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Fatty Liver and Impaired Hepatic Metabolism Alter the Congenerspecific Distribution of Polychlorinated Biphenyls (PCBs) in Mice with A Liver-specific Deletion of Cytochrome P450 Reductase

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Abstract

Polychlorinated biphenyls (PCBs) are persistent organic pollutants that are linked to adverse health outcomes. PCB tissue levels are determinants of PCB toxicity; however, it is unclear how factors, such as an altered metabolism and/or a fatty liver, affect PCB distribution in vivo. We determined the congener-specific disposition of PCBs in mice with a liver-specific deletion of cytochrome P450 reductase (KO), a model of fatty liver with impaired hepatic metabolism, and wild-type (WT) mice. Eight-week-old male WT (M_{WT} , n=3), male KO (M_{KO} , n=5), female WT (F_{WT} , n=4), and female KO mice (F_{KO} , n=4) were exposed orally to Aroclor 1254. PCBs were quantified in adipose, blood, brain, and liver tissues by gas chromatography-mass spectrometry. The ΣPCB levels followed the rank order adipose $>$ liver \sim brain $>$ blood in WT and adipose \sim liver $>$ brain $>$ blood in KO mice. PCB levels were much higher in the liver of KO than WT mice, irrespective of the sex. A comparison across exposure groups revealed minor genotype and sex-dependent differences in the PCB congener profiles (cos $\Theta > 0.92$). Within each exposure group, tissue profiles showed small differences between tissues (cos $\Theta = 0.85$ to 0.98). These differences were due to a decrease in metabolically more labile PCB congeners and an increase in congeners resistant to metabolism. The tissue-to-blood ratio of PCBs decreased for adipose, increased for the brain, and remained constant for the liver with an increase in chlorination. While these ratios did

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

not follow the trends predicted using a composition-based model, the agreement between experimental and calculated partition coefficients was reasonable. Although the distribution of PCBs differs between KO and WT mice, the magnitude of the partitioning of PCBs from the blood into tissues can be approximated using composition-based models.

Graphical Abstract

Capsule:

Mice with a liver-specific deletion of cytochrome P450 reductase display an altered congenerspecific distribution of PCBs due to a fatty liver and impaired hepatic metabolism.

Keywords

composition-based model; disposition; tissue to blood partition coefficients; persistent organic pollutants; transgenic mice

1. Introduction

Exposure to polychlorinated biphenyls (PCBs), a group of 209 persistent organic pollutants, is associated with adverse human health effects, including cancer, developmental neurotoxicity, and effects on the immune system (ATSDR, 2000). Individual PCB congeners can selectively affect a myriad of cellular targets in humans (Pencikova et al., 2018; Sethi et al., 2019); however, humans are exposed to complex mixtures of >150 PCB congeners via the diet, dermally or by inhalation (Ampleman et al., 2015; ATSDR, 2000; Chen et al., 2017; Schecter et al., 2010). Moreover, occupational exposures to PCBs remain a concern (Haga et al., 2018). In contrast, in vitro and in vivo studies of PCBs typically employ individual congeners or simplified and technical PCB mixtures. The disposition of individual PCBs in laboratory animals, especially after oral exposure, is well investigated. Several studies also report the tissue distribution of PCBs after inhalation exposure in rats (Hu et al., 2010; Hu et al., 2012; Hu et al., 2013). In mammals and other species, many PCB congeners are oxidized by hepatic and extrahepatic cytochrome P450 enzymes to hydroxylated metabolites (OH-PCBs) (Grimm et al., 2015). These metabolites can undergo further biotransformation to, for example, dihydroxylated metabolites, sulfate and glucuronide conjugates, and other PCB metabolites and, depending on the degree of chlorination are excreted with the feces or urine (Birnbaum, 1985).

Toxicokinetic models have been developed for individual PCB congeners or simple PCB mixture in rodent models following oral (Kania-Korwel et al., 2012; Kania-Korwel et al., 2010) and intravenous exposure (Birnbaum, 1983; Lutz et al., 1984). These studies provide insights into the levels of PCBs in target tissues, such as the adipose tissue, blood, liver, or brain, thus allowing the translation of results from toxicity studies with PCBs to humans. As a rule of thumb, PCB levels are directly proportional to the fat content of a tissue, and, in wild-type rodents, follow the general order adipose > liver > brain > blood (Kania-Korwel et al., 2005; Milanowski et al., 2010; Weisbrod et al., 2000). However, the partitioning of PCBs from the blood into tissues can differ from this general trend, depending on the animal model, the experimental design, or other factors.

A number of computational models have been developed to predict the partition coefficients in tissues in the absence of experimental data, including tissue composition-based models (Endo et al., 2013; Schmitt, 2008). These models use readily available information, including the quantities of the biological components (e.g., lipids, protein, and water) in tissues and the partition coefficients of PCBs (e.g., lipid/water partitioning coefficient), to estimate the distribution of PCBs in vivo. For example, a PP-LFER-composition-based model was adopted to determine the liver : blood partitioning coefficients of mono- to hexachlorinated PCBs and the adipose : plasma partition coefficients of di- to octachlorinated PCBs in rats (Endo et al., 2013). This model can be used to predict the partitioning of PCBs between animal tissues of interest for all PCB congeners with published Abraham solvation parameters (van Noort et al., 2010) and established PP-LFERs describing the correlations between the chemical descriptors and tissue/water partition coefficients (Endo et al., 2012; Endo et al., 2011; Endo and Goss, 2011; Geisler et al., 2012). Moreover, this model can estimate the tissue : blood partitioning of PCBs in animal models that the quantities of the biological components are altered due to genetic or environmental factors.

