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Sexual Dimorphism in the Expression of Cytochrome P450 Enzymes in Rat Heart, Liver, Kidney, Lung, Brain, and Small Intestine^S

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ABSTRACT

Cytochrome P450 (P450) enzymes are monooxygenases that are expressed hepatically and extrahepatically and play an essential role in xenobiotic metabolism. Substantial scientific evidence indicates sexspecific differences between males and females in disease patterns and drug responses, which could be attributed, even partly, to differences in the expression and/or activity levels of P450 enzymes in different organs. In this study, we compared the mRNA and protein expression of P450 enzymes in different organs of male and female Sprague-Dawley rats by real-time polymerase chain reaction and western blot techniques. We found significant sex- and organ-specific differences in several enzymes. Hepatic Cyp2c11, Cyp2c13, and Cyp4a2 showed male-specific expression, whereas Cyp2c12 showed female-specific expression. Cyp2e1 and Cyp4f enzymes demonstrated higher expression in the female heart and kidneys compared with males; however, they showed no significant sexual dimorphism in the liver. Male rats showed higher hepatic and renal Cyp1b1 levels. All assessed enzymes were found in the liver, but some were not expressed in other organs. At the protein expression level, CYP1A2, CYP3A, and CYP4A1 demonstrated higher expression levels in the females in several organs, including the liver. Elucidating sex-specific differences in P450 enzyme levels could help better understand differences in disease pathogeneses and drug responses between males and females and thus improve treatment strategies.

SIGNIFICANCE STATEMENT

This study characterized the differences in the mRNA and protein expression levels of different cytochrome P450 (P450) enzymes between male and female rats in the heart, liver, lung, kidney, brain, and small intestine. It demonstrated unique sex-specific differences in the different organs. This study is considered a big step towards elucidating sex-specific differences in P450 enzyme levels, which is largely important for achieving a better understanding of the differences between males and females in the disease's processes and treatment outcomes.

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Introduction

Cytochrome P450 (P450) is a superfamilyof membrane-bound hydrophobic heme enzymes that play a pivotal role in health, homeostasis, and metabolism. P450 enzymes are expressed in almost all biologic systems (El-Sherbeni and El-Kadi, 2017). They are so called because their heme pigment absorbs light at a wavelength of 450 nm following reduction and exposure to carbon monoxide (Lynch and Price, 2007). The discovery of P450 enzymes started in the early 1950s and continued until the 1960s (El-Sherbeni and El-Kadi, 2017). P450 enzymes are classified into families, subfamilies, and individual enzymes based on the structural homology of their amino acid sequences (Nebert et al., 1987; Elbekai and El-Kadi, 2006). Microsomal P450s, which are attached to the endoplasmic reticulum membrane, comprise the majority of human P450 enzymes and catalyze a wide array of biologic reactions (El-Sherbeni and El-Kadi, 2017).

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P450 enzymes are mainly expressed in the liver, but they are also found in several extrahepatic tissues, including the heart (Elbekai and El-Kadi, 2006), kidney (Fan et al., 2015), lungs (Hukkanen et al., 2002), brain (Dutheil et al., 2008), and other tissues (Zhu and Zhang, 2012; Alonso et al., 2015; Ibrahim et al., 2020). The induction or inhibition of P450 enzymes by xenobiotics or disease states is a major mechanism underlying drug-drug and drug-disease interactions. Moreover, genetic polymorphisms in P450 genes could lead to differences in P450 enzymes that might explain individual and ethnic variations in disease pathogeneses and drug responses (Manikandan and Nagini, 2018). Given their significant biologic effects, P450 enzymes have been the focus of many clinical, experimental, and drug development studies.

P450 enzymes play an essential role in the detoxification and activation of both xenobiotics and endogenous molecules. In humans, there are 18 P450 families with more than 50 individual P450 isoenzymes, nine of which are involved in the metabolism of several drugs (CYP1A2, CYP2C9, CYP2C19, CYP2D6, CYP3A4, and CYP3A5) (Wilkinson, 2005). In addition to their essential role in xenobiotic metabolism, P450 enzymes are also largely involved in the synthesis and metabolism of endogenous molecules such as steroids and prostaglandins. For example, P450s metabolize polyunsaturated fatty acids like arachidonic acid (AA) by the insertion of either an epoxide or hydroxyl group, based on which they can be classified into P450 epoxygenases or hydroxylases, respectively. AA metabolism by P450 enzymes to give multiple epoxy and

ABBREVIATIONS: AA, arachidonic acid; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; GH, growth hormone; P450, cytochrome P450; PCR, polymerase chain reaction; SD, Sprague-Dawley.

TABLE 1
Rat cytochrome P450 enzymes and their human orthologs

Rat gene	Human Ortholog
Cyp1a1	CYP1A1
Cyp1a2	CYP1A2
Cyp1b1	CYP1B1
Cyp2a1	· —
Cyp2b1	CYP2B6
Cyp2b2	_
Cyp2c6	CYP2C19
Cyp2c11	CYP2C9
Cyp2c12	_
Cyp2c13	_
Cyp2c23	_
Cyp2d2	_
Cyp2d3	CYP2D6
Cyp2d4	_
Cyp2e1	CYP2E1
Cyp2j3	_
Cyp2j4	CYP2J2
Cyp2j10	_
Cyp3a1	CYP3A5
<i>Cyp3a2</i>	-
Cyp3a9	CYP3A7
Cyp3a18	_
Cyp3a23	CYP3A5
Cyp4a1	CYP4A11
Cyp4a2	_
Cyp4a3	_
Cyp4a8	_
Cyp4f1	CYP4F12
Cyp4f4	CYP4F8
Cyp4f5	_
Cyp4f6	CYP4F3

hydroxy metabolites has been extensively studied in health and diseases, particularly in the cardiovascular system (Elbekai and El-Kadi, 2006; Shoieb et al., 2019; Gerges and El-Kadi, 2022).

To date, a large body of evidence points out to significant differences between males and females in the pathogenesis and outcomes of different diseases, as well as in drug metabolism and responses (Dahan et al., 2008; Waxman and Holloway, 2009; Ngo et al., 2014; Regitz-Zagrosek and Kararigas, 2017; Holingue et al., 2020; Tramunt et al., 2020; Madla et al., 2021; Gerges and El-Kadi, 2022). Moreover, it was established previously that there are significant sex differences in the expression or activity of several drugmetabolizing enzymes in animals and humans and that this could be responsible for differences in clinical drug effects between men and women (Waxman and Holloway, 2009). For example, multiple studies have shown higher mRNA and protein expression levels of hepatic CYP3A4 in women than in men, which could explain the higher clearance rates of CYP3A4 substrates in women (Tanaka, 1999; Greenblatt and Von Moltke, 2008; Waxman and Holloway, 2009). On the other hand, some CYP1A2, CYP2E1, and CYP2D6 substrates were found to have higher clearance rates in men than in women (Franconi et al., 2007; Schwartz, 2007). Cardiovascular diseases are among the diseases that demonstrate significant sex-specific discrepancies, which could be mediated, even in part, by different expression or activity levels of cardiac P450 enzymes and their metabolites (Gerges and El-Kadi, 2022).

Elucidating sex differences in the expression levels of P450 enzymes in different organs could help explain observed sex differences in diseases and drug effects, decrease the incidence of adverse effects and improve the efficacy of different medications, and approach precision medicine. Thus, the current study is one of a series of studies aiming at investigating sex-specific differences in the expression and activity levels of different P450s, as well as the levels of their metabolites. The aim of the current study was to compare the mRNA and protein expression levels of different P450 enzymes in the heart, liver, lung, kidney, brain, and small intestine between male and female rats.

TABLE 2 Rat primer sequences

Gene	Forward Primer	Reverse Primer
Cyp1a1	CCAAACGAGTTCCGGCCT	TGCCCAAACCAAAGAGAATGA
Cyp1a2	CGCCCAGAGCGGTTTCTTA	TCCCAAGCCGAAGAGCATC
Cyp1b1	GCTTTACTGTGCAAGGGAGACA	GGAAGGAGGATTCAAGTCAGGA
Cyp2a1	CACAGGGCAGCTCTATGACA	CAGACCCAGCAAAGAAGAGG
Cyp2b1	AACCCTTGATGACCGCAGTAAA	TGTGGTACTCCAATAGGGACAAGATC
Cyp2b2	CCATCCCTTGATGATCGTACCA	AATTGGGGCAAGATCTGCAAA
Cyp2c6	CCTGCTGAAGTGTCCAGAGG	CCCATCTAAAAAGTGGCCAG
Cyp2c11	CACCAGCTATCAGTGGATTTGG	GTCTGCCCTTTGCACAGGAA
Cyp2c12	TATAAACTCAATACGTTCTGAG	TTTTACATTAACTTCAGAAACTG
Cyp2c13	CTGGCAATCATGGTGACTGA	GAAACTCCTTGCTGTCATGC
Cyp2c23	GATGCTGTCTTCCGTCATGC	GTAATAGGCTTGATGTCAAG
Cyp2d2	CTACTGCCCATCTATAATCA	CCAAAGCTCTCCTTCAATGT
Cyp2d3	ACCAATGCTGTCATCCATGAGGT	GCTGGACTAGAATTTCTTTCCTT
Cyp2d4	GACCAGTCGGGCTTTGGACCAC	CGAAGGCCTTCTTTCCAGAG
Cyp2e1	AAAGCGTGTGTGTTGGAGAA	AGAGACTTCAGGTTAAAATGCTGCA
Cyp2j3	CATTGAGCTCACAAGTGGCTTT	CAATTCCTAGGCTGTGATGTCG
Cyp2j4	GCTCGGACCTTCATTCCACA	GATCGTGGCTACCAGAGAGC
Cyp2j10	TTGAACTTAGCAGAGGGGCTG	TCATACTCAAAGCGCTCCCC
Cyp3a1	GGAAATTCGATGTGGAGTGC	AGGTTTGCCTTTCTCTTGCC
Cyp3a2	GCTCTTGATGCATGGTTAAAGATTTG	ATCACAGACCTTGCCAACTCCTT
Cyp3a9	GGACGATTCTTGCTTACAGG	ATGCTGGTGGGCTTGCCTTC
Cyp3a18	CAACTACGGTGATGGCATGT	CACTCGGTTCTTCTGGTTTG
Cyp3a23	ATGTTCCCTGTCATCGAACAGTATG	TTCACAGGGACAGGTTTGCCT
Cyp4a1	TTGAGCTACTGCCAGATCCCAC	CCCATTTTTGGACTTCAGCACA
Cyp4a2	CTCGCCATAGCCATGCTTATC	CCTTCAGCTCATTCATGGCAATT
Cyp4a3	CTCGCCATAGCCATGCTTATC	CCTTCAGCTCATTCATGGCAATC
Cyp4a8	TGTGGTATCATGAGTGGCTCG	CTTCAGCACGCAGGTCCTTA
Cyp4f1	CCCCCAAGGCTTTTTGATG	GAGCGCAACGGCAGCT
Cyp4f4	CAGGTCTGAAGCAGGTAACTAAGC	CCGTCAGGGTGGCACAGAGT
Cyp4f5	AGGATGCCGTGGCTAACTG	GGCTCCAAGCAGCAGAAGA
Cyp4f6	TCACTTGACCTTGATGAAGAACAAC	AAGAGAGGTGGATATCACGGAAG
B-actin	CCAGATCATGTTTGAGACCTTCAA	GTGGTACGACCAGAGGCATACA
Gapdh	CAAGGTCATCCATGACAACTTTG	GGGCCATCCACAGTCTTCTG

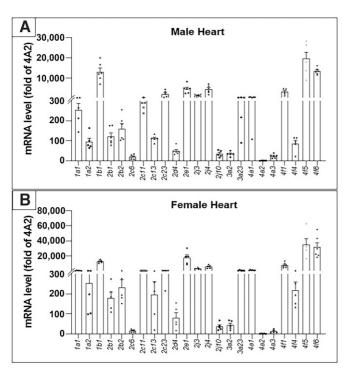


Fig. 1. The mRNA expression of different P450 enzymes in male (A) and female (B) rat heart relative to the least expressed. The mRNA expression of P450 enzymes was determined in the heart of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β-actin housekeeping gene. Results are presented as mean plus or minus S.E.M, n = 4-6.

Material and Methods

Nomenclature. The nomenclature used throughout the manuscript is following the Guidelines for Formatting Gene and Protein Names, released in 2014. Briefly,

enzyme symbols were written in sentence case and italicized when referring to genes or mRNA of mice or rats and were capitalized and nonitalicized when referring to the proteins (https://www.biosciencewriters.com/Guidelines-for-Formatting-Gene-and-Protein-Names.aspx).

Animals. Adult (8 weeks old) male (260–280 g, n=6) and female (200–220g, n=6) Sprague-Dawley (SD) rats were purchased from Charles River Canada (Montreal, QC, Canada). All animals were allowed access to food and water ad libitum throughout the experiment period and were maintained on a 12-hour light/dark cycle. Rats were kept in the animal facility for an acclimatization period of 1 week, after which they were euthanized under isoflurane anesthesia. The liver, heart, lung, kidney, brain, and small intestine (20 cm extending from the stomach distally) were isolated and immediately frozen in liquid nitrogen and then stored at -80° C. All procedures involving experimental animals were performed in accordance with the Guide for the Care and Use of Laboratory Animals as adopted and promulgated by the US National Institutes of Health and were approved by Alberta Health Sciences Animal Policy and Welfare Committee.

