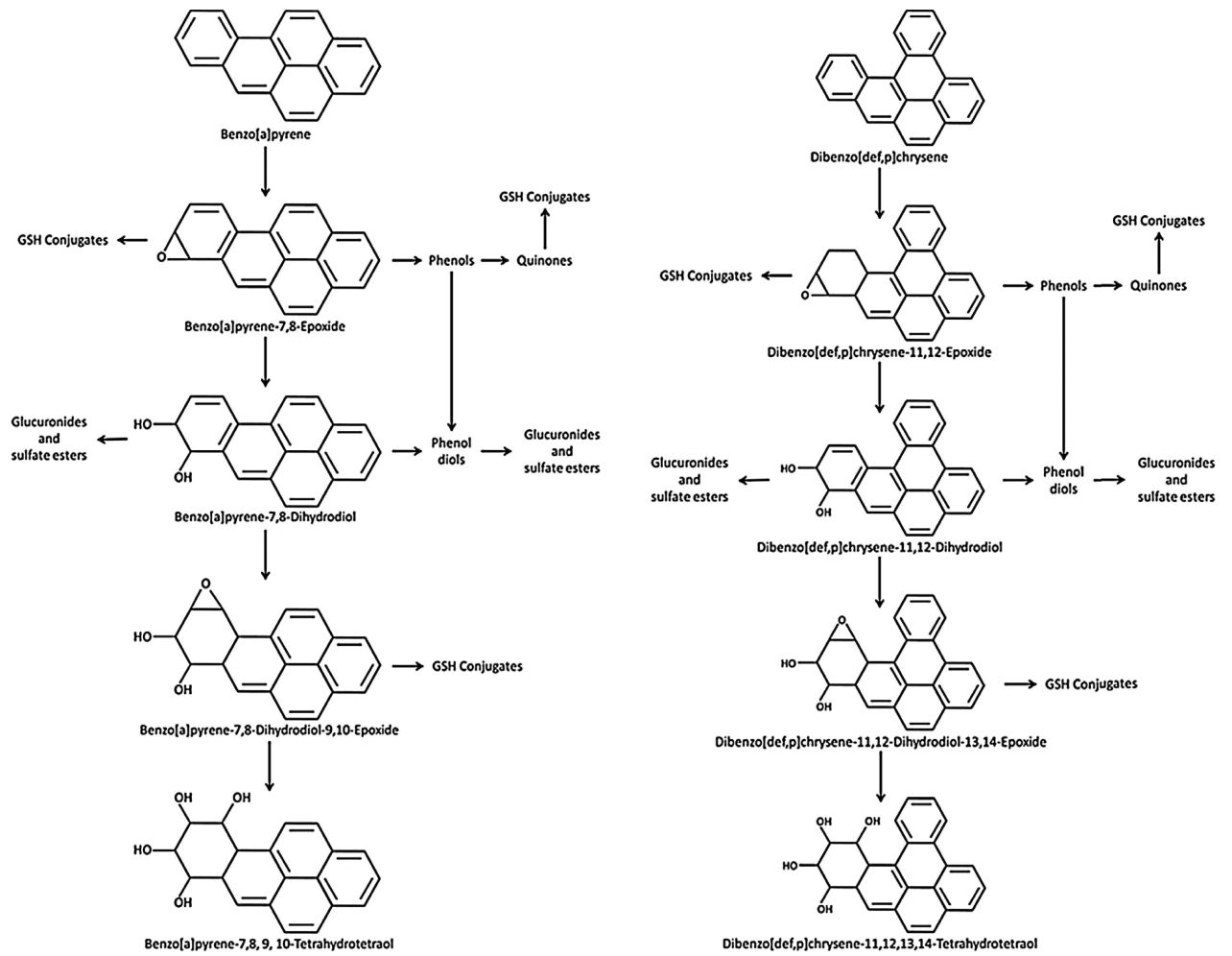


Fig. 1.
Metabolic pathway of benzo[a]pyrene (B[a]P) and dibenzo[def,p]chrysene (DBC).