Factors affecting the tissue : blood partitioning of PCBs include an altered (hepatic) metabolism and redistribution away from the site of metabolism to tissues with a high-fat content (Birnbaum, 1983; Soontornchat et al., 1994). The induction of hepatic cytochrome P450 enzymes will more rapidly eliminate episodic and some persistent PCB congeners (Imsilp and Hansen, 2005). In sub-chronic studies, growth dilution can contribute to the distribution of PCBs (Hansen and Welborn, 1977). Also, changes in the composition of body compartments can shift the distribution of PCBs. Starvation shifts the partitioning of PCB 153 from adipose to other tissues (Wyss et al., 1982). Similarly, PCB 153 is mobilized from fat tissues during lactation and redistributed to the breast milk (Spindler-Vomachka and Vodicnik, 1984). Fatty liver caused by exposure to PCB 126 produces a redistribution of PCBs from the adipose tissue to the liver (Chu et al., 1994; Van Birgelen et al., 1994). Similarly, fatty liver induced by a high-fat diet is expected to result in the sequestration of PCBs in the mouse liver, a process that may underly the obesogenic effects of PCBs (Wahlang et al., 2013; Wahlang et al., 2014). Moreover, transgenic animals can have body weights or tissue compositions that differ from wild-type animals and affect PCB disposition (Milanowski et al., 2010; Wu et al., 2015; Wu et al., 2020).

The objective of the present study was to determine how a fatty liver and an impaired hepatic PCB metabolism alters the PCB congener profiles in target tissues following exposure to a complex PCB mixture. PCB congener profiles, levels, and tissue : blood ratios were determined in adipose tissue, blood, brain, and liver in male and female KO and congenic WT mice exposed to a complex PCB mixture. The KO mouse model displays an impaired hepatic xenobiotic metabolism and a fatty liver with altered fatty acid profiles (Gu et al., 2005; Weng et al., 2005). Aroclor 1254 was selected for this study because the disposition and toxicity of this technical PCB mixture are well characterized (ATSDR, 2000). Moreover, Aroclor 1254 contains PCB 91 and PCB 136 (Frame, 1997). The disposition and toxicokinetics of both PCB congeners have been studied in KO and congenic WT mice after a single oral exposure to the individual PCB congeners (Li et al., 2019; Wu et al., 2015; Wu et al., 2020). We adopted a comparable study design to facilitate comparison across studies.

2. Materials and Methods

2.1. Chemicals and Reagents

Aroclor 1254 (lot #: KB 05–612) was provided by Dr. Larry Hansen. A detailed characterization of this Aroclor 1254 batch has been reported previously (Zhao et al., 2010). Additional information regarding other chemicals and reagents is provided in the supplementary material.

2.2. Animal exposures

Alb-Cre+/−/Cprlox+/+ mice with a liver-specific deletion of the cytochrome P450 oxidoreductase (EC 1.6.2.4) gene (KO) and congenic Alb-Cre^{-/-}/Cpr^{lox+/+} (WT) mice were obtained from Dr. Xinxin Ding (University of Arizona, AZ, USA) to establish a breeding colony at the University of Iowa (Gu et al., 2003; Wu et al., 2003). Details regarding the maintenance of the breeding colony are provided in the supplementary material. The Institutional Animal Care and Use Committee of the University of Iowa approved all animal procedures (protocol #: 1206120). Daily animal welfare-assessments were performed by laboratory personnel, and no adverse outcomes were observed throughout the study. Eightweek-old mice were randomly divided into four exposure groups, including male WT $(M_{\rm WT})$ n=3), male KO (M_{KO} , n=5), female WT (F_{WT} , n=4), and female KO (F_{KO} , n=4). The animals were exposed around 9 am once by oral gavage to Aroclor 1254 (16 mg/kg body weight in corn oil, 10 mL/kg body weight), respectively. This acute dosing paradigm lays the groundwork for future toxicokinetic studies. The dose was selected based on a previous animal study and ensured a robust detection of PCBs in target tissues (Kania-Korwel et al., 2005). In parallel, animals $(n=3)$ were exposed to corn oil $(10 \text{ mL/kg}$ body weight) alone to assess the background contamination of the animals. The animals were euthanized 24 h after PCB exposure by carbon dioxide asphyxiation followed by thoracotomy and the removal of vital organs as confirmation of death. Whole blood was collected by cardiac puncture and stored at − 20 °C in glass tubes with 80 μL of ethylenediaminetetraacetic acid solution (EDTA, 7.5% w/w). Tissues (abdominal adipose tissue, brain, and liver) were collected and stored at −80 °C.

2.3. PCB extraction

PCBs were extracted from adipose $(0.05-0.09 \text{ g})$, brain $(0.10-0.15)$, liver $(0.06-0.18 \text{ g})$ and blood (0.55–0.73 g) with a pressurized liquid extraction method (Kania-Korwel et al., 2007; Wu et al., 2015) on the Dionex ASE200 system (Dionex, Sunnyvale, CA, USA). Extraction cells (33 mL) loaded with pre-combusted Florisil (12 g) and diatomaceous earth (DE, 2 g) were extracted with a mixture of hexane and acetone (1:1, v/v) on an ASE200 at 100 °C and 1500 psi (10 MPa) with preheat equilibration for 6 min, 60% of cell flush volume, and 1 static cycle of 5 min. The tissue was homogenized with the top DE layer, and the DE-tissue mixture was placed back into the extraction cell. Blood samples were loaded directly onto the cellulose filter of the cell. Surrogate standards, d_5 -PCB 30 and d_5 -PCB 65 (80 ng each), were added to every sample before the extraction. The cells were extracted with the same parameters described above. The extracts were concentrated to ~0.5 mL on a TurboVap II (Biotage LLC, NC, USA), transferred to a pre-combusted tube, and treated with 2 mL of concentrated sulfuric acid for lipid removal. The extract was transferred to a new glass tube, evaporated under a gentle stream of nitrogen to ~50 μL, and transferred to a gas chromatography vial. The internal standards (volume correctors), ${}^{13}C_{12}$ -PCB 9 and ${}^{13}C_{12}$ -PCB 194 (50 ng each), were spiked to every sample before analysis.