Chemicals. The TRIzol reagent used for mRNA extraction was Invitrogen brand (Thermo Fisher Scientific, Carlsbad, CA). High Capacity cDNA Reverse Transcription Kit and SYBR Green PCR Master Mix were purchased from Applied Biosystems (Foster City, CA). Real-time polymerase chain reaction (PCR) primers were formulated by and purchased from Integrated DNA Technologies (Coralville, IA). Trans-Blot Turbo RTA Transfer Kit and 2X Laemmli Sample Buffer were purchased from Bio-Rad Laboratories (Hercules, CA). CYP1A2, CYP3A, and CYP4A1 mouse monoclonal primary antibodies were purchased from Santa Cruz Biotechnology (Dallas, TX); CYP2C23, CYP2E1, and CY4F2 rabbit polyclonal primary antibodies were purchased from Abcam (Cambridge, UK); and CYP2J rabbit polyclonal primary antibody was purchased from MilliporeSigma (St. Louis, MO). Chemiluminescence western blotting detection reagents (enhanced chemiluminescence) were obtained from Cytiva (Marlborough, MA). All other chemicals used were obtained from Sigma Aldrich (St. Louis, MO).

RNA Extraction and cDNA Synthesis. RNA extraction and cDNA synthesis were performed according to the method described by Elshenawy and El-Kadi (2015).

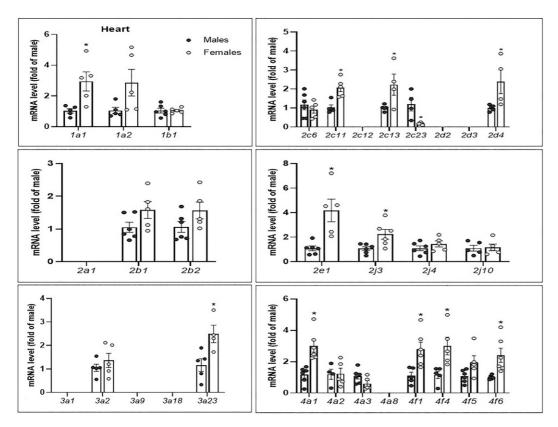


Fig. 2. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat heart. The mRNA expression of P450 enzymes was determined in the heart of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

GAPDH

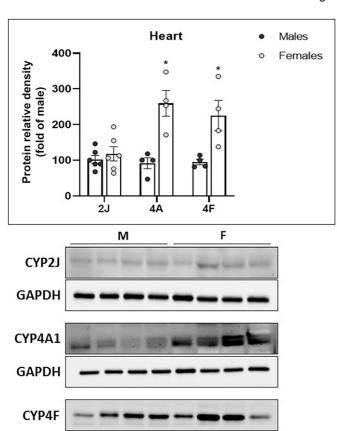


Fig. 3. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat heart. The protein expression of P450 enzymes was determined in the heart of adult male and female Sprague-Dawley rats by western blot and normalized to glyceraldehyde-3-phosphate dehydrogenase (GAPDH) house-keeping protein. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

Quantification of mRNA Expression by Real-Time PCR. The resulting cDNA was subject to PCR amplification using 384-well optical reaction plates in the QuantStudio 5 (Applied Biosystems). The 20 µL reaction mix contained 0.04 µL of 10 µM forward primers and 0.04 µL of 10 µM reverse primers (20 nM final concentration of each primer), 10 µL SYBR Green Universal Master Mix, 8.92 µL of nuclease-free water, and 1 µL cDNA sample. Thermocycling conditions were as described in previous reports (Shoieb et al., 2022). The rat P450 enzymes and their human orthologs are listed in Table 1. Rat primer sequences used in this study are listed in Table 2. Analysis of the real-time PCR data were performed using the relative gene expression ($\Delta\Delta$ CT) method (Livak and Schmittgen, 2001). In short, the fold change in the level of target genes between female and male rats, corrected for the level of the housekeeping gene, was determined using the following equation: Fold change = $2^{-\Delta(\Delta Ct)}$, where ΔCt = Ct(target gene) – Ct(housekeeping gene) and $\Delta(\Delta Ct) = \Delta Ct(females)$ – mean $\Delta Ct(males)$. For the comparison of all genes' expression within the same organ, fold change was calculated relative to the least expressed gene.

Preparation of Microsomal Protein. Microsomal fractions were prepared by differential centrifugation of the homogenized organs. Briefly, a weighed mass of each organ was homogenized in cold sucrose solution (0.25 M in distilled water, 0.5 g tissue in 2 mL sucrose solution) containing protease inhibitor cocktail (5 μ L/1 mL sucrose solution). The homogenate was centrifuged at 10,000g for 20 minutes. The resulting supernatant was centrifuged again at 100,000g for 60 minutes to obtain the microsomal pellet. The pellets were dissolved in the homogenization sucrose solution containing protease inhibitor cocktail and stored at -80° C. The Lowry method was used to determine microsomal

protein concentrations using bovine serum albumin as a standard (Lowry et al., 1951)

Western Blot Analysis. We determined the protein expression of important P450-metabolizing enzymes (CYP1A2, CYP3A) as well as some main arachidonic acid epoxygenases (CYP2C23, CYP2J) and hydroxylases (CYP2E1, CYP4A1, and CYP4F) using denaturing gel electrophoresis. Briefly, isolated proteins from the different organs of male and female rats (15 µg from the liver; 50 μg from the kidney, lung, and brain; 60 μg from the heart; and 75 μg from the small intestine) were diluted with an equal amount of 2X Laemmli Sample Buffer, boiled for 5 minute, and separated by 10% SDS-PAGE as described by Shoieb et al. (2022). Then, the blots were incubated with the primary antibody: mouse anti-rat CYP1A2 (sc-53241), rabbit anti-rat CYP2C23 (ab53944), rabbit anti-rat CYP2E1 (ab28146), rabbit anti-rat CYP2J (ABS1605), mouse anti-rat CYP3A (sc-271033), mouse anti-rat CYP4A1 (sc-53248), and rabbit anti-human CYP4F2 (ab230709) for 2 hours or overnight at 4°C. Then, blots were incubated with a horseradish peroxidase-conjugated horse anti-mouse or goat anti-rabbit IgG secondary antibody for 45 minutes at room temperature. Bands were visualized using the ChemiDocTM Imaging System (Bio-Rad Laboratories, Hercules, CA) using the enhanced chemiluminescence method.

Statistical Analysis. All results are presented as mean plus or minus S.E.M. Comparisons between male and female groups were carried out using unpaired student t tests. Differences were considered significant at P < 0.05. All statistical analyses and graphs plotting were performed using GraphPad Prism software, version 8.4.3. (GraphPad Software, Inc. La Jolla, CA).

Results

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Heart. The mRNA expression levels of different P450 enzymes were determined by real-time PCR. Some P450 enzymes were found to not be expressed in the heart (Cyp2a1, Cyp2c12, Cyp2d2, Cyp2d3, Cyp3a1, Cyp3a9, Cyp3a18, and Cyp4a8). Cyp4f5 is the most highly expressed P450 in the hearts of both male and female rats, whereas Cyp4a2 is the least expressed (Fig. 1). Generally, female hearts showed higher P450 expression levels than male hearts. Cyp2e1 showed the most marked difference, being nearly fourfold higher in female hearts than male hearts. Cyp1a1, Cyp1a2, Cyp4a1, Cyp4f1, and Cyp4f4 are all approximately threefold higher in female than male hearts. Cyp4a2, Cyp4a3, and Cyp4f5 showed no significant difference, whereas all other Cyp4 family members showed significantly higher expression in female hearts. Only Cyp2c23 was significantly higher in male hearts than female hearts (5.3-fold) (Supplemental Material). Fig. 2 shows the mRNA expression levels of different P450 enzymes in male hearts compared with female hearts.

Since mRNA expression does not always correlate with protein levels of enzymes, we measured the protein expression of certain P450 enzymes (CYP1A2, CYP2C23, CYP2E1, CYP2J, CYP3A, CYP4A1, and CYP4F) in all organs to investigate sex-specific differences at the protein level. As shown in Fig. 3, CYP2J, CYP4A, and CYP4F enzymes were detected in the heart. In agreement with the mRNA results, CYP4A and CYP4F enzymes showed higher protein levels in female rat hearts (2.8 and 2.4-fold higher than the male level, respectively). However, CYP2J showed no significant difference (Fig. 3).

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Liver. All the investigated 31 P450 enzymes were found to be expressed in the liver. The expression of *Cyp2c12* and *Cyp3a18* was found to be limited to the liver, with no extrahepatic expression. *Cyp2c23* was the highest-expressed P450 in the male liver, whereas *Cyp2e1* was the highest in the female liver. *Cyp4a8* was the lowest in both sexes (Fig. 4). In contrast to the heart, *Cyp2c11* and *Cyp2c13* levels in the liver are male specific, with dramatically higher expression in males, around 1700- and 1300-fold the female hepatic expression levels, respectively. In contrast, *Cyp2c12* in female livers is nearly 200-fold compared with male livers. In the CYP3 family, *Cyp3a9* is nearly

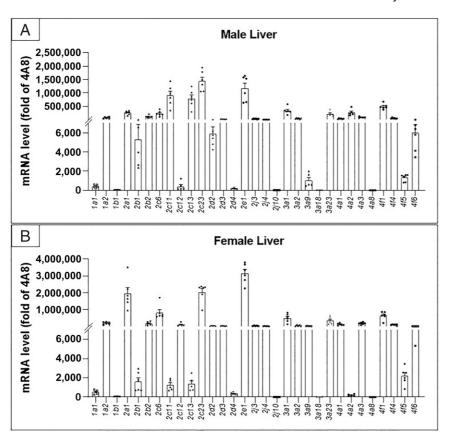


Fig. 4. The mRNA expression of different P450 enzymes in male (A) and female (B) rat liver relative to the least expressed. The mRNA expression of P450 enzymes was determined in the liver of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β-actin housekeeping gene. Results are presented as mean plus or minus S.E.M, n = 4-6.

10-fold higher in the female liver, whereas *Cyp3a18* is nearly 25-fold higher in the male liver. Most CYP4 family enzymes are significantly higher in male livers, especially *Cyp4a2*, which is also

male specific: 3000-fold in the male liver compared with the female liver (Supplemental Material). Fig. 5 shows the mRNA expression levels of different P450 enzymes in male versus female livers.

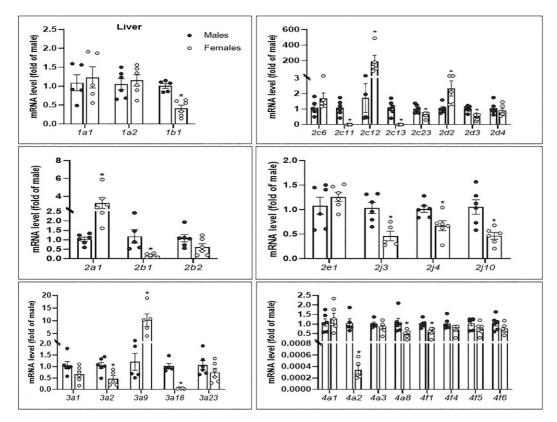


Fig. 5. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat liver. The mRNA expression of P450 enzymes was determined in the liver of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

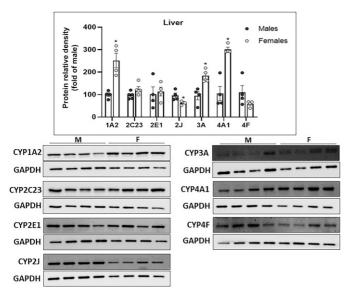


Fig. 6. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat liver. The protein expression of P450 enzymes was determined in the liver of adult male and female Sprague-Dawley rats by western blot and normalized to GAPDH housekeeping protein. Results are presented as mean plus or minus S.E.M., n = 4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

As shown in Fig. 6, all the assessed enzymes were found to be expressed in the liver. CYP1A2, CYP3A, and CYP4A1 were significantly higher in female rat livers (2.5-fold, 1.9-fold, and 2.9-fold,

respectively) compared with the male livers. In contrast, CYP2J was found to be significantly higher in the male rat liver (1.5-fold female expression level), in agreement with the mRNA result (Fig. 6).

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Kidney. Cyp2c23 was the most highly expressed, whereas Cyp3a1 was the least expressed, P450 in both male and female kidneys (Fig. 7). Cyp2c12, Cyp2c13, and Cyp3a18 are not expressed in the kidney. Several enzymes demonstrated similar sexual dimorphism in the kidney to that shown in the heart, such as Cyp1a1, Cyp2e1, Cyp4f1, Cyp4f4, and Cyp4f6, which all showed higher expression in the female organs. Cyp1a1 showed the greatest difference, with the female expression level 26-fold the male level. However, similar to the liver and in contrast to the heart, Cyp2c11 expression was significantly higher in male versus female kidneys (13-fold), and in contrast to both heart and liver, Cyp2c23 expression was significantly higher in female kidneys (approximately twofold). In addition, Cyp4a8 showed nearly 7.5-fold higher expression in the females' kidney, whereas it was not expressed in the heart (Supplemental Material). Fig. 8 shows the mRNA expression levels of different P450 enzymes in male versus female kidneys.