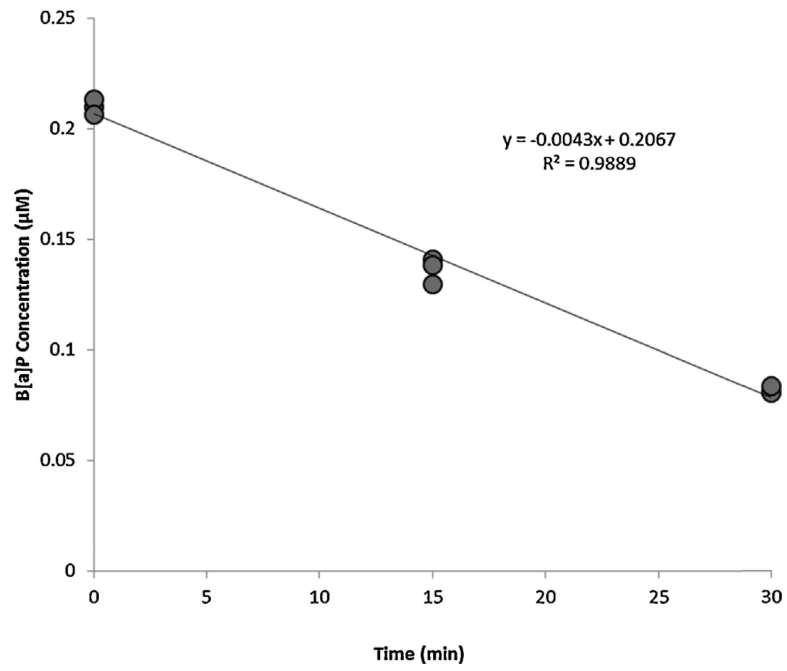


Fig. 2. Representative B[a]P substrate depletion over 30 min reaction period in the female human at an initial velocity (V_0) of 0.0043 nmol/mL/min.