2.4. Gas chromatographic determinations

Samples were analyzed using an Agilent 7890A gas chromatograph coupled with an Agilent 5975C Inert Mass Selective Detector (MSD) operated in electronic ionization mode. An SLB-5MS capillary GC column (30 m length, 250 μm inner diameter, 0.25 μm film thickness; Supelco, Bellefonte, PA, USA) was used in the select ion monitoring (SIM) mode (Holland et al., 2016; Hu et al., 2015; Li et al., 2018). The temperature program was as follows: 80 °C, hold for 1 min, 2 °C/min to 160 °C/min, 1 °C/min to 170 °C, hold for 15 min, 1 °C/min to 180 °C, hold for 15 min, 1 °C/min to 245 °C, t10 °C/min to 300 °C, and hold for 15 min. The injector temperature was 280 °C, and the operating temperatures were 280 °C, 230 °C and 150 °C for transfer line, source, and quadrupole, respectively. The flow rate of carrier gas helium was 1.1 mL/min. By this congener specific method, 162 individual or coeluting peaks of PCBs can be separated and quantified. Twenty-eight channels were set up in GC-MS SIM mode (Table S1). The MSD was linear over the concentration range encountered in this study (Table S2). Datasets with quality assurance/quality control data (i.e., results from all method blanks, tissue blanks, and ongoing recovery and precision standards analyzed in parallel with all samples) and data for individual animals are openly available in Iowa Research Online at<http://doi.org/10.25820/data.006117>(Li et al., 2020). For Figures with PCB congener profiles, see Figs. S1–S4. The peak corresponding to PCB 138, PCB 158, PCB 163, and PCB 164 is abbreviated as PCB 138 throughout the manuscript.

2.5. Model prediction of the tissue-blood partitioning of PCBs

A composition-based model (Endo et al., 2013) was used for the prediction of the partition coefficients of PCBs between tissues and whole blood. In this model, tissues and blood are assumed to contain five biological components, including albumin, muscle protein, membrane lipid, storage lipid, and water. The volume fractions and lipid content of all

biological components, both in tissues and blood, were taken from published experimental values (Table S3) (Endo et al., 2013; Schmitt, 2008; Wu et al., 2015). The partitioning of PCBs between the biological components was determined with published Abraham solvation parameters (van Noort et al., 2010) and polyparameter linear free energy relationships (PP-LFERs) describing the correlations between the chemical descriptors and tissue/water partition coefficients (Endo et al., 2012; Endo et al., 2011; Endo and Goss, 2011; Geisler et al., 2012). The equation used for the calculations of tissue-blood partitioning coefficients ($K_{tissue/blood}$) were as follows (Endo et al., 2013):

$$
K_{tissue/blood} = \frac{K_{tissue/water}}{K_{blood/water}}
$$

=
$$
\frac{K_{ap/w}f_{ap}^{tissue} + K_{mp/w}f_{mp}^{tissue} + K_{sl/w}f_{sl}^{tissue} + K_{ml/w}f_{ml}^{tissue} + f_{w}^{tissue}}{K_{ap/w}f_{ap}^{blood} + K_{mp/w}f_{mp}^{blood} + K_{sl/w}f_{sl}^{blood} + K_{ml/w}f_{ml}^{blood} + f_{w}^{blood}}
$$
⁽¹⁾

Where $K_{tissue/water}$ and $K_{blood/water}$ are the partition coefficients of PCBs between tissue and water and between blood and water, respectively. $K_{ap/w}$, $K_{mp/w}$, $K_{s l/w}$ and $K_{m l/w}$ are the partition coefficients of PCBs between albumin protein and water, muscle protein and water, storage lipid and water, and membrane lipid and water, respectively. f_{ap}^{tissue} , f_{mp}^{tissue} , f_{sl}^{tissue} , f_{ml}^{tissue} and f_{w}^{tissue} are the volume fractions of albumin protein, muscle protein, storage lipid, membrane lipid, and water in tissue, respectively. f_{ap}^{blood} , f_{mp}^{blood} , f_{sl}^{blood} , f_{ml}^{blood} and f_{w}^{blood} are the volume fractions of albumin protein, muscle protein, storage lipid, membrane lipid, and water in the blood, respectively.

2.6. Data visualization and statistical analysis

PCB mass percentages and wet weight-adjusted tissue levels are presented as mean \pm standard deviation (Tables S4–S9). Median values are provided for experimental tissue partition coefficients. Differences in PCBs congener profiles were analyzed and visualized using the Heatmap function and Sparse partial least squares-discriminant analysis (sPLS-DA), as implement by MetaboAnalyst 4.0 (Chong et al., 2018). Significant differences between exposure groups were analyzes using Student's t-test (Tables S10–S13). Two-way ANOVA was used to compare the PCB profiles, based on tissue levels, with respect to homolog group, Class, or congener between sex or genotypes. If the interaction term, for instance, between classes and sex, is significant, then the profiles on classes are different between sex (Table S14). These analyses were performed with R (version 3.6.3). Similarity coefficients cos Θ comparing average congener profiles (Tables S15–S17) were calculated are described (Davis, 1986). The similarity coefficient uses the cosine of the angle between two multivariable vectors, such as PCB profiles). A value of 0.0 describes two completely different profiles, and 1.0 describes two identical profiles.