At the protein expression level, CYP2C23, CYP2E1, CYP3A, CYP4A1, and CYP4F2 were detected in the kidney. Similar to the mRNA expression, CYP2E1 was found to be significantly higher in the female rat kidney (1.7-fold the male expression level). CYP2C23 also appears to be slightly higher in females, but the difference did not achieve statistical significance. Moreover, CYP3A protein levels are significantly higher in the female kidney (2.4-fold), in agreement with CYP3A2 mRNA result (Fig. 9).

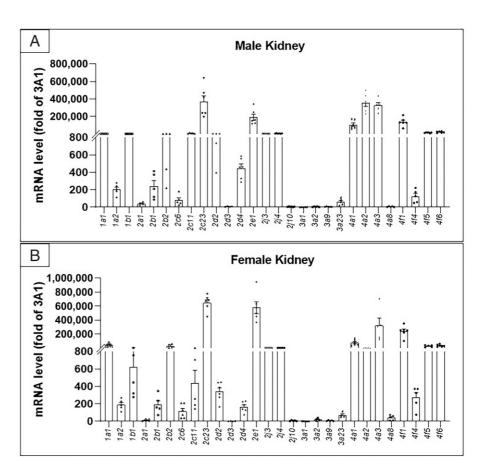


Fig. 7. The mRNA expression of different P450 enzymes in male (A) and female (B) rat kidney relative to the least expressed. The mRNA expression of P450 enzymes was determined in the kidney of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M, n=4-6.

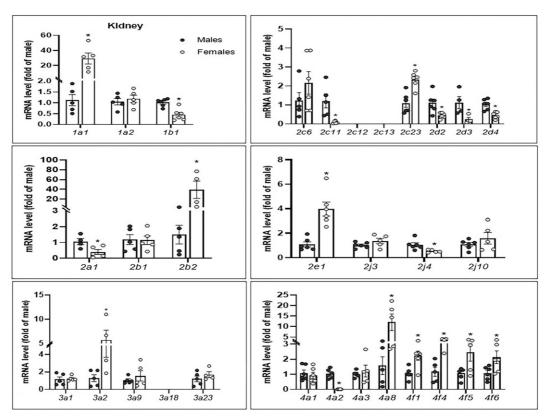


Fig. 8. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat kidney. The mRNA expression of P450 enzymes was determined in the kidney of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Lung. The most expressed P450 mRNA in the lungs of both male and female rats was *Cyp2b1*, with a marked difference between it and the second enzyme (*Cyp2e1*) (11.5-fold

in males and 27-fold in females). On the other hand, *Cyp2j10* and *Cyp3a2* were the least expressed (Fig. 10). In addition to *Cyp2c13* and *Cyp4a2* and except *Cyp3a9*, all the enzymes that were not expressed in the heart were also found to not be expressed in the lung. Except *Cyp1a2* and

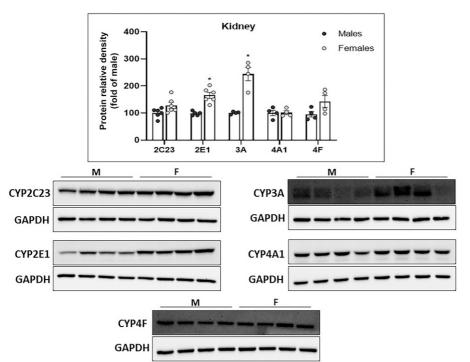


Fig. 9. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat kidney. The protein expression of P450 enzymes was determined in the kidney of adult male and female Sprague-Dawley rats by western blot and normalized to GAPDH housekeeping protein. Results are presented as mean plus or minus S.E.M., n = 4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

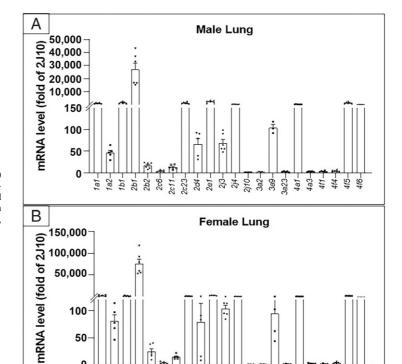


Fig. 10. The mRNA expression of different P450 enzymes in male (A) and female (B) rat lung relative to the least expressed. The mRNA expression of P450 enzymes was determined in the lung of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M, n=4-6.

Cyp2b1, all other P450 enzymes expressed in the lung did not show statistically significant sex-specific differences between males and females. Cyp1a2 is around 1.8-fold higher in male lung, whereas Cyp2b1 is

2.3-fold higher in female lung (Supplemental Material). Fig. 11 demonstrates the male versus female mRNA expression levels of different P450 enzymes in the lungs.

264 – 261 – 2/3 – 2/4 – 2/4 – 2/10 – 3/4 –

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4a1-

2c23

262-206-206-

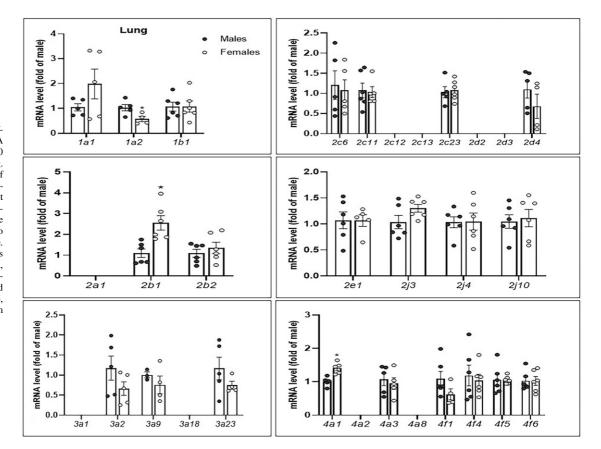


Fig. 11. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat lung. The mRNA expression of P450 enzymes was determined in the lung of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M., n = 4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

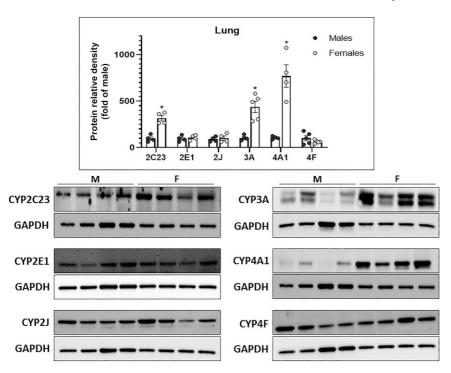


Fig. 12. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat lung. The protein expression of P450 enzymes was determined in the lung of adult male and female Sprague-Dawley rats by western blot and normalized to GAPDH housekeeping protein. Results are presented as mean plus or minus S.E.M., n = 4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

CYP2C23, CYP3A, and CYP4A1 were all found to be significantly higher in the female lung at the protein expression level (3.2-, 4.2-, and 7.3-fold the male expression level, respectively). On the other hand, CYP2E1, CYP2J, and CYP4F demonstrated no significant difference between males and females, and CYP1A2 was not detected (Fig. 12).

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Brain. Cyp2j3 was found to be the most expressed enzyme in the brain of male and female rats, whereas Cyp4a8 and Cyp2c13 were the least expressed (Fig. 13). Cyp2c12, Cyp2d2, Cyp2d3, Cyp3a18, and Cyp4a2 were found to not be expressed in the brain. Similar to the liver, the brain mRNA expression levels of Cyp2c11 and Cyp2c13 were significantly higher in males than in females but with a less marked difference (1.8- and 3.7-fold, respectively). Unlike all other organs, Cyp1a1 and Cyp4a3 in the brain were significantly higher in the males (approximately two- and sixfold, respectively). On the other hand, Cyp4f4 and Cyp4f5 mRNA levels were significantly higher in the females (1.7- and 1.4-fold, respectively) (Supplemental Material). The mRNA expression levels of different P450 enzymes in male versus female brain are shown in Fig. 14.

At the protein expression level, CYP2J and CYP4F enzymes were detected in the brain, and both showed no significant sex-specific difference (Fig. 15).

Sex-Specific Differences in the mRNA and Protein Expression Levels of P450 Enzymes in the Small Intestine. At the mRNA expression level, Cyp2b1 and Cyp2b2 were found to be the most highly expressed P450 enzymes in the small intestine of female and male rats, respectively, whereas Cyp3a1 was the least expressed in both sexes, similar to the kidney (Fig. 16). Cyp3a18, Cyp2c12, Cyp2d2, Cyp4a2, and Cyp4a3 were found to not be expressed in the small intestine. Only Cyp2c13 showed significant sex-specific difference in the small intestine, being 1.4-fold higher in the males compared with the females (Supplemental Material). The male-versus-female mRNA expression of P450 enzymes in the small intestine is demonstrated in Fig. 17.

At the protein expression level, CYP1A2 and CYP3A were demonstrated to be significantly higher in the females: 2.4-fold and 3.7-fold, respectively. This result is in agreement with other organs such as the liver, kidney, and lung. CYP4A1 was also detected at the protein

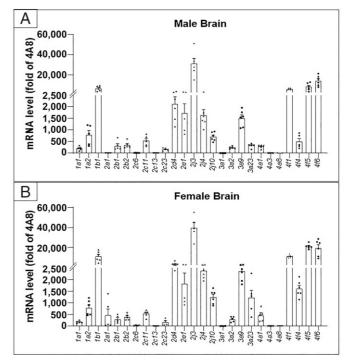


Fig. 13. The mRNA expression of different P450 enzymes in male (A) and female (B) rat brain relative to the least expressed. The mRNA expression of P450 enzymes was determined in the brain of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β -actin housekeeping gene. Results are presented as mean plus or minus S.E.M, n = 4-6.

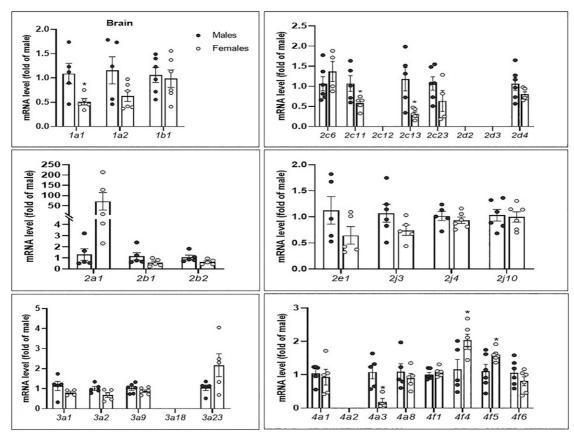


Fig. 14. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat brain. The mRNA expression of P450 enzymes was determined in the brain of adult male and female Sprague-Dawley rats by real-time PCR and normalized to β-actin housekeeping gene. Results are presented as mean plus or minus S.E.M., n = 4–6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

expression level but showed no significant difference between males and females, similar to CYP4A1 expression in the kidney (Fig. 18).

Discussion

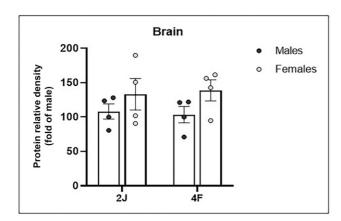
Previously, many studies were conducted on male animals only, and results were generalized to both sexes (Holdcroft, 2007; Lee, 2018). Ignoring sex differences could have undesirable consequences such as increased side effects or decreased efficacy. Thus, sex-specific differences are gaining increasing attention in research, and more studies are starting to include female groups (Wald and Wu, 2010; Lee, 2018). Previous studies have demonstrated sex-specific differences in the expression or activity levels of different P450 enzymes (Waxman and Holloway, 2009; Zhang et al., 2011). However, most of these studies reported just a few enzymes and investigated sexual dimorphism in response to inducers or inhibitors. In this study, we investigated and compared the mRNA and protein expression of different P450s in the heart, liver, lung, kidney, brain, and small intestine of male and female SD rats.

Higher expression levels of CYP1A1 were previously reported in the lungs (Lingappan et al., 2013, 2016) and hearts (Zhang et al., 2015) of female versus male mice. Moreover, CYP1A1 was detectable in the lungs and kidneys of female but not male SD rats and was undetectable in the liver of both sexes (Iba et al., 1999). Our results showed significantly higher cardiac and renal *Cyp1a1* mRNA levels in female rats. Hepatic CYP1A2 activity was reported to be higher in male than in female SD rats (Fonsart et al., 2008). However, a study found higher CYP1A2 levels in female compared with male human liver samples

(Zhang et al., 2011), whereas other studies gave conflicting results (Nafziger and Bertino, 1989; Ou-Yang et al., 2000; Zanger and Schwab, 2013). A study in 2016 demonstrated higher *Cyp1a2* brain mRNA levels in female Wistar rats (Nagai et al., 2016), whereas our results showed no significant difference in the brain. In our study, *Cyp1a2* mRNA expression showed significant sex difference only in the lung, being male dominant, whereas CYP1A2 protein was female dominant in the liver and the intestine.