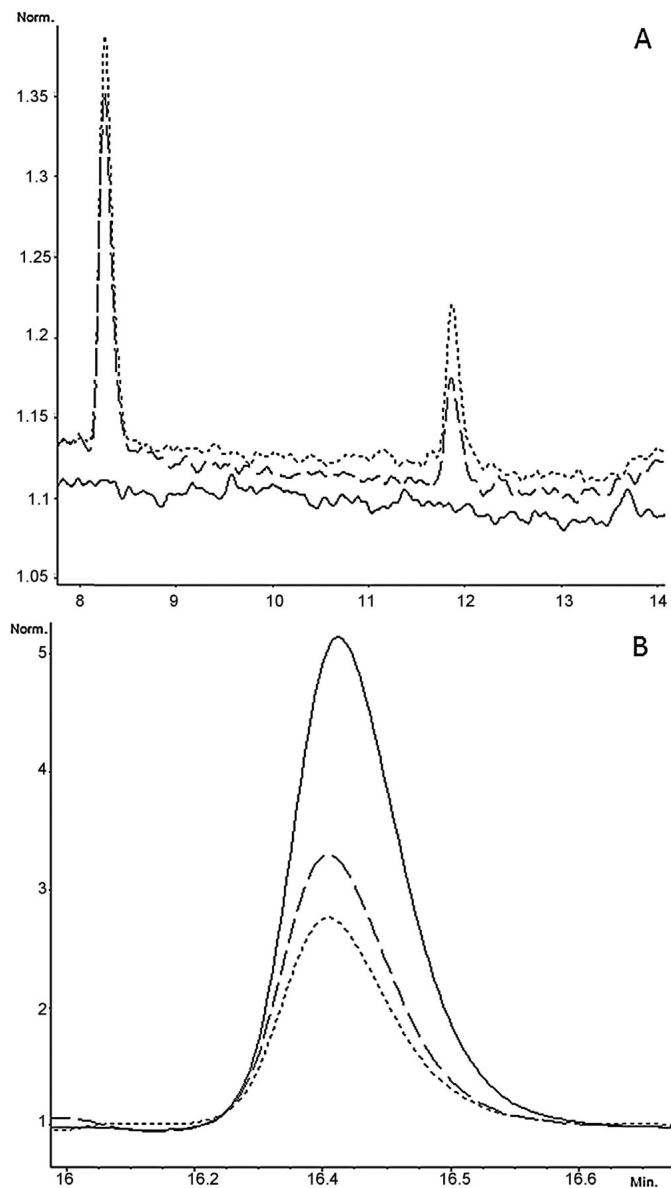


Fig. 3.

(A) Chromatograph overlay of B[a]P metabolites produced at times 0 min (----), 15 min (---), and 30 min (- - -). B[a]P tetraol (retention time 8.3 min) and diol (retention time 12 min) metabolite peak areas increase over reaction time as parent B[a]P compound is consumed. (B) Chromatograph overlay of B[a]P parent compound at times 0 min (----), 15 min (---), and 30 min (- - -). B[a]P concentration decreases over reaction time due to formation of metabolites.