3. Results and Discussion

3.1. Levels of total PCBs (Σ**PCB) in tissues**

In WT mice, the Σ PCB levels followed the rank order adipose > liver ~ brain > blood (Fig. 1). This rank order is consistent with earlier studies reporting an accumulation of PCBs and

An enrichment of ΣPCB was observed in the liver of KO compared to WT mice, irrespective of the sex (M_{WT} vs. M_{KO} , $p=0.0010$; F_{WT} vs. F_{KO} , $p=0.0004$; Table S10). An enrichment of PCBs in the liver of KO mice was also observed when PCBs were analyzed on a homolog, Class, or congener basis, as discussed below. Similarly, an accumulation of PCB in the liver was observed in disposition studies of single PCB congener in KO and WT mice (Wu et al., 2015; Wu et al., 2020). The accumulation of PCBs in the liver of KO mice is a consequence of the higher lipid content and the elevated protein levels in the liver of KO compared to WT mice (Weng et al., 2005; Wu et al., 2015). The higher lipid content in the liver of KO mice is caused by an impaired hepatic fat metabolism (Gu et al., 2005; Weng et al., 2005). In addition to an increased lipid content in the liver, the expression of hepatic proteins that bind PCBs, such as cytochrome P450 enzymes (Kania-Korwel et al., 2008; Kennedy et al., 1981), is increased in the liver to compensate for the liver-specific deletion of cpr (Gu et al., 2003). Dioxin-like PCBs and structurally related compounds are sequestered in the liver because they bind to particular cytochrome P450 isoforms (Chen et al., 2003; Diliberto et al., 1999). It is likely that cytochrome P450 enzymes in the liver from KO mice bind PCBs in the absence of metabolic conversion, thus contributing to the hepatic accumulation of PCBs in KO mice.

3.2. Comparison of PCB homologs, Classes, and congeners across exposure groups

S5), PCB 91 or PCB 136 (Wu et al., 2015; Wu et al., 2020).

3.2.1. Distribution of PCB homologs in tissues from different exposure groups and Aroclor 1254—The homolog composition of the PCB residues from blood, brain, and liver, but not adipose tissue, showed differences compared to the homolog composition of Aroclor 1254 (Table S4). Typically, the percentage of lower chlorinated homologs decreased, whereas the percentage of higher chlorinated homologs increased in these tissues compared to Aroclor 1254. Different shifts in the homolog composition were observed for WT compared to KO mice. For example, hexachlorinated PCBs were the major constituents in the brain from F_{WT} mice, whereas pentachlorinated PCBs are the major homolog group in Aroclor 1254 (Table S4).

PCB homolog profiles expressed on a tissue wet weight basis (Fig. 1 and Table S7), or as a weight percentage (Table S4), also displayed genotype and sex-dependent differences. The homolog composition in the blood, brain, and liver was significantly different between F_{WT} and F_{KO} mice (Table S7 and S11). Differences in the homolog composition of M_{WT} and MKO mice were only observed in the liver. Sex differences in the homolog profiles were observed in the blood and liver of WT mice and the brain of KO mice.

The changes in the homolog mass profiles translated into sex and genotype-dependent differences in wet weight-adjusted homolog levels. Pronounced changes were observed for F_{WT} and F_{KO} mice (Fig. 1, Table S7, and S11). Levels of tetra- ($p=0.0128$) and

pentachlorinated PCBs ($p=0.0163$) were lower, and levels of heptachlorinated PCBs ($p=0.0189$) were higher in blood from F_{WT} mice compared to F_{KO} group (Fig. 1a2, Table S11). Moreover, levels of tetra-, penta-, hexa-, and heptachlorinated PCBs were lower in the brain of F_{WT} than F_{KO} mice (Fig. 1a3, Table S11). For example, levels of pentachlorinated PCBs in the brain of F_{KO} mice were 1,110±70 ng/g tissue, which is almost 2-fold higher than the levels in F_{WT} mice (630±70 ng/g tissue) (Table S7).

Consistent with the higher ΣPCB levels described above, the levels of all homolog groups were more elevated in the liver of F_{KO} compared to F_{WT} mice (p 0.0020). Similarly, the levels of all PCB homologs were higher in M_{KO} vs. M_{WT} mice (p 0.0374).

3.2.2. Distribution of PCB Classes in tissues from different exposure groups and Aroclor 1254—Although lower chlorinated PCBs are generally more rapidly metabolized than higher chlorinated PCBs, the metabolism and elimination of PCBs also depend on the chlorine substitution pattern (Birnbaum, 1985). Therefore, we compared differences in the mass percentage of PCBs congeners with a 4-, 3,4-, and 3,4,5- substitution pattern and zero or one ortho chlorine substituent (Class A); PCB congeners with two or more ortho chlorine substituents and a 2,4- or 2,3,4- substitution pattern (Class B); PCBs congeners with a 2,4,5-substitution pattern (Class C); and PCB congeners that are considered to be episodic because they are rapidly eliminated following exposure (Imsilp and Hansen, 2005). Class D congeners have adjacent, unsubstituted C-atoms and, typically, contain a 2-, 2,3-, 2,5- and 2,3,6-substitution pattern in at least one phenyl ring (Class D) (Haraguchi et al., 2005; Matthews and Tuey, 1980). Overall, the mass percentage of PCBs belonging to Class C increased relative to Aroclor 1254 in all tissues from all four exposure groups (Fig. 1; Table S5).

We also observed significant differences in the distribution of PCB Classes, expressed both as tissue levels (Fig. 1; Table S8) and weight percentage (Table S5), by genotype and sex. For example, the levels (Fig. 1) and weight percentage (Table S5) of more persistent PCBs (i.e., Classes A, B, and C) appeared to be higher in the blood and brain of WT compared to KO mice. When comparing levels in F_{WT} and F_{KO} mice, this difference was statistically significant for Class A PCBs in the brain ($p=0.0096$), for Class B and Class C PCBs in the blood ($p=0.0319$ and $p=0.0333$, respectively), and for Class D PCBs in the blood ($p=0.0044$) and the brain ($p<0.0001$).