CYP1B1 is constitutively expressed in several tissues, most importantly in the heart (Maayah et al., 2015). A previous study showed that treatment of embryonic rat cardiomyocytes with growth hormone (GH) in a pulsatile pattern, which mimics the male secretory pattern, significantly decreased Cyp1a1 and increased Cyp1b1 expression compared with the constant treatment pattern, which mimics the female pattern. Moreover, they found higher Cyp1b1 mRNA levels in male mice hearts compared with female mice (Zhang et al., 2015). Acute doxorubicin exposure in mice also caused a male-specific increase in cardiac Cyp1b1 (Grant et al., 2017). In our study, cardiac Cyp1b1 showed no significant difference, but hepatic and renal Cyp1b1 levels were significantly higher in male rats. Interestingly, human hepatic CYP1B1 was also previously reported to be significantly higher in men (Yang et al., 2012). Hepatic CYP2A1 was previously found to be female dominant in rats (Martignoni et al., 2006). We found higher hepatic Cyp2a1 expression in females but higher renal expression in males.

The CYP2C family is known to be highly abundant in the rat liver (Martignoni et al., 2006). CYP2C11 and CYP2C13 were previously reported to be male-specific enzymes in the liver, spleen, and bone marrow, whereas CYP2C12 was reported to be female specific (Thangavel



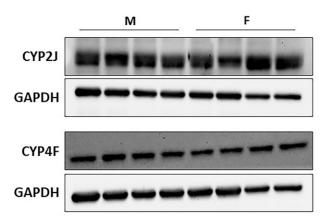


Fig. 15. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat brain. The protein expression of P450 enzymes was determined in the brain of adult male and female Sprague-Dawley rats by western blot and normalized to GAPDH housekeeping protein. Results are presented as mean plus or minus S.E.M., n = 4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

et al., 2007; Huang et al., 2011; Babelova et al., 2015). In addition, previous studies demonstrated significantly lower metabolism of CYP2C substrates in female compared with male Wistar (Ohhira et al., 2006) and SD rats (Fukuno et al., 2018). CYP2C11 sexual dimorphism is attributed to sex differences in the circulating GH profile. Female SD rats could express CYP2C11 after hypophysectomy and infusion with GH in the pulsatile male secretory fashion (Legraverend et al., 1992; Thangavel et al., 2007; Banerjee et al., 2021). We also found significantly higher *Cyp2c11* levels in male versus female kidney and brain; however, interestingly, it was found to be female dominant in the heart. CYP2C23 protein was previously reported to be highly abundant in the liver and kidney of SD rats, whereas it was undetectable in the heart and lungs (El-Sherbeni et al., 2013). We detected CYP2C23 in the lung with a higher expression in female rats.

Our results showed female-dominant expression of *Cyp2e1* in the heart and kidney. In mice, in contrast, renal CYP2E1 expression was found to be higher in males (Freeman et al., 1992; Speerschneider and Dekant, 1995), whereas cardiac CYP2E1 showed no sexual dimorphism (Zhang et al., 2015). However, acute doxorubicin exposure was associated with a female-specific increase in heart CYP2E1 in mice (Grant et al., 2017). Hepatic CYP2J2 levels were found to be significantly higher in female subjects compared with male subjects (Yang et al., 2012). In contrast, our results showed male-dominant hepatic expression of CYP2J enzymes. We found *Cyp2j4* to be significantly higher in male kidneys. A previous study demonstrated significantly higher CYP2J5

levels in male versus female mice kidneys (Ma et al., 2004). As for cardiac expression, we found significantly higher Cyp2j3 levels in female rat hearts compared with males. In line with that, treatment of rat cardiomyocytes with GH in a male secretory pattern significantly decreased Cyp2j3 expression compared with the female pattern. Mouse Cyp2j11 is also higher in female than in male hearts (Zhang et al., 2015). However, cardiac CYP2J protein levels showed no significant difference.

The CYP3A subfamily of enzymes is considered the most important among human drug-metabolizing enzymes. Several studies have shown that women have significantly higher hepatic and intestinal CYP3A enzyme activities compared with men (Tanaka, 1999; Greenblatt and Von Moltke, 2008; Krogstad et al., 2020), as well as higher hepatic CYP3A4 levels (Lamba et al., 2010; Yang et al., 2010, 2012). CYP3A7, the human ortholog of rat Cyp3a9, was previously found to have significantly higher gene expression in female hepatic samples compared with males (Yang et al., 2012). In agreement with human data, our results showed higher hepatic *Cyp3a9* mRNA and CYP3A protein levels in female rats; however, we found *Cyp3a2* mRNA to be higher in male rats. CYP3A2 was previously found to be induced by zolmitriptan in male but not female SD rats (Yu et al., 2008).

A study in 2003 reported *Cyp3a9* mRNA levels to be significantly higher in the livers and lungs of female SD rats compared with male rats and that its expression is affected by ovariectomy and subsequent estrogen administration (Anakk et al., 2003). In agreement, we found significantly higher *Cyp3a9* mRNA and CYP3A protein levels in the livers of female rats. In the lung, kidney, and small intestine, *Cyp3a9* mRNA levels showed no significant difference, whereas CYP3A protein levels were significantly higher in the females. Similar to our findings, previous reports have identified hepatic CYP3A2 and CYP3A18 to be male-dominant isoenzymes and CYP3A9 to be a female-dominant isoenzyme in Wistar (Robertson et al., 1998) and SD rats (Kushida et al., 2021).

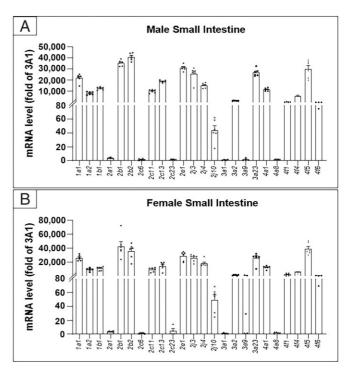


Fig. 16. The mRNA expression of different P450 enzymes in male (A) and female (B) rat small intestine relative to the least expressed. The mRNA expression of P450 enzymes was determined in the small intestine of adult male and female Sprague-Dawley rats by real-time PCR and normalized to Gapdh house-keeping gene. Results are presented as mean plus or minus S.E.M, n=4-6.

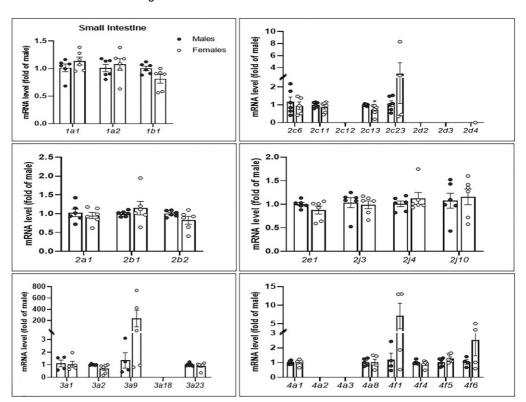


Fig. 17. Sex-specific differences in the mRNA expression levels of P450 enzymes in the rat small intestine. The mRNA expression of P450 enzymes was determined in the small intestine of adult male and female Sprague-Dawley rats by real-time PCR and normalized to Gapdh housekeeping gene. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

CYP4A enzymes play an important role in the ω -hydroxylation of AA (El-Sherbeni and El-Kadi, 2017). In line with our results, hepatic Cyp4a2 was previously reported to be significantly higher in male rats of Fischer 344 and obese ZSF1 strains (Sundseth and Waxman, 1992; Babelova et al., 2015). Moreover, the induction of hepatic Cyp4a by endotoxin was found to be male specific in SD and Fischer 344 rats (Mitchell et al., 2001). Although hepatic Cyp4a1 showed no sex-specific difference at the mRNA level, hepatic Cyp4a1 protein levels were found to be significantly higher in female rats.

We found renal *Cyp4a2* levels to be significantly higher in male rats. Similarly, previous studies showed significantly higher renal CYP4A2 levels in male versus female Fischer 344 and SD rats (Sundseth and Waxman, 1992; Bleicher et al., 2001). Interestingly, a previous study demonstrated that treatment of SD rats with dihydrotestosterone lowered the renal *Cyp4a1* levels and enhanced *Cyp4a2/3* levels (Nakagawa et al., 2003). In addition, clofibrate administration significantly enhanced renal *Cyp4a2* expression only in male SD rats (Bleicher et al., 2001). Another study showed an increase of renal CYP4A protein in female SD rats treated with dihydrotestosterone (Zhou et al., 2005).

CYP4F enzymes appear to be female-dominant enzymes. We found significantly higher *Cyp4f* levels in the heart and kidney of females versus males, and *Cyp4f*4 and *Cyp4f*5 were significantly higher in female brains. At the protein expression level, we found significantly higher CYP4F levels in the heart of female versus male rats. Similarly, a study in 2002 found significantly higher expression levels of CYP4F enzymes in female versus male SD rats in the liver, kidney, lung, and brain and found a significant decrease in hepatic and renal CYP4F expression levels in female rats after ovariectomy, which was significantly restored by estrogen treatment (Kalsotra et al., 2002).

In conclusion, there are significant sex-specific differences in the expression levels of different P450 enzymes. Elucidating sex-specific differences in P450s is crucial for explaining the differences between males and females in diseases processes and treatment outcomes. This study has some limitations. First, species discrepancies in the basal and

inducible levels of enzymes could complicate the translation of the results to humans (Hammer et al., 2021). Moreover, P450 expression levels could differ among different rat strains (Nishiyama et al., 2016).

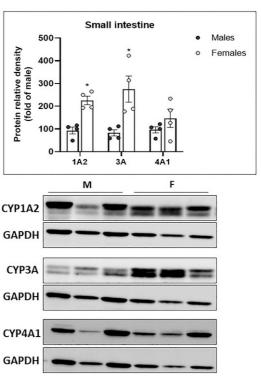


Fig. 18. Sex-specific differences in the protein expression levels of some P450 enzymes in the rat small intestine. The protein expression of P450 enzymes was determined in the small intestine of adult male and female Sprague-Dawley rats by western blot and normalized to GAPDH housekeeping protein. Results are presented as mean plus or minus S.E.M., n=4-6. Data were analyzed using an unpaired student t test. *P < 0.05, significant difference from male rats.

Finally, expression levels of enzymes are not necessarily correlated to their activity levels. However, rats are still considered to be valuable models for preclinical development and have previously been used for the study of sex differences in drug-metabolizing enzymes and the mechanisms underlying these differences (Waxman and Holloway, 2009; Jung et al., 2015; Blais et al., 2017). Moreover, several rat P450 enzymes (e.g., CYP1A1, CYP1A2, CYP2E1, CYP2J3, CYP4F1) show high degrees of structural similarity to their human orthologs (Hammer et al., 2021). Thus, despite the study limitations and some results that are different from human data, we believe that our results could still give valuable insights regarding sex-specific differences in human P450 enzymes. Additional studies investigating the activity of different P450 enzymes and levels of their metabolites in males and females are important for having better insight into sex-specific discrepancies and their potential clinical and therapeutic implications.

Authorship Contributions

Participated in research design: Gerges, El-Kadi.

Conducted experiments: Gerges.

Performed data analysis: Gerges, El-Kadi.

Wrote or contributed to the writing of the manuscript: Gerges, El-Kadi.

References

- Alonso, S, Su M, Jones JW, Ganguly S, Kane MA, Jones RJ, and Ghiaur G (2015) Human bone marrow niche chemoprotection mediated by cytochrome P450 enzymes. Oncotarget 6:14905–14912.
- Anakk S, Ku CY, Vore M, and Strobel HW (2003) Insights into gender bias: rat cytochrome P450 3A9. *J Pharmacol Exp Ther* **305**:703–709.
- Babelova A, Burckhardt BC, Salinas-Riester G, Pommerenke C, Burckhardt G, and Henjakovic M (2015) Next generation sequencing of sex-specific genes in the livers of obese ZSF1 rats. Genomics 106:204-213.
- Banerjee S, Hayes AM, and Shapiro BH (2021) Early expression of requisite developmental growth hormone imprinted cytochromes P450 and dependent transcription factors. *Endocr Connect* 10:1167–1179.
- Blais EM, Rawls KD, Dougherty BV, Li ZI, Kolling GL, Ye P, Wallqvist A, and Papin JA (2017) Reconciled rat and human metabolic networks for comparative toxicogenomics and biomarker predictions. *Nat Commun* 8:14250.
- Bleicher KB, Pippert TR, Glaab WE, Skopek TR, Sina JF, and Umbenhauer DR (2001) Use of real-time gene-specific polymerase chain reaction to measure RNA expression of three family members of rat cytochrome P450 4A. J. Biochem Mol Toxicol. 15:133–142.
- members of rat cytochrome P450 4A. *J Biochem Mol Toxicol* **15**:133–142.