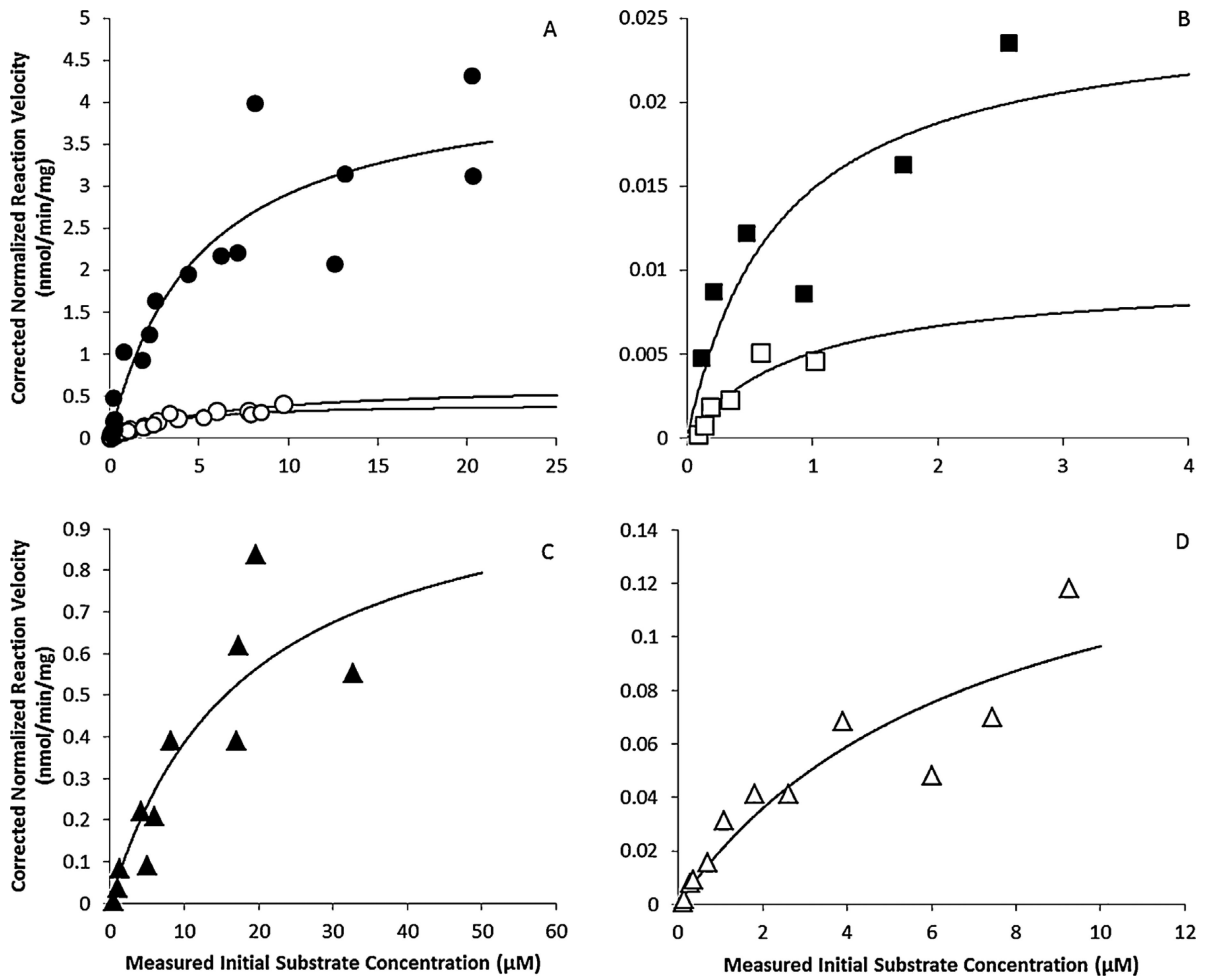


Fig. 4. Michaelis–Menten kinetic analyses of substrate depletion for (A) B[a]P in the naïve mouse (●) and DBC in naïve (○) and pregnant (●) mouse; (B) B[a]P (■) and DBC (□) in female human; (C) B[a]P (▲) in male rat and (D) DBC (△) in male rat.

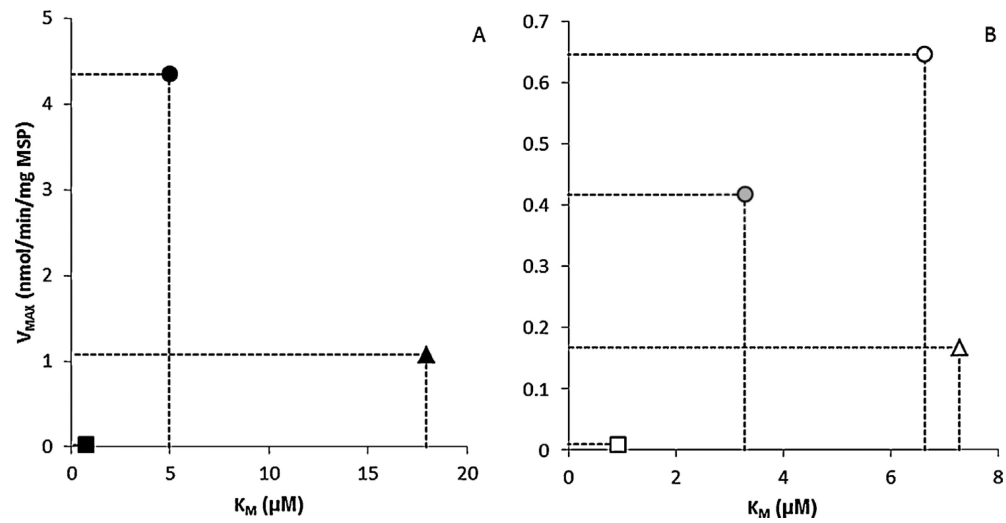


Fig. 5. Comparison of maximum velocities (V_{MAX}) and Michaelis–Menten constants (K_M) calculated from (A) B[a]P depletion data in naive female mouse (●), male SD rat (▲) and female human (■), and (B) DBC depletion data in naïve (○), pregnant female mouse (●), male SD rat (△) and female human (□).