3.2.3. Distribution of PCB congeners in adipose tissue from different

exposure groups and Aroclor 1254—Differences in PCB profiles in adipose and other tissues (see below), organized by genotype, and Aroclor 1254 are shown in Fig. 2 (for a comparison of PCB levels by genotype, see Fig. S6; for a comparison of PCB profiles by tissue from mice within an exposure group, see Fig. S7). PCB congener profiles in adipose tissues were different from the PCB profile of Aroclor 1254 (cos $\Theta = 0.84$ to 0.88; Table S15). In contrast, PCB profiles in adipose tissue were quite similar across all four exposure groups (cos Θ = 0.97 to 1.0; Table S16), with the most pronounced difference observed between F_{WT} and F_{KO} mice. Further analysis of individual PCB congeners revealed minor genotype and, for some congeners, sex-dependent differences in the adipose tissue (Fig. 2a). Persistent PCB congeners, for example, PCB 118 (Class A) and PCB 180 (Class C),

contained a relatively higher percentage in adipose tissue from WT mice. In contrast, PCB congeners belonging to Class D, such as PCB 70, PCB 95, and PCB 136, showed the opposite trend, with lower percentages of these PCB congeners being present in the adipose tissue from WT compared to KO mice. Moreover, the percentage of many Class D congeners were lower in F_{WT} mice compared to the other exposure groups.

Analysis of the PCB congener composition, based on tissue wet weights, revealed significant differences only between F_{WT} and F_{KO} mice ($p \le 0.001$; Table S9 and S13). Consistent with the changes in the mass profiles, tissue levels of PCB 70, a Class D congener, were lower in F_{WT} than F_{KO} mice (600 \pm 200 ng/g and 1,100 \pm 300 ng/g, respectively; $p=0.0142$), whereas levels of PCB 180, a Class C congener, were higher in F_{WT} than F_{KO} mice (250±90 ng/g vs. 120±20 ng/g, respectively; $p=0.0307$).

3.2.4. Distribution of PCB congeners in whole blood from different exposure

groups and Aroclor 1254—The main PCB congeners detected in blood from WT and KO mice are also frequently found in human serum (Grimm et al., 2015; Sethi et al., 2019). The overall PCB congener profiles in blood were different from Aroclor 1254 (cos $\Theta = 0.72$) to 0.74; Table S15) and revealed minor genotype-dependent differences in the blood (cos Θ $= 0.98$ to 0.99, Table S16). F_{WT} mice were an exception and displayed a PCB congener profile that was different from the profiles observed in the blood from the other exposure groups (cos Θ = 0.92 to 0.97). A comparison of the wet weight-based PCB congener composition also revealed statistically significant differences between F_{WT} and F_{KO} mice ($p \le 0.0001$) and F_{WT} and M_{WT} mice ($p \le 0.0001$) (Table S14).

The percentage of some congeners (e.g., PCB 70) was higher in KO compared to WT mice (Fig. 2b). Different congeners (e.g., PCB 99 and PCB 180) had a lower percentage in KO compared to WT mice. In the blood from F_{WT} mice, PCB congeners belonging to Class D were present at a lower mass percentage, and more persistent PCB congeners from Classes A, B, and C were present at a higher mass percentage compared to M_{WT} , F_{KO} , and M_{KO} . Similar trends between KO and WT mice were observed for these congeners when comparing PCB levels adjusted for wet weight (Fig. S6b). For example, PCB 95 levels, expressed on a wet weight basis, were lower in the blood from F_{WT} mice $(8\pm 1 \text{ ng/g})$ compared to F_{KO} (14 \pm 3 ng/g; p =0.0034) and M_{WT} mice (13 \pm 3 ng/g; p =0.0163).

3.2.5. Distribution of PCB congeners in the brain from different exposure groups and Aroclor 1254—PCB 118, PCBs 132/153, and PCB 138 were major PCB congeners detected in the brain of WT and KO mice. These PCBs were also prevalent PCB congeners detected in the brain from postnatal day 31 rats exposed to Aroclor 1254 throughout development via the maternal (Yang et al., 2009). PCB 138, PCB 153, and PCB 118 were also major PCB congeners in postmortem human brain samples (Dewailly et al., 1999; Mitchell et al., 2012). Unlike the present study, these studies analyzed a small set of PCB congeners and, thus, provide limited insights into the congener profiles present in the rodent or human brain.

PCB residues in the brain from all exposure groups were distinctively different from the Aroclor 1254 profile (cos Θ =0.80 to 0.86; Table S15) and showed genotype and sex-

dependent differences (cos $\Theta = 0.96$ to 0.99; Table S16) (Fig. 2c). A comparison of the wet weight-based PCB congener composition also revealed statistically significant differences between F_{WT} and F_{KO} mice ($p \le 0.0001$) and F_{KO} and M_{KO} mice ($p \le 0.0001$) (Table S14). Based on their mass percentage and tissue wet weights, many congeners belonging to Class D (e.g., PCB 136) were lower in F_{WT} mice compared to the other exposure groups, especially F_{KO} mice, an observation that is toxicologically relevant. For example, PCB 95 and PCB 136 are potent sensitizers of the ryanodine receptors (Pessah et al., 2006), cellular targets implicated in PCB-induced developmental neurotoxicity (Holland et al., 2016; Pessah et al., 2010; Pessah et al., 2019). PCB mixtures (i.e., Aroclor 1254) and PCB 95 display a non-monotonic dose-response relationship in rodent studies of PCB developmental neurotoxicity (Wayman et al., 2012; Yang et al., 2009). Thus, depending on the experimental design and the dose, relatively small differences in the levels of neurotoxic PCB congeners in the developing brain may significantly affect neurotoxic outcomes. In the present study, levels of PCB 95 and PCB 136 were 1.4- and 3.0-fold higher in the brain from F_{KO} compared to F_{WT} mice, respectively ($p \le 0.0001$ and $p \le 0.0001$, respectively; Tables S9 and 14).