 Dahan A, Kest B, Waxman AR, and Sarton E (2008) Sex-specific responses to opiates: animal and human studies. *Anesth Analg* **107**:83–95.
- Dutheil F, Beaune P, and Loriot MA (2008) Xenobiotic metabolizing enzymes in the central nervous system: Contribution of cytochrome P450 enzymes in normal and pathological human brain. *Biochimie* **90**:426–436.
- El-Sherbeni AA, Aboutabl ME, Zordoky BNM, Anwar-Mohamed A, and El-Kadi AOS (2013) Determination of the dominant arachidonic acid cytochrome p450 monooxygenases in rat heart, lung, kidney, and liver: protein expression and metabolite kinetics. *AAPS J* 15:112–122.
- El-Sherbeni AA and El-Kadi AOS (2017) Microsomal cytochrome P450 as a target for drug discovery and repurposing. *Drug Metab Rev* 49:1–17.
- Elbekai RH and El-Kadi AOS (2006) Cytochrome P450 enzymes: central players in cardiovascular health and disease. *Pharmacol Ther* **112**:564–587.
- Elshenawy OH and El-Kadi AOS (2015) Modulation of aryl hydrocarbon receptor regulated genes by acute administration of trimethylarsine oxide in the lung, kidney and heart of C57BL/6 mice. *Xenobiotica* **45**:930–943.
- Fan F, Muroya Y, and Roman RJ (2015) Cytochrome P450 eicosanoids in hypertension and renal disease. Curr Opin Nephrol Hypertens 24:37–46.
- Fonsart J, Menet MC, Declèves X, Galons H, Crété D, Debray M, Scherrmann JM, and Noble F (2008) Sprague-Dawley rats display metabolism-mediated sex differences in the acute toxicity of 3,4-methylenedioxymethamphetamine (MDMA, ecstasy). *Toxicol Appl Pharmacol* 230: 117–125.
- Franconi F, Brunelleschi S, Steardo L, and Cuomo V (2007) Gender differences in drug responses *Pharmacol Res* 55:81–95.
- Freeman JE, Stirling D, Russell AL, and Wolf CR (1992) cDNA sequence, deduced amino acid sequence, predicted gene structure and chemical regulation of mouse Cyp2e1. *Biochem J* 281:689–695.
- Fukuno S, Nagai K, Horii A, Yamamoto K, and Konishi H (2018) Pharmacokinetics and metabolic elimination of tolbutamide in female rats: Comparison with male rats. *Biopharm Drug Dispos* 39:321–327.
- Gerges SH and El-Kadi AOS (2022) Sex differences in eicosanoid formation and metabolism: A possible mediator of sex discrepancies in cardiovascular diseases. *Pharmacol Ther* 234:108046.
 Grant MKO, Seelig DM, Sharkey LC, and Zordoky BN (2017) Sex-dependent alteration of cardiac
- cytochrome P450 gene expression by doxorubicin in C57Bl/6 mice. *Biol Sex Differ* **8**:1–13. Greenblatt DJ and von Moltke LL (2008) Gender has a small but statistically significant effect on clearance of CYP3A substrate drugs. *J Clin Pharmacol* **48**:1350–1355.
- Hammer H, Schmidt F, Marx-Stoelting P, Pötz O, and Braeuning A (2021) Cross-species analysis of hepatic cytochrome P450 and transport protein expression. Arch Toxicol 95:117–133.
- Holdcroft A (2007) Gender bias in research: how does it affect evidence based medicine? J R Society 100:2–3.

- Holingue C, Budavari AC, Rodriguez KM, Zisman CR, Windheim G, and Fallin MD (2020) Sex Differences in the Gut-Brain Axis: Implications for Mental Health. *Curr Psychiatry Rep* 22:83.
- Huang HJ, Tsai ML, Chen YW, and Chen SH (2011) Quantitative shot-gun proteomics and MS-based activity assay for revealing gender differences in enzyme contents for rat liver microsome. J Proteomics 74:2734–2744.
- Hukkanen J, Pelkonen O, Hakkola J, and Raunio H (2002) Expression and regulation of xenobiotic-metabolizing cytochrome P450 (CYP) enzymes in human lung. Crit Rev Toxicol 32: 391–411
- Iba MM, Fung J, Thomas PE, and Park Y (1999) Constitutive and induced expression by pyridine and β-naphthoflavone of rat CYP1A is sexually dimorphic. Arch Toxicol 73:208–216.
- Ibrahim M, MacFarlane EM, Matteo G, Hoyeck MP, Rick KRC, Farokhi S, Copley CM, O'Dwyer S, and Bruin JE (2020) Functional cytochrome P450 1A enzymes are induced in mouse and human islets following pollutant exposure. *Diabetologia* 63:162–178.
- Jung JW, Choi MR, Kwon YS, Jeong JS, Son M, and Kang HE (2015) Gender differences in corydaline pharmacokinetics in rats. Xenobiotica 45:456–463.
- Kalsotra A, Anakk S, Boehme CL, and Strobel HW (2002) Sexual dimorphism and tissue specificity in the expression of CYP4F forms in Sprague Dawley rats. *Drug Metab Dispos* 30: 1022–1028.
- Krogstad V, Peric A, Robertsen I, Kringen MK, Wegler C, Angeles PC, Hjelmesæth J, Karlsson C, Andersson S, Artursson P, et al. (2020) A Comparative Analysis of Cytochrome P450 Activities in Paired Liver and Small Intestinal Samples from Patients with Obesity. *Drug Metab Dispos* 48:8–17.
- Kushida H, Matsumoto T, Ikarashi Y, Nishimura H, and Yamamoto M (2021) Gender differences in plasma pharmacokinetics and hepatic metabolism of geissoschizine methyl ether from Uncaria hook in rats. J Ethnopharmacol 264:113354.
- Lamba V, Panetta JC, Strom S, and Schuetz EG (2010) Genetic predictors of interindividual variability in hepatic CYP3A4 expression. J Pharmacol Exp Ther 332:1088–1099.
- Lee SK (2018) Sex as an important biological variable in biomedical research. *BMB Rep* **51**:167–173.
- Legraverend C, Mode A, Westin S, Ström A, Eguchi H, Zaphiropoulos PG, and Gustafsson JÅ (1992) Transcriptional regulation of rat P-450 2C gene subfamily members by the sexually dimorphic pattern of growth hormone secretion. *Mol Endocrinol* 6:259–266.
- Lingappan K, Jiang W, Wang L, Couroucli XI, Barrios R, and Moorthy B (2013) Sex-specific differences in hyperoxic lung injury in mice: implications for acute and chronic lung disease in humans. Toxicol Appl Pharmacol 272:281–290.
- Lingappan K, Jiang W, Wang L, and Moorthy B (2016) Sex-specific differences in neonatal hyperoxic lung injury. Am J Physiol Lung Cell Mol Physiol 311:L481–L493.
- Livak KJ and Schmittgen TD (2001) Analysis of relative gene expression data using real-time quantitative PCR and the 2(-Δ Δ C(T)) Method. Methods 25:402–408.
- Lowry OH, Rosebrough NJ, Farr, Al, and Randall RJ (1951) Protein measurement with the Folin phenol reagent. J Biol Chem 193:265–275.
- Lynch T and Price A (2007) The effect of cytochrome P450 metabolism on drug response, interactions, and adverse effects. Am Fam Physician 76:391–396.
- Ma J, Graves J, Bradbury JA, Zhao Y, Swope DL, King L, Qu W, Clark J, Myers P, Walker V, et al. (2004) Regulation of mouse renal CYP2J5 expression by sex hormones. *Mol Pharmacol* 65:730–743.
- Maayah ZH, Elshenawy OH, Althurwi HN, Abdelhamid G, and El-Kadi AOS (2015) Human fetal ventricular cardiomyocyte, RL-14 cell line, is a promising model to study drug metabolizing enzymes and their associated arachidonic acid metabolites. J Pharmacol Toxicol Methods 71: 33-41.
- Madla CM, Gavins FKH, Merchant HA, Orlu M, Murdan S, and Basit AW (2021) Let's talk about sex: Differences in drug therapy in males and females. *Adv Drug Deliv Rev* 175:113804.
- Manikandan P and Nagini S (2018) Cytochrome P450 Structure, Function and Clinical Significance: A Review. Curr Drug Targets 19:38–54.
- Martignoni M, Groothuis GMM, and de Kanter R (2006) Species differences between mouse, rat, dog, monkey and human CYP-mediated drug metabolism, inhibition and induction. Expert Opin Drug Metab Toxicol 2:875–894.
- Mitchell SR, Sewer MB, Kardar SS, and Morgan ET (2001) Characterization of CYP4A induction in rat liver by inflammatory stimuli: dependence on sex, strain, and inflammation-evoked hypophagia. *Drug Metab Dispos* 29:17–22.
- Nafziger AN and Bertino Jr JS (1989) Sex-related differences in theophylline pharmacokinetics. Eur J Clin Pharmacol 37:97–100.
- Nagai K, Fukuno S, Suzuki H, and Konishi H (2016) Higher gene expression of CYP1A2, 2B1 and 2D2 in the brain of female compared with male rats. *Pharmazie* 71:334–336.
- Nakagawa K, Marji JS, Schwartzman ML, Waterman MR, and Capdevila JH (2003) Androgenmediated induction of the kidney arachidonate hydroxylases is associated with the development of hypertension. Am J Physiol Regul Integr Comp Physiol 284:R1055–R1062.
- Nebert DW, Adesnik M, Coon MJ, Estabrook RW, Gonzalez FJ, Guengerich FP, Gunsalus IC, Johnson EF, Kemper B, Levin W, et al. (1987) The P450 gene superfamily: recommended nomenclature. DNA 6:1-11.
- Ngo ST, Steyn FJ, and McCombe PA (2014) Gender differences in autoimmune disease. Front Neuroendocrinol 35:347–369.
- Nishiyama Y, Nakayama SMM, Watanabe KP, Kawai YK, Ohno M, Ikenaka Y, and Ishizuka M (2016) Strain differences in cytochrome P450 mRNA and protein expression, and enzymatic activity among Sprague Dawley, Wistar, Brown Norway and Dark Agouti rats. J Vet Med Sci 78:675–680.
- Ohhira Š, Enomoto M, and Matsui H (2006) Sex difference in the principal cytochrome P-450 for tributyltin metabolism in rats. *Toxicol Appl Pharmacol* 210:32–38.
- Ou-Yang DS, Huang SL, Wang W, Xie HG, Xu ZH, Shu Y, and Zhou HH (2000) Phenotypic polymorphism and gender-related differences of CYP1A2 activity in a Chinese population. Br J Clin Pharmacol 49:145–151.
- Regitz-Zagrosek V and Kararigas G (2017) Mechanistic pathways of sex differences in cardiovascular disease. *Physiol Rev* 97:1–37.
- Robertson GR, Farrell GC, and Liddle C (1998) Sexually dimorphic expression of rat CYP3A9 and CYP3A18 genes is regulated by growth hormone. *Biochem Biophys Res Commun* **242**:57–60.
- CYPACI8 genes is regulated by growtn normone. *Biochem Biophys Res Commun* 242:57–60. Schwartz JB (2007) The current state of knowledge on age, sex, and their interactions on clinical pharmacology. *Clin Pharmacol Ther* 82:87–96.
- Shoieb SM, Alammari AH, Levasseur J, Silver H, Dyck JRB, and El-kadi AOS (2022) Ameliorative Role of Fluconazole Against Abdominal Aortic Constriction Induced Cardiac Hypertrophy in Rats. J Cardiovasc Pharmacol 79:833–845.

- Shoieb SM, El-Sherbeni AA, and El-Kadi AOS (2019) Subterminal hydroxyeicosatetraenoic acids: crucial lipid mediators in normal physiology and disease states. Chem Biol Interact 299:140-150.
- Speerschneider P and Dekant W (1995) Renal tumorigenicity of 1,1-dichloroethene in mice: the role of male-specific expression of cytochrome P450 2E1 in the renal bioactivation of 1, 1-dichloroethene. *Toxicol Appl Pharmacol* 130:48–56.
- Sundseth SS, and Waxman DJ (1992) Sex-dependent expression and clofibrate inducibility of cytochrome P450 4A fatty acid ω-hydroxylases. Male specificity of liver and kidney CYP4A2 mRNA and tissue-specific regulation by growth hormone and testosterone. *J Biol Chem* 267:3915–3921.
- Tanaka E (1999) Gender-related differences in pharmacokinetics and their clinical significance. J Clin Pharm Ther 24:339–346.
- Thangavel C, Dhir RN, Volgin DV, and Shapiro BH (2007) Sex-dependent expression of CYP2C11 in spleen, thymus and bone marrow regulated by growth hormone. *Biochem Pharmacol* 74:1476–1484.
- Tramunt B, Smati S, Grandgeorge N, Lenfant F, Amal JF, Montagner A, and Gourdy P (2020) Sex differences in metabolic regulation and diabetes susceptibility. *Diabetologia* **63**:453–461.
- Wald C, and Wu C (2010) Biomedical research. Of mice and women: the bias in animal models. Science 327:1571–1572. DOI: 10.1126/science.327.5973.1571.
- Waxman DJ and Holloway MG (2009) Sex differences in the expression of hepatic drug metabolizing enzymes. *Mol Pharmacol* **76**:215–228.
- Wilkinson GR (2005) Drug metabolism and variability among patients in drug response. N Engl J Med 352:2211–2221.
- Yang L, Li Y, Hong H, Chang C-W, Guo L-W, Lyn-Cook B, Shi L, and Ning B (2012) Sex differences in the expression of drug-metabolizing and transporter genes in human liver. J Drug Metab Toxicol 3:1000119.