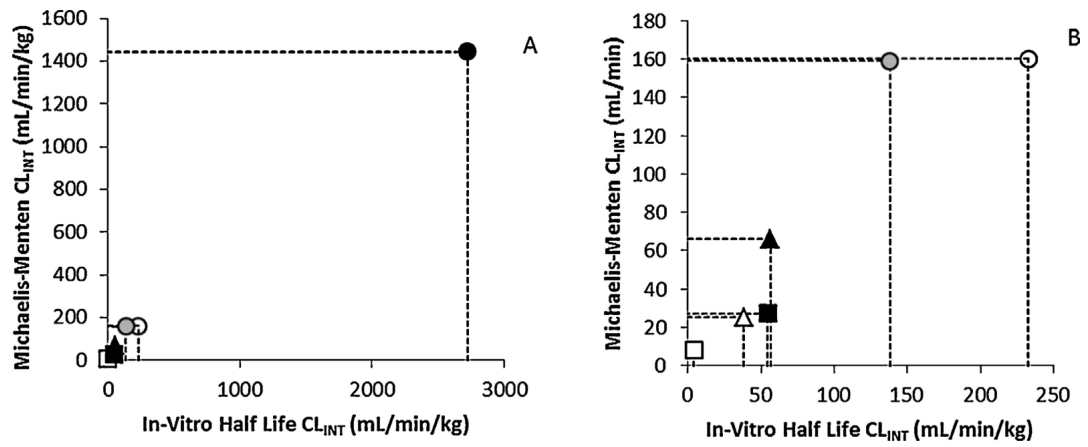


Fig. 6. Comparison of organismal CL_{INT} values obtained via Michaelis–Menten and *in vitro* half-life methods of B[a]P in naive female mouse (●), male SD rat (▲) and female human (■), and DBC in naïve (○), pregnant female mouse (●), male SD rat (△) and female human (□). (A) All organisms and (B) B[a]P in female mouse removed to show detail.

Table 1

Kinetic parameters of measured initial substrate concentration ranges ($[S]_0$, μM) of B[a]P and DBC metabolism in the naïve and pregnant B6129SF1/J mouse, male Sprague-Dawley rat and female human. Maximum enzyme velocity (V_{max}). Michaelis constant (K_m). Michaelis-Menten and *In vitro* half-life intrinsic clearance rates (CL_{INT}) scaled to the hepatic tissue and organismal level.

Chemical	Organism	Measured [S] range (μM)	V_{max} (nmol/min/mg MSP)	K_m (μM)	Tissue level CL_{INT} (ml/min/g tissue)		Organism level CL_{INT} (ml/min/kg body weight)	
					MM	$t_{1/2}$	MM	$t_{1/2}$
B[a]P	Naïve mouse	0.143-20.4	4.35 \pm 0.628	4.97 \pm 1.96	26.28 \pm 14.14	48.53 \pm 32.79	1445.2 \pm 778.0	2699.0 \pm 1803.4
	Male rat	0.161-90.9	1.08 \pm 0.401	17.97 \pm 12.87	1.8 \pm 1.96	1.40 \pm 0.61	65.90 \pm 71.68	51.34 \pm 22.19
	Female human	0.119-2.56	0.025 \pm 0.007	0.725 \pm 0.581	1.06 \pm 1.17	2.13	27.16 \pm 29.98	54.69
DBC	Naïve mouse	0.049-9.72	0.646 \pm 0.055	6.63 \pm 1.01	2.92 \pm 0.71	4.59 \pm 0.53	160.71 \pm 38.94	252.18 \pm 29.26
	Pregnant mouse	0.024-8.5	0.417 \pm 0.075	3.28 \pm 1.38	3.81 \pm 2.28	3.31 \pm 1.75	159.43 \pm 95.51	138.50 \pm 72.96
	Male rat	0.107-9.27	0.167 \pm 0.064	7.28 \pm 4.99	0.69 \pm 0.73	1.01 \pm 0.30	25.14 \pm 26.85	37.05 \pm 10.96
	Female human	0.087-1.02	0.009 \pm 0.005	0.915 \pm 0.823	0.32 \pm 0.46	0.16	8.19 \pm 11.73	4.23