3.2.6. Distribution of PCB congeners in the liver from different exposure groups and Aroclor 1254—PCB profiles in the liver were different from the profile observed for Aroclor 1254 (Fig. 2d) (cos $\Theta = 0.77$ to 0.88; Table S15). A comparison of PCB congener profiles in the liver of M_{WT} , M_{KO} , and F_{KO} mice showed only small differences (cos $\Theta = 0.97$ to 1.0; Table S16). As with blood and brain tissue, more pronounced differences were observed for F_{WT} mice compared to the other exposure groups (cos Θ = 0.92 to 0.93; Table S16). A comparison of the PCB congener composition, expressed as tissue weight-adjusted tissue levels, revealed significant differences between KO and WT mice of the same sex ($p<0.001$ for both male and female mice). Moreover, levels of individual PCB congeners were always higher in KO compared to WT mice, irrespective of the sex, because of the higher fat and protein content of the liver of KO mice (Fig. S6d) (Gu et al., 2005; Gu et al., 2003; Weng et al., 2005).

There were also differences in the PCB congener composition between male and female mice of the same genotype ($p=0.0430$ and $p<0.0001$ when comparing WT and KO mice, respectively) (Table S14). Female mice tended to have higher levels of specific PCB congeners in the liver than male mice of the same genotype (p >0.05; Tables S9 and S13). These differences are consistent with the increased susceptibility of female mice to PCBinduced hepatotoxicity (Wahlang et al., 2019).

3.2.7. Factors contributing to sex and genotype-dependent differences in the PCB tissue distribution—The shifts in the composition of PCB homologs, Classes, and individual congeners are driven by several factors. As discussed above, the liver-specific deletion of CPR changes the liver composition in KO mice, thus resulting in a pronounced accumulation of PCBs in the liver of KO mice. Besides, many of the observed changes in the PCB profiles in tissues are due to the more rapid metabolism of lower chlorinated PCBs, in particular congeners belonging to Class D, in both WT and KO mice (Lucier et al., 1978; Matthews and Anderson, 1975). A relative decrease in lower chlorinated PCBs and an

increase in higher chlorinated PCB in WT compared to KO mice is consistent with the impaired hepatic metabolism caused by the lack of CPR activity in the liver KO mice (Weng et al., 2005; Wu et al., 2015). Besides, differences in the initial distribution and redistribution of PCBs to the adipose tissue, as well as their (extrahepatic) metabolism (Saghir et al., 1999), likely contribute to subtle differences in the PCB distribution at the time point investigated.

Our study also observed subtle sex differences in the disposition of PCBs, especially in WT mice. Such sex differences, especially in the liver (Fig. 1), are consistent with the higher expression of Cyp2b10, a cytochrome P450 isoform implicated in the metabolism on PCB congeners belonging to Class D (Wu et al., 2013a), and other hepatic and extrahepatic cytochrome P450 enzymes in the female compared to the male mouse liver (Renaud et al., 2011). Indeed, PCB 136 is more rapidly oxidized in precision-cut tissue slices from male compared to female rats (Wu et al., 2013b). However, no sex-dependent differences in PCB levels have been reported in earlier disposition studies in rodents after developmental PCB exposure (Dziennis et al., 2008; Kania-Korwel et al., 2017; Miller et al., 2010; Yang et al., 2009). Alternatively, sex differences in the wet weight-adjusted PCB levels in the liver may be due to different fat content in the liver of female vs. male mice or other factors that we could not account for this the present study.

3.3. Differences in PCB profiles in tissues from the same exposure group

Laboratory and biomonitoring studies generally assume that the profiles and levels of PCB in the blood approximate tissue profiles and levels. For example, the distribution of PCBs in juvenile rats shows that serum levels are predictive of liver levels (Soontornchat et al., 1994); however, studies that investigate the congener-specific distribution of PCBs across tissues are limited, especially for lower chlorinated PCBs. In the present study, PCB profiles displayed modest differences between tissues within each exposure group, as shown in heatmaps organized by tissue (Fig. S7). Tissue-to-tissue similarity coefficients ranged from 0.85 to 0.98 (Table S17; Fig. S7). Several other animal studies also reported modest intertissue differences in the PCB profiles after oral or inhalation exposure (cos Θ = 0.80) (Hu et al., 2015; Hu et al., 2012; Kodavanti et al., 1998). These comparisons support the assumption that PCB profiles in the blood are a reasonable approximation of PCB tissue profiles in mice and other mammals.

sPLS-DA was used to investigate differences in the PCB profiles across tissues within the same exposure group (Fig. 3; Figs. S8–S10). sPLS-DA comparing the PCB congener profiles from FWT mice showed clear groupings according to tissue, with three principal components (PCs) accounting for 84.6% of the data variance (Fig. 3). PC1 separated adipose tissue from the brain and liver. Based on their loadings, PCB congeners belonging to Class D (i.e., PCB 52 and PCB 70) were higher in adipose tissue, but lower in brain and liver. PCBs belonging to Class B (i.e., PCB 138) and Class C (i.e., PCB 146 and PCB 183) made a lower contribution to the PCB profile in adipose tissue, but a higher contribution in the brain and liver. PC2 separated blood from adipose tissue, with some Class D congener (i.e., PCB 139/149 and PCB 151) being higher in blood, but lower in adipose tissue and liver. PC3 separated the liver from other tissues because of lower relative levels of Class D congeners

(i.e., PCBs 134/143 and PCB 136). Similar differences in the PCB congener profiles across tissues were observed for the other exposure groups despite the impaired hepatic metabolism in KO mice (Figs. S8–S10). Overall, differences in the PCB congener profiles across tissues from mice from the same exposure group were due to differences in the percentage of metabolically more labile PCB congeners (Class D) compared to congeners resistant to metabolites (Class A through C). In addition, time-course experiments suggest that the initial distribution vs. redistribution from highly perfused to storage tissues also influences how PCB congener profiles change over time (Birnbaum, 1983; Hu et al., 2010).