- Yang X, Zhang B, Molony C, Chudin E, Hao K, Zhu J, Gaedigk A, Suver C, Zhong H, Leeder JS, et al. (2010) Systematic genetic and genomic analysis of cytochrome P450 enzyme activities in human liver. Genome Res 20:1020–1036.
- Yu L, Lu S, Zhao N, Ni S, Yao T, and Zeng S (2008) Male-specific induction of CYP3A2 in rats by zolmitriptan. J Pharm Pharmacol 60:1601–1607.
- Zanger UM and Schwab M (2013) Cytochrome P450 enzymes in drug metabolism: regulation of gene expression, enzyme activities, and impact of genetic variation. *Pharmacol Ther* 138:103–141.Zhang F, Yu X, He C, Ouyang X, Wu J, Li J, Zhang J, Duan X, Wan Y, and Yue J (2015) Effects of sexually dimorphic growth hormone secretory patterns on arachidonic acid metabolizing enzymes in rodent heart. *Toxicol Appl Pharmacol* 289:495–506.
- Zhang Y, Klein K, Sugathan A, Nassery N, Dombkowski A, Zanger UM, and Waxman DJ (2011) Transcriptional profiling of human liver identifies sex-biased genes associated with polygenic dyslipidemia and coronary artery disease. *PLoS One* 6:e23506.
- Zhou Y, Lin S, Chang H-H, Du J, Dong Z, Dorrance AM, Brands MW, and Wang M-H (2005) Gender differences of renal CYP-derived eicosanoid synthesis in rats fed a high-fat diet. Am J Hypertens 18:530–537.
- Zhu Y and Zhang QY (2012) Role of intestinal cytochrome p450 enzymes in diclofenac-induced toxicity in the small intestine. J Pharmacol Exp Ther 343:362–370.

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Sexual Dimorphism in the Expression of Cytochrome P450 Enzymes in Rat Heart, Liver, Kidney, Lung, Brain, and Small Intestine

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Supplemental information: Supplemental tables.

Supplemental Material:

S.1 Calculation of the fold change in the heart level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta CT$ method:

Table S.1.1: *Cyp1a1* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp1a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	29.13337135	20.92013931	8.213232	-0.35456	1.278592
Male	29.66686821	20.3425827	9.324286	0.756497	0.591932
IVIGIC	29.91246796	21.15733528	8.755133	0.187344	0.878221
	28.86815262	20.42703056	8.441122	-0.12667	1.091768
	28.72581673	20.62064743	8.105169	-0.46262	1.378041
		Mean ΔCt (control)	8.567788		
			Fold gene	expression	1.043711
			Standa	rd deviation	0.316523
		Sta	ndard error	of the mean	0.141553
		Relative	e standard er	ror of mean	13.5625
	28.59449959	22.33119392	6.263306	-2.30448	4.939903
	29.1534977	20.97496605	8.178532	-0.38926	1.309718
Female					
remaie	28.30037117	21.40462685	6.895744	-1.67204	3.186658
	28.07941055	20.5268631	7.552547	-1.01524	2.02124
	26.61247826	19.77409935	6.838379	-1.72941	3.31592
	Fold gene expression				
	Standard deviation				
	Standard error of the mean				
		Relative	e standard er	ror of mean	21.00355

Table S.1.2: Cyp1a2 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	30.19035339	21.05346298	9.13689	-0.85739	1.811758	
0.0-1-	30.82837486	20.3425827	10.48579	0.491512	0.711279	
Male	31.35821342	21.15733528	10.20088	0.206598	0.866579	
	30.26600075	20.42703056	9.83897	-0.15531	1.113661	
	30.92951965	20.62064743	10.30887	0.314592	0.804079	
	Mean ΔCt (control) 9.994281					
			Fold gene	expression	1.061471	
			Standar	d deviation	0.445119	
		Sta	ndard error o	f the mean	0.199063	
		Relative	standard err	or of mean	18.75351	
	29.95394897	22.33119392	7.622755	-2.37153	5.174881	
	30.75047493	20.97496605	9.775509	-0.21877	1.163742	
Female						
remaie	30.23438263	21.40462685	8.829756	-1.16452	2.241594	
	28.30106926	20.5268631	7.774206	-2.22007	4.659175	
	29.61742592	19.77409935	9.843327	-0.15095	1.110303	
Fold gene expression					2.869939	
Standard deviation					1.931085	
	Standard error of the mean				0.863608	
	Relative standard error of mean					

Table S.1.3: Cyp1b1 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1b1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	26.03538895	22.85683632	3.178553	0.297267	0.813792	
	27.12061119	24.93114853	2.189463	-0.69182	1.615323	
na-l-	27.05721092	24.19585228	2.861359	-0.01993	1.013908	
Male	27.26329994	24.80513954	2.45816	-0.42312	1.340829	
	26.72787857	23.04715919	3.680719	0.799434	0.574575	
	26.9718399	24.05238152	2.919458	0.038173	0.973887	
	Mean ΔCt (control) 2.881285					
			Fold gene	expression	1.055386	
			Standard	d deviation	0.372268	
		Sta	ndard error o	f the mean	0.151978	
		Relative	standard err	or of mean	14.4002	
	27.11470032	24.01935005	3.09535	0.214065	0.862105	
	26.84157181	24.15325165	2.68832	-0.19297	1.143111	
Famala						
Female	25.97097778	23.04361534	2.927362	0.046077	0.968566	
	25.44727898	22.94629478	2.500984	-0.3803	1.301614	
	26.67259979	23.86953545	2.803064	-0.07822	1.055715	
Fold gene expression					1.066222	
Standard deviation				0.167803		
Standard error of the mean				0.075044		
Relative standard error of mean					7.038304	

Table S.1.4: Cyp2b1 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2b1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	28.87459373	19.83474731	9.039846	-0.59072	1.506003
	29.72143173	20.69773674	9.023695	-0.60688	1.522958
Mala	30.55892372	20.32675362	10.23217	0.601599	0.659023
Male	30.96835899	21.10294151	9.865417	0.234846	0.849775
	28.85180855	19.21535873	9.63645	0.005879	0.995933
	30.71287537	20.72702789	9.985847	0.355276	0.78172
Mean ΔCt (control) 9.630571					
			Fold gene	expression	1.052569
			Standa	rd deviation	0.374016
		St	andard error	of the mean	0.152691
		Relativ	e standard er	ror of mean	14.50653
	29.4179821	21.04896927	8.369013	-1.26156	2.397546
	29.8570919	20.39573288	9.461359	-0.16921	1.124444
Female					
remaie	28.93009186	19.21967506	9.710417	0.079846	0.946159
	28.53790855	19.59550095	8.942408	-0.68816	1.611231
	29.09913063	20.33984947	8.759281	-0.87129	1.829298
Fold gene expression					1.581735
Standard deviation				0.578959	
Standard error of the mean				0.258919	
Relative standard error of mean					16.36927

Table S.1.5: Cyp2b2 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2b2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	28.32958603	19.83474731	8.494839	-0.75395	1.686407
	29.66993904	20.69773674	8.972202	-0.27659	1.211328
Mala	29.96926117	20.32675362	9.642508	0.393716	0.761167
Male	29.95347214	21.10294151	8.850531	-0.39826	1.317918
	28.94464874	19.21535873	9.72929	0.480498	0.71673
	30.53040886	20.72702789	9.803381	0.554589	0.680851
Mean ΔCt (control) 9.248792					
			Fold gen	e expression	1.0624
			Standa	ard deviation	0.408084
		St	andard error	of the mean	0.1666
		Relativ	e standard e	rror of mean	15.68144
	29.01791382	21.04896927	7.968945	-1.27985	2.428132
	29.2552166	20.39573288	8.859484	-0.38931	1.309765
Female					
remaie	28.43643379	19.21967506	9.216759	-0.03203	1.022452
	28.55935287	19.59550095	8.963852	-0.28494	1.218359
	28.69308472	20.33984947	8.353235	-0.89556	1.860327
	Fold gene expression				
	Standard deviation				0.572678
	Standard error of the mean				0.256109
		Relativ	e standard e	rror of mean	16.33551

Table S.1.6: Cyp2c6 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	33.16339493	21.7662735	11.39712	-1.01135	2.015803	
	33.71570969	22.06328773	11.65242	-0.75605	1.688865	
na-l-	35.91555405	23.92972374	11.98583	-0.42265	1.340384	
Male	35.42041779	21.89413071	13.52629	1.117811	0.460792	
	34.54964447	21.03207779	13.51757	1.10909	0.463586	
	34.05702209	21.68539238	12.37163	-0.03685	1.025869	
	Mean ΔCt (control) 12.40848					
			Fold gene	expression	1.165883	
			Standar	d deviation	0.63817	
		Star	ndard error o	f the mean	0.260532	
		Relative	standard err	or of mean	22.34629	
	34.42850876	22.53539467	11.89311	-0.51536	1.429353	
	35.80165863	22.48576355	13.3159	0.907419	0.533138	
Familia						
Female	35.7972374	22.18027306	13.61696	1.208488	0.432722	
	34.45858002	22.10305214	12.35553	-0.05295	1.037383	
	34.17071915	21.98749924	12.18322	-0.22526	1.168985	
	Fold gene expression					
	Standard deviation				0.424945	
	Standard error of the mean				0.190041	
	Relative standard error of mean					

Table S.1.7: Cyp2c11 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c11</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	29.3860054	21.7662735	7.619732	-0.6333	1.551104	
	30.49220085	22.06328773	8.428913	0.175886	0.885224	
Male						
iviale	30.49056244	21.89413071	8.596432	0.343405	0.788179	
	29.34132957	21.03207779	8.309252	0.056225	0.961778	
	29.99619865	21.68539238	8.310806	0.057779	0.960742	
	Mean ΔCt (control) 8.253027					
			Fold gene	expression	1.029405	
			Standa	rd deviation	0.300182	
		Sta	ndard error	of the mean	0.134245	
		Relative	standard er	ror of mean	13.04105	
	30.12242508	22.53539467	7.58703	-0.666	1.586664	
	29.54848671	22.48576355	7.062723	-1.1903	2.282008	
Female						
remaie	29.34716606	22.18027306	7.166893	-1.08613	2.123044	
	28.89202118	22.10305214	6.788969	-1.46406	2.758833	
	29.53164482	21.98749924	7.544146	-0.70888	1.634536	
	Fold gene expression					
	Standard deviation				0.486119	
	Standard error of the mean					
	Relative standard error of mean					

Table S.1.8: Cyp2c13 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c13</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	31.44035339	21.7662735	9.67408	-0.03493	1.024509	
	31.48086929	22.06328773	9.417582	-0.29143	1.223854	
Male						
iviale	31.56737709	21.89413071	9.673246	-0.03577	1.025101	
	30.75460434	21.03207779	9.722527	0.013514	0.990677	
	31.74302292	21.68539238	10.05763	0.348618	0.785336	
	Mean ΔCt (control) 9.709013					
			Fold gene	expression	1.009896	
			Standard	d deviation	0.15574	
		Stan	dard error o	f the mean	0.069649	
		Relative s	standard err	or of mean	6.896638	
	32.34845352	22.53539467	9.813059	0.104046	0.93042	
	30.33805656	22.48576355	7.852293	-1.85672	3.621833	
Female						
remale						
	30.55036163	22.10305214	8.447309	-1.2617	2.397787	
	30.70465279	21.98749924	8.717154	-0.99186	1.988747	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
		Relative s	standard err	or of mean	24.88953	

Table S.1.9: Cyp2c23 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	29.78787422	24.93114853	4.856726	-0.55644	1.470633
Mole	28.60475159	24.19585228	4.408899	-1.00426	2.00592
Male	30.24746704	24.80513954	5.442327	0.029164	0.979988
	29.99185944	23.04715919	6.9447	1.531537	0.345909
	Mean ΔCt (control) 5.413163				
			Fold gene	expression	1.200612
			Standa	rd deviation	0.707254
		Star	ndard error	of the mean	0.353627
		Relative	standard er	ror of mean	29.45389
	31.64385796	24.01935005	7.624508	2.211345	0.215933
	31.83003235	24.15325165	7.676781	2.263618	0.208249
	32.59623718	25.18678856	7.409449	1.996285	0.250645
Female	30.94094086	23.04361534	7.897326	2.484162	0.178728
	31.66618919	22.94629478	8.719894	3.306731	0.101059
	31.81600189	23.86953545	7.946466	2.533303	0.172743
Fold gene expression					0.187893
	Standard deviation				
	Standard error of the mean				
Relative standard error of mean					11.07562

Table S.1.10: *Cyp2d4* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp2d4)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	31.50374603	20.44839668	11.05535	-0.07904	1.056314	
Male	31.64796257	20.65575409	10.99221	-0.14218	1.103572	
iviale	32.53544617	21.05667305	11.47877	0.344384	0.787644	
	31.02033997	20.1381073	10.88223	-0.25216	1.190986	
	31.90005112	20.63667107	11.26338	0.128991	0.914471	
	Mean ΔCt (control) 11.13439					
			Fold gene	expression	1.010597	
			Standard	d deviation	0.159861	
		Star	ndard error o	f the mean	0.071492	
		Relative	standard err	or of mean	7.074255	
	31.13170242	21.61183929	9.519863	-1.61453	3.062109	
Female	33.99454498	23.12124252	10.8733	-0.26109	1.198381	
remaie						
	31.12455177	20.55188942	10.57266	-0.56173	1.476034	
	30.08351898	20.89955521	9.183964	-1.95042	3.864884	
Fold gene expression					2.400352	
	Standard deviation					
	Standard error of the mean				0.637827	
Relative standard error of mean					26.57221	