3.4. Tissue-to-blood distribution of PCBs within each exposure group

Tissue : blood ratios, a measure of how much PCB is distributed from the blood into a target tissue, followed the rank order adipose : blood (median: 201; range: 12–841) >> liver : blood (median: 22 ; range: $7-213$) > brain : blood (median: 11; range: $3-34$) in WT mice. This rank order is consistent with other studies in mice and other mammalian models (Lutz et al., 1984). The differences between liver : blood vs. brain : blood partition coefficients were more pronounced in KO mice because of the higher lipid content in the liver, with adipose : blood (median: 71; range: $40-470$) ~ liver : blood (median: 91; range $47-481$) >> brain : blood (median: 5; range 3–28). With the exception of the liver : blood partition coefficients in KO mice, the partition coefficients in this study are within the range of partition coefficients observed in animal studies and in humans (Lutz et al., 1984; Wolff et al., 1982). For example, adipose to plasma partition coefficients of 24 PCB congeners in occupationally exposed individuals ranged from 50 to 370 (Wolff et al., 1982).

The partitioning of PCBs between tissues and blood changed with increasing degree of chlorination. For example, tetra- to hexachlorinated PCBs appeared to have a much higher affinity for adipose tissue than higher chlorinated homologs in F_{WT} mice at the time point investigated (Fig. 4). The opposite was true for the brain, where the partitioning into the brain appeared to increase with an increase in chlorination. The partitioning of PCBs from the blood into the liver from F_{WT} mice seemed to be comparable across homolog groups. Similar trends were observed for M_{WT} , M_{KO} , and F_{KO} mice (Figs. S11–S13). The decrease in the adipose : blood partitioning by the homolog group is unexpected based on a structurebased model of the tissue-to-blood partitioning of PCBs (Parham et al., 1997). Besides, the partitioning of individual congeners from the blood into tissues was congener-specific and displayed considerable variability (Figs. 4a–c). Thus, congener specific differences in the blood-to-tissue partitioning exist at specific time points that may influence toxic outcomes following PCB exposure. These differences are not captured when, as discussed above, only PCB profiles are compared between tissues. Especially systematic studies of tissue : blood ratios in human postmortem samples remain limited (Artacho-Cordón et al., 2015; Ploteau et al., 2016; Whitcomb et al., 2005), and further congener-specific studies are needed, especially for lower chlorinated PCB congeners that, unlike persistent legacy PCBs, are still inadvertently produced and represent a current public health concern (Grimm et al., 2015).

3.5. Comparison of the experimental and theoretical tissue-to-blood distribution of PCBs

The theoretical tissue : blood ratios, determined with a published model (Endo et al., 2013), are shown for all 209 PCB congeners in Fig. 4d. The model did not predict the congener-

specific variability of the partition coefficients. Moreover, the model predicted an increase, not a decrease in the adipose : blood ratios with increasing degree of chlorination. These differences between the calculated and experimental values are not entirely surprising because the model does not take structural features into account that, for example, influence the rate of metabolism of individual PCB congeners. Moreover, individual PCBs probably had not reached steady-state levels in our study. Although PCBs, like other lipophilic compounds, are expected to readily partition into the brain, the model overestimated the partitioning of PCBs into the brain. Similarly, earlier studies noted that the levels of PCBs in the brain are lower than expected based on the lipid content of the brain due to the unique lipid composition of the brain (Bachour et al., 1998; Dewailly et al., 1999). Besides, the distribution on PCB 153 in the rat brain is not uniform (Saghir et al., 2000). These differences in local PCB levels could also be due to brain region-specific variations in the tissue composition.

Despite these differences, the predicted and experimental partition coefficients were in reasonable agreement with each other (i.e., deviations were within one order of magnitude, Fig. 4e) and matched the calculated adipose : blood partition coefficients in rats (Endo et al., 2013). The model correctly predicted the more pronounced partitioning of PCBs into the liver of KO mice, as indicated by the distinct clusters of the liver : blood partition coefficients of KO vs. WT mice in Fig. 4e. While not perfect, composition-based models are, therefore, an effective tool to approximate tissue levels based on blood PCB levels, which are experimentally more accessible, especially in humans. However, more detailed, PCB congener-specific studies of the PCB levels in human tissues and blood from the same donors are needed to establish whether or not the model used in our research also allows a rough estimate of PCB tissue levels based on experimentally determined blood levels. Additional work is also necessary to better predict the partitioning of PCBs into brain tissues.