Table S.1.11: *Cyp2e1* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2e1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	25.65829277	20.44839668	5.209896	0.795858	0.576001		
	25.30113029	21.11756516	4.183565	-0.23047	1.17322		
na.t.	24.88606644	20.65575409	4.230312	-0.18373	1.135814		
Male	26.14256287	21.05667305	5.08589	0.671851	0.627701		
	23.60245705	20.1381073	3.46435	-0.94969	1.931456		
	24.94688797	20.63667107	4.310217	-0.10382	1.074616		
	Mean ΔCt (control) 4.414038						
			Fold gene	expression	1.086468		
			Standar	d deviation	0.488907		
		Sta	ndard error o	f the mean	0.199595		
		Relative	standard err	or of mean	18.37104		
	24.98876572	21.86377716	3.124989	-1.28905	2.443671		
	23.17398262	21.61183929	1.562143	-2.8519	7.21948		
Female							
Female	22.96647835	20.75888824	2.20759	-2.20645	4.615376		
	22.78383446	20.55188942	2.231945	-2.18209	4.538115		
	24.2583828	20.89955521	3.358828	-1.05521	2.078022		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.1.12: Cyp2j3 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	26.68292427	21.19024277	5.492682	-0.5398	1.453772	
	27.1415844	21.41108322	5.730501	-0.30198	1.232836	
0.0.1.	27.69579124	20.65575409	7.040037	1.007555	0.497388	
Male	27.2524395	21.7329464	5.519493	-0.51299	1.427004	
	26.89409065	20.46039772	6.433693	0.401211	0.757223	
	27.20636559	21.22787857	5.978487	-0.054	1.038136	
	•	Mean ΔCt (control)	6.032482			
			Fold gene	expression	1.067726	
			Standard	d deviation	0.381561	
		Sta	ndard error o	f the mean	0.155772	
		Relative	standard err	or of mean	14.58912	
	27.23317146	21.80290604	5.430265	-0.60222	1.518047	
	26.30326462	21.69820595	4.605059	-1.42742	2.689659	
Female	27.68857956	23.57600403	4.112576	-1.91991	3.783986	
remale	27.19836998	21.27204323	5.926327	-0.10616	1.076356	
	26.26378822	21.57160187	4.692186	-1.3403	2.532032	
	26.77512932	21.63104248	5.144087	-0.8884	1.851116	
Fold gene expression						
Standard deviation					0.969315	
Standard error of the mean					0.395721	
	Relative standard error of mean					

Table S.1.13: *Cyp2j4* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.88828468	21.19024277	3.698042	-0.79473	1.734756	
	25.57577705	21.41108322	4.164694	-0.32808	1.255343	
Mala	27.3547802	23.41542816	3.939352	-0.55342	1.467563	
Male	26.62948799	21.7329464	4.896542	0.403767	0.755882	
	26.09538651	20.46039772	5.634989	1.142214	0.453064	
	25.85091019	21.22787857	4.623032	0.130257	0.913669	
	Mean ΔCt (control) 4.492775					
			Fold gene	expression	1.096713	
			Standa	rd deviation	0.476182	
		St	andard error	of the mean	0.194401	
		Relativ	e standard er	ror of mean	17.72575	
	26.51693726	21.80290604	4.714031	0.221256	0.857818	
	25.38801003	21.69820595	3.689804	-0.80297	1.74469	
Female						
remaie	25.78979492	21.27204323	4.517752	0.024977	0.982836	
	24.92934799	21.57160187	3.357746	-1.13503	2.19623	
	25.56069183	21.63104248	3.929649	-0.56313	1.477467	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					16.96565	

Table S.1.14: *Cyp2j10* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	33.66636276	21.19024277	12.47612	0.984301	0.505471
	32.99433136	21.41108322	11.58325	0.091429	0.938593
Male	31.50067139	20.65575409	10.84492	-0.6469	1.565802
iviale	32.44746399	21.7329464	10.71452	-0.7773	1.713922
	33.06817245	21.22787857	11.84029	0.348475	0.785414
Mean ΔCt (control) 11.49182					
			Fold gene	expression	1.10184
			Standa	rd deviation	0.517769
		Sta	ndard error	of the mean	0.231553
		Relative	standard er	ror of mean	21.01512
	32.1833725	21.80290604	10.38047	-1.11135	2.160482
	32.82197189	21.69820595	11.12377	-0.36805	1.29061
Female					
remaie	33.14074326	21.27204323	11.8687	0.376881	0.770101
	33.77165222	21.57160187	12.20005	0.708231	0.61207
	33.09469986	21.63104248	11.46366	-0.02816	1.019712
Fold gene expression					1.170595
Standard deviation					0.610203
Standard error of the mean					0.272891
Relative standard error of mean					23.31216

Table S.1.15: *Cyp3a2* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp3a2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	33.68899918	22.86250496	10.82649	-0.63033	1.547923	
D.C. L.	34.66284943	22.36817551	12.29467	0.837846	0.559478	
Male	34.50880051	23.12959099	11.37921	-0.07762	1.055275	
	34.19684982	22.78125954	11.41559	-0.04124	1.028996	
	34.2376442	22.8694725	11.36817	-0.08866	1.063379	
	Mean ΔCt (control) 11.45683					
			Fold gen	e expression	1.05101	
			Standa	ard deviation	0.349708	
		S	tandard error	of the mean	0.156394	
		Relati	ive standard e	rror of mean	14.88035	
	35.65810394	25.21422005	10.44388	-1.01294	2.018025	
	34.7634964	23.18468285	11.57881	0.121986	0.918922	
Female						
Female	35.52947235	23.29874611	12.23073	0.773898	0.584835	
	32.70921326	22.336092	10.37312	-1.08371	2.119475	
	32.52546692	21.34239578	11.18307	-0.27376	1.208952	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.1.16: Cyp3a23 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	29.25986862	21.7662735	7.493595	-0.92593	1.899909
	29.97011375	22.06328773	7.906826	-0.5127	1.426717
Male					
ividie	31.88621712	21.89413071	9.992086	1.572561	0.336211
	29.0117569	21.03207779	7.979679	-0.43985	1.35646
	30.41083336	21.68539238	8.725441	0.305915	0.808929
		Mean ΔCt (control)	8.419526		
			Fold gene	expression	1.165645
			Standard	d deviation	0.603929
		Sta	ndard error o	f the mean	0.270085
		Relative	standard err	or of mean	23.17044
	30.167696	22.53539467	7.632301	-0.78722	1.725751
	29.09408951	22.48576355	6.608326	-1.8112	3.50934
Female					
remale	29.3111763	22.18027306	7.130903	-1.28862	2.442947
	29.20411301	21.98749924	7.216614	-1.20291	2.302038
	Fold gene expression				
	Standard deviation				
	Standard error of the mean				
Relative standard error of mean					14.90955

Table S.1.17: *Cyp4a1* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp4a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	28.21507835	21.19024277	7.024836	-0.74626	1.677438	
	28.80516052	21.41108322	7.394077	-0.37702	1.298654	
Male	30.39070892	20.65575409	9.734955	1.96386	0.256342	
iviale	28.91800117	21.7329464	7.185055	-0.58604	1.501121	
	27.7267971	20.46039772	7.266399	-0.5047	1.418824	
	29.24912453	21.22787857	8.021246	0.250151	0.840808	
	Mean ΔCt (control) 7.771095					
			Fold gene	expression	1.165531	
			Standa	rd deviation	0.526995	
		St	andard error	of the mean	0.215145	
		Relativ	e standard er	ror of mean	18.45895	
	28.3837204	21.80290604	6.580814	-1.19028	2.281971	
	28.31004715	21.69820595	6.611841	-1.15925	2.233418	
Female	29.10734367	23.57600403	5.53134	-2.23975	4.723168	
remale	27.51332474	21.27204323	6.241282	-1.52981	2.887484	
	27.57800484	21.57160187	6.006403	-1.76469	3.398014	
	28.09321594	21.63104248	6.462173	-1.30892	2.477562	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					12.93248	

Table S.1.18: *Cyp4a2* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp4a2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	36.9072113	21.3415451	15.56567	-0.92179	1.894459
Male	37.74897385 36.7227478	21.55198669 20.66747475	16.19699 16.05527	-0.29046 -0.43218	1.223034 1.34927
	38.39699173	20.26511002 Mean ΔCt (control)	18.13188 16.48745	1.64443	0.319873
				expression	1.196659
		St	Standard andard error o	f the mean	0.653112 0.326556
		Relativ	e standard err	or of mean	27.28896
	37.12036896	20.45787621	16.66249	0.175041	0.885743
Female	39.28878021	23.97945786	15.30932	-1.17813	2.262832
remate	39.78359222	21.98205376	17.80154	1.314086	0.40218
	38.68133163	21.74609947	16.93523	0.44778	0.73317
	36.59297562	20.99609566	15.59688	-0.89057	1.853911
Fold gene expression					
Standard deviation					0.791615
Standard error of the mean					0.354021
Relative standard error of mean					28.83923

Table S.1.19: *Cyp4a3* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp4a3)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.75954056	21.93746185	12.82208	0.799525	0.574538	
	33.94125748	22.16553879	11.77572	-0.24684	1.186601	
Male	33.30577087	22.12517929	11.18059	-0.84196	1.792486	
iviale	34.62602615	22.91962242	11.7064	-0.31615	1.245004	
	33.64625549	21.66824532	11.97801	-0.04454	1.031357	
	35.2222023	22.54968262	12.67252	0.649966	0.637295	
	Mean ΔCt (control) 12.02255					
			Fold gene	expression	1.07788	
			Standa	rd deviation	0.447224	
		Sta	ndard error	of the mean	0.182578	
		Relative	standard er	ror of mean	16.93864	
	34.84775925	23.05732346	11.79044	-0.23212	1.174558	
	35.12662888	22.89714622	12.22948	0.206929	0.86638	
Female						
remale	35.63805008	21.75164413	13.88641	1.863852	0.274742	
	35.51810455	21.95197487	13.56613	1.543576	0.343034	
	35.82263947	22.31215477	13.51048	1.487931	0.356523	
Fold gene expression					0.603047	
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					29.48072	

Table S.1.20: *Cyp4f1* mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp4f1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	27.99660683	21.93746185	6.059145	1.07968	0.473134	
	27.13925743	22.16553879	4.973719	-0.00575	1.003991	
na.i.	26.42750168	22.12517929	4.302322	-0.67714	1.598969	
Male	28.21333313	22.91962242	5.293711	0.314246	0.804271	
	25.93667221	21.66824532	4.268427	-0.71104	1.636981	
	Mean ΔCt (control) 4.979465					
			Fold gene	expression	1.103469	
			Standa	rd deviation	0.506679	
		S	tandard error	of the mean	0.226594	
		Relati	ve standard er	ror of mean	20.53466	
	26.77590942	23.05732346	3.718586	-1.26088	2.396417	
	26.09575272	22.89714622	3.198606	-1.78086	3.436305	
Female	27.04598045	24.28857994	2.757401	-2.22206	4.665605	
Female	25.25510406	21.75164413	3.50346	-1.476	2.781773	
	26.04194069	21.95197487	4.089966	-0.8895	1.852533	
	26.57153511	22.31215477	4.25938	-0.72008	1.647278	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.1.21: Cyp4f4 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	31.81648636	22.16553879	9.650948	-0.62922	1.54673
Male	32.06219864	22.12517929	9.937019	-0.34315	1.268523
iviale	34.97869492	22.91962242	12.05907	1.778904	0.291405
	31.31488037	21.66824532	9.646635	-0.63353	1.55136
	32.65685272	22.54968262	10.10717	-0.173	1.127399
		Mean ΔCt (control)	10.28017		
			Fold gene	expression	1.157083
			Standaı	rd deviation	0.517199
		Sta	ndard error	of the mean	0.231298
		Relative	standard er	ror of mean	19.98977
	31.54910088	23.05732346	8.491777	-1.78839	3.454295
	31.76424599	22.89714622	8.8671	-1.41307	2.663031
200					
Female	30.97224236	21.75164413	9.220598	-1.05957	2.084311
	29.91296196	21.95197487	7.960987	-2.31918	4.990491
	31.72716331	22.31215477	9.415009	-0.86516	1.821542
Fold gene expression					
Standard deviation					1.275657
Standard error of the mean					0.570491
Relative standard error of mean					18.99905

Table S.1.22: Cyp4f5 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	23.18008041	20.44839668	2.731684	0.413446	0.750828		
	22.81097031	21.11756516	1.693405	-0.62483	1.542031		
Mala	22.67821503	20.65575409	2.022461	-0.29578	1.227545		
Male	22.84887886	21.05667305	1.792206	-0.52603	1.439963		
	22.48275757	20.1381073	2.34465	0.026413	0.981859		
	23.961689	20.63667107	3.325018	1.006781	0.497656		
		Mean ΔCt (control)	2.318237				
			Fold gene	expression	1.073314		
			Standa	rd deviation	0.405379		
		S	tandard error	of the mean	0.165495		
		Relati	ve standard er	ror of mean	15.41908		
	23.19177437	21.86377716	1.327997	-0.99024	1.986516		
	22.12547493	21.61183929	0.513636	-1.8046	3.493327		
Female							
Female	22.42471123	20.75888824	1.665823	-0.65241	1.571796		
	22.01683044	20.55188942	1.464941	-0.8533	1.806624		
	23.30839348	20.89955521	2.408838	0.090601	0.939131		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relati	ve standard er	ror of mean	21.55602		