4. Conclusions

The present study investigated how an altered hepatic metabolism and a fatty liver caused by the liver-specific deletion of cytochrome P450 reductase affect PCB distribution in mice. The KO mouse model used in this study was developed for research on the role of hepatic metabolism on the systemic toxicity of toxicants; however, the impaired hepatic metabolism also results in a fatty liver and other compensatory changes in the liver and other organs. The impaired hepatic metabolism affected the disposition of PCBs, consistent with earlier studies with individual PCB congeners. However, the altered composition of the liver had a much more prominent effect on the distribution of PCBs, with wet weight-adjusted PCB levels being similar in the liver and adipose tissue from KO mice. In contrast, PCB levels followed the rank order adipose > liver in WT mice. This observation demonstrates that compensatory changes in genetic models can result in an altered disposition of toxicants, such as PCBs, and represent a potential pitfall of genetic models for mechanistic toxicity studies. It is, therefore, essential to assess the disposition of a toxicant when working with a genetic model. In silico approaches, for example, the composition-based model used in this study, represent a time and cost-effective alternative to the measurement of the levels of a toxicant. Despite the difference in the experimental and theoretical partition ratios, our model

provided a reasonable approximation of the partition coefficients of all 209 PCB congeners and predicted the more pronounced accumulation of PCBs in the liver of KO mice. This approach can be used to approximate the distribution of PCBs into the liver of genetic steatosis models, such as ob/ob mice, and diet-induced obesity models and, thus, aid in the interpretation of toxicity studies with PCBs and other lipophilic compounds.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- **•** Fatty liver and impaired hepatic metabolism alter PCB tissue distribution in mice
- **•** PCBs accumulate in the liver of cytochrome P450 reductase-null mice
- **•** Tissue partitioning of PCBs does not always increase with degree of chlorination
- **•** Composition-based models approximate the magnitude of PCBs' tissue partitioning

Fig. 1.

PCB homolog composition, based on PCB homolog tissue levels (μ g/g wet weight), in (a1) adipose, (a2) blood, (a3) brain, and (a4) liver and PCB Class composition in (b1) adipose, (b2) blood, (b3) brain, and (b4) liver showed statistically significant genotype and sexdependent differences, as determined using two-way ANOVA. Total PCB levels (ΣPCB, μg/g wet weight) showed statistically significant differences between WT and KO mice in the male liver ($p=0.0010$), female liver ($p=0.0004$), and female brain ($p=0.0002$) (Table S13). The composition of Aroclor 1254 by homolog group and Class is shown for comparison in panels (a5) and (b5). Class A: PCBs congeners with a 4-, 3,4-, and 3,4,5 substitution pattern and zero or one ortho chlorine substituent. Class B: PCB congeners with two or more ortho chlorine substituents and a 2,4- or 2,3,4- substitution pattern. Class C: PCBs congeners with a 2,4,5-substitution pattern. Class D: PCB congeners with, in contrast to Classes A through C, are readily metabolized and, for example, a 2-, 2,3-, 2,5- and 2,3,6 substitution pattern in at least one phenyl ring (Haraguchi et al., 2005). Levels of PCB congeners were measured by GC-MS in tissues from male wild-type $(M_{WT}, n=3)$, male knockout (M_{KO} , n=5), female wild-type (F_{WT} , n=4), and female knockout (F_{KO} , n=4) mice exposed orally to Aroclor 1254.

Fig. 2.

A comparison of the PCB congener profiles in (a) adipose tissue, (b) blood, (c) brain and (d) liver from male wild-type (M_{WT}, n=3), male knockout (M_{KO}, n=5), female wild-type (F_{WT}, $n=4$), and female knockout (F_{KO} , $n=4$) mice reveals genotype-dependent differences in the distribution of individual PCB congeners in these tissues. The congener profile of Aroclor 1254 is shown for comparison. The small panels illustrate differences in the mass percentages of representative PCB congeners. These congeners shown here were selected because they illustrate specific differences in the mass composition and, in most cases, are also toxicologically relevant. Heatmaps were generated after removing features with >25% missing values and autoscaling features of the original data using the Heatmap function as implement by MetaboAnalyst 4.0 (Chong et al., 2018). For PCB congener profiles and tissue levels, see the Supplementary material or (Li et al., 2020), respectively. A, Aroclor 1254.

Fig. 3.

Sparse partial least squares - discriminant analysis (sPLS-DA)(Lê Cao et al., 2011) revealed differences in the PCB congener profiles between tissues from F_{WT} mice (n=4), with three principal components (PCs) accounting for 84.6% of the data variance. These differences were due to tissue-specific changes in the relative levels of PCB congeners that are more readily metabolized (Class D) vs. PCB congeners that are more resistant to metabolism (Classes A through C). PC1 separated adipose tissue from the brain and liver; PC2 separated blood from adipose tissue; PC3 separated the liver from other tissues. sPLS-DA was performed with PCB congener profiles using MetaboAnalyst 4.0 after removing variables for a threshold of 25%, cubic root transformation, and autoscaling features (Chong et al., 2018). For analogous analyses of PCB congener profiles from $M_{\rm WT}$, $M_{\rm KO}$, and $F_{\rm KO}$ mice, see Figs. S8–S10 in the Supplementary material.

Fig. 4.

Tissue : blood ratios ($K_{tissue/blood}$, $(ng/g)/(ng/g)$) of different PCB homolog groups in female wild type mice (F_{WT} , n=4) show different trends with an increasing degree of chlorination: (a) Adipose : blood ratios decrease, (b) brain : blood ratios increase, and (c) liver : blood ratios of PCBs remain comparable with increasing degree of chlorination. Similar trends were observed with all other exposure groups (Figures S11–S13). (d) In contrast to the experimental findings, predicted tissue : blood ratios (i.e., adipose : blood, brain : blood, and liver : blood ratios) increase with an increasing degree of chlorination. Data for the tissue composition were obtained from the literature (Endo et al., 2013). (e) Despite this discrepancy, the predicted tissue : blood ratios are in reasonable agreement with the experimental values (for the underlying tissue composition data, see Table S3). Each symbol represents the tissue : blood ratio of an individual PCB congener (Li et al., 2020).