Table S.1.23: Cyp4f6 mRNA expression in the heart of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	23.12285233	20.44839668	2.674456	-0.10512	1.075587		
	24.0228138	21.11756516	2.905249	0.125669	0.916579		
Mala	23.4430542	20.65575409	2.7873	0.007721	0.994663		
Male	24.04642677	21.05667305	2.989754	0.210174	0.864433		
	22.64389992	20.1381073	2.505793	-0.27379	1.208977		
	23.45159721	20.63667107	2.814926	0.035347	0.975797		
		Mean ΔCt (control)	2.779579				
			Fold gene	expression	1.006006		
			Standa	rd deviation	0.122573		
		S	tandard error	of the mean	0.05004		
		Relati	ve standard er	ror of mean	4.974129		
	22.92057037	21.86377716	1.056793	-1.72279	3.300733		
	23.2191391	21.61183929	1.6073	-1.17228	2.253675		
Famala	23.83021355	23.12124252	0.708971	-2.07061	4.200638		
Female	22.76478958	20.75888824	2.005901	-0.77368	1.709623		
	22.65598297	20.55188942	2.104094	-0.67549	1.597135		
	23.19455338	20.89955521	2.294998	-0.48458	1.39918		
	Fold gene expression						
	Standard deviation						
		S	tandard error	of the mean	0.454506		
		Relati	ve standard er	ror of mean	18.85787		

S.2 Calculation of the fold change in the liver level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta CT$ method:

Table S.2.1: *Cyp1a1* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp1a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	27.1903019	21.88467026	5.305632	-0.67301	1.594398	
	27.54239464	21.40517998	6.137215	0.158571	0.895912	
Male	28.3407402	21.32709694	7.013643	1.035	0.488016	
iviale	27.38173103	22.04953957	5.332191	-0.64645	1.565314	
	27.83932114	21.73478508	6.104536	0.125893	0.916437	
		Mean ΔCt (control)	5.978643			
			Fold gene	expression	1.092015	
			Standa	rd deviation	0.4771	
		Sta	ndard error	of the mean	0.213366	
		Relative	e standard er	ror of mean	19.53872	
	25.96431923	20.89143944	5.07288	-0.90576	1.873536	
	27.02539635	20.72109985	6.304296	0.325653	0.797937	
Female						
remaie	26.12791252	21.08501244	5.0429	-0.93574	1.912876	
	27.49214745	20.66811371	6.824034	0.84539	0.55656	
	26.22910309	20.28797722	5.941126	-0.03752	1.026346	
			Fold gene	expression	1.233451	
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.2.2: Cyp1a2 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	19.89443588	21.88467026	-1.99023	-0.31153	1.241024		
	20.33478546	21.40517998	-1.07039	0.608309	0.655965		
Male	20.1892395	21.32709694	-1.13786	0.540846	0.687368		
iviale	19.78228569	22.04953957	-2.26725	-0.58855	1.503735		
	20.22178459	21.73478508	-1.513	0.165703	0.891494		
	20.02577019	22.11924934	-2.09348	-0.41478	1.333092		
		Mean ΔCt (control)	-1.6787				
			Fold gene	expression	1.052113		
			Standar	d deviation	0.356194		
	Standard error of the mean						
		Relative s	standard err	or of mean	13.82128		
	18.56800842						
	18.61347771	20.72109985	21.88467026 -1.99023 -0.31153 21.40517998 -1.07039 0.608309 21.32709694 -1.13786 0.540846 22.04953957 -2.26725 -0.58855 21.73478508 -1.513 0.165703 22.11924934 -2.09348 -0.41478 Mean ΔCt (control) -1.6787 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 20.89143944 -2.32343 -0.64473	1.346224			
Female	19.01803207	20.0019474	-0.98392	0.694788	0.6178		
remale	19.36067581	21.08501244	-1.72434	-0.04563	1.032136		
	19.1503315	20.66811371	-1.51778	0.160921	0.894454		
	18.05700874	20.28797722	-2.23097	-0.55227	1.466386		
	Standard deviation						
		Stan	dard error o	f the mean	0.149656		
		Relative s	standard err	or of mean	12.97508		

Table S.2.3: Cyp1b1 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1b1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	32.04344559	21.88467026	10.15878	0.216345	0.860743		
	31.14527512	21.40517998	9.740095	-0.20233	1.150559		
Male	31.12107658	21.32709694	9.79398	-0.14845	1.108378		
	31.87917519	22.04953957	9.829636	-0.11279	1.081321		
	31.92444992	21.73478508	10.18966	0.247235	0.84251		
		Mean ΔCt (control)	9.94243				
			Fold gene	expression	1.008702		
			Standa	rd deviation	0.14564		
		Sta	ndard error	of the mean	0.065132		
		Relative	e standard er	ror of mean	6.457025		
	32.96253967	20.89143944	12.0711	2.12867	0.228669		
	32.2551651	20.72109985	11.53407	1.591635	0.331795		
Famala	Male 32.04344559 21.88467026 10.15878 0.216345 31.14527512 21.40517998 9.740095 -0.20233 31.12107658 21.32709694 9.79398 -0.14845 31.87917519 22.04953957 9.829636 -0.11279 31.92444992 21.73478508 10.18966 0.247235 Mean ΔCt (control) 9.94243 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 32.96253967 20.89143944 12.0711 2.12867	0.145156					
remale	31.74835968	21.08501244	10.66335	0.720917	0.606712		
	31.26067543	20.66811371	10.59256	0.650132	0.637222		
	31.07182503	20.28797722	10.78385	0.841418	0.558095		
	Fold gene expression						
Standard deviation					0.210247		
	Standard error of the mean						
		Relative	e standard er	ror of mean	20.53708		

Table S.2.4: *Cyp2a1* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.4530735	22.33282661	-3.87975	-0.4252	1.34276		
	16.77977371	20.00734138	-3.22757	0.226984	0.854419		
Mala	18.48085594	22.09959602	-3.61874	-0.16419	1.120536		
iviale	17.07998276	20.5872097	-3.50723	-0.05268	1.037187		
	16.21551514	18.93590355	-2.72039	0.734163	0.601167		
	16.51438141	20.28801346	-3.77363	-0.31908	1.247535		
		Mean ΔCt (control)	-3.45455				
			Fold gene	expression	1.033934		
			Standa	rd deviation	0.271302		
		Star	ndard error	of the mean	0.110758		
		Relative	standard er	ror of mean	10.71233		
	14.4031105	19.47748566	-5.07438	-1.61982	3.073375		
	Male	5.897587					
Famala	15.90442657	20.04458618	-4.14016	-0.68561	1.60838		
remale	15.78352928	20.70932961	-4.9258	-1.47125	2.772618		
	14.43437099	19.17353249	-4.73916	-1.28461	2.436162		
	14.34364605	19.71036148	-5.36672	-1.91216	3.763732		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relative	standard er	ror of mean	18.48873		

Table S.2.5: Cyp2b1 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2b1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	25.49092293	22.33282661	3.158096	0.746702	0.595964		
	21.14013481	20.00734138	1.132793	-1.2786	2.426036		
Male	24.68083382	22.09959602	2.581238	0.169844	0.888939		
iviale	22.44049072	20.5872097	1.853281	-0.55811	1.472342		
	22.26746559	18.93590355	3.331562	0.920168	0.528448		
		Mean ΔCt (control)	2.411394				
			Fold gene	expression	1.182346		
			Standa	rd deviation	0.788729		
		Sta	ndard error	of the mean	0.35273		
		Relative	standard er	ror of mean	29.83311		
	23.88619041	19.47748566	4.408705	1.997311	0.250466		
	27.43149376	21.30310631	6.128387	3.716993	0.076045		
Female	26.25716782	20.04458618	6.212582	3.801188	0.071735		
remale	24.9078331	20.70932961	4.198503	1.787109	0.289752		
	25.42348862	19.17353249	6.249956	3.838562	0.0699		
	24.61631966	19.71036148	4.905958	2.494564	0.177444		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
		Relative	standard er	ror of mean	25.70708		

Table S.2.6: Cyp2b2 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2b2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	20.05709267	22.33282661	-2.27573	-0.12226	1.088437		
	16.89903641	20.00734138	-3.1083	-0.95483	1.93835		
Male	19.86379814	22.09959602	-2.2358	-0.08232	1.058721		
iviale	19.27927017	20.5872097	-1.30794	0.845536	0.556504		
	17.24253845	18.93590355	-1.69337	0.460111	0.72693		
	17.98830032	20.28801346	-2.29971	-0.14624	1.106679		
		Mean ΔCt (control)	-2.15348				
			Fold gene	expression	1.07927		
			Standar	d deviation	0.476803		
		Star	ndard error o	f the mean	0.194654		
		Relative	standard err	or of mean	18.03569		
	17.26698112	19.47748566	-2.2105	-0.05703	1.040321		
	20.46083832	21.30310631	-0.84227	1.311208	0.402983		
Familia	20.24387932	20.04458618	0.199293	2.352769	0.19577		
Female	18.24297714	20.70932961	-2.46635	-0.31288	1.242182		
	19.45895195	19.17353249	0.285419	2.438895	0.184425		
	18.26835251	19.71036148	-1.44201	0.711467	0.610699		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean					29.50253		

Table S.2.7: Cyp2c6 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)			
	18.30450439	22.33282661	-4.02832	-0.86967	1.827239			
	17.07195473	20.00734138	-2.93539	0.22327	0.856622			
Mole	18.48781204	22.09959602	-3.61178	-0.45313	1.369005			
iviale	17.60372734	20.5872097	-2.98348	0.175174	0.88566			
	16.7560215	18.93590355	-2.17988	0.978775	0.507411			
	17.07493019	20.28801346	-3.21308	-0.05443	1.038446			
		Mean ΔCt (control)	-3.15866					
			Fold gene	expression	1.08073			
			Standa	rd deviation	0.460247			
		Sta	ndard error	of the mean	0.187895			
		Relative	standard er	ror of mean	17.38594			
	15.96260357	19.47748566	-3.51488	-0.35623	1.280072			
Famala	16.42355156	20.04458618	-3.62103	-0.46238	1.377811			
remale	17.28323555	20.70932961	-3.42609	-0.26744	1.203668			
	18.30450439 22.33282661 -4 17.07195473 20.00734138 -2 18.48781204 22.09959602 -3 17.60372734 20.5872097 -2 16.7560215 18.93590355 -2 17.07493019 20.28801346 -3 Mean ΔCt (control) -3 Female	-3.38921	-0.23056	1.173288				
	15.87950802	19.71036148	-3.83085	-0.6722	1.593497			
Fold gene expression								
Standard deviation					0.169288			
		Sta	ndard error	of the mean	0.075708			
		Relative	Relative standard error of mean					

Table S.2.8: Cyp2c11 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2c11)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	16.40513229	22.33282661	-5.92769	-0.80642	1.748871		
	15.24024487	20.00734138	-4.7671	0.354174	0.782318		
no.1.	16.46564293	22.09959602	-5.63395	-0.51268	1.426701		
Male	15.21088886	20.5872097	-5.37632	-0.25505	1.193378		
	15.03774738	18.93590355	-3.89816	1.223114	0.428357		
	15.16361237	20.28801346	-5.1244	-0.00313	1.002172		
	'	Mean ΔCt (control)	-5.12127				
			Fold gen	e expression	1.096966		
			Standa	ard deviation	0.468621		
		S	tandard error	of the mean	0.191314		
		Relati	ve standard e	rror of mean	17.44026		
	26.28154755	21.30310631	4.978441	10.09971	0.000911		
Female	26.10057068	20.04458618	6.055984	11.17725	0.000432		
remaie	25.51708221	20.70932961	4.807753	9.929023	0.001026		
	25.46457291	19.17353249	6.29104	11.41231	0.000367		
	25.4253521	19.71036148	5.714991	10.83626	0.000547		
	Fold gene expression						
	Standard deviation						
		S	tandard error	of the mean	0.000132		
	Relative standard error of mean						

Table S.2.9: Cyp2c12 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c12</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	30.055336	22.33282661	7.722509	1.003765	0.498697		
	24.32423019	20.00734138	4.316889	-2.40186	5.284823		
Male							
ividie	28.42908859	20.5872097	7.841879	1.123135	0.459095		
	26.90167809	18.93590355	7.965775	1.247031	0.421314		
	26.03468132	20.28801346	5.746668	-0.97208	1.961661		
		Mean ΔCt (control)	6.718744				
			Fold gene	expression	1.725118		
			Standaı	rd deviation	2.093698		
		Sta	ındard error (of the mean	0.93633		
		Relative	e standard er	ror of mean	54.27628		
	19.28835678	19.47748566	-0.18913	-6.90787	120.0817		
	19.08397102	21.30310631	-2.21914	-8.93788	490.4218		
Female	19.26565742	20.04458618	-0.77893	-7.49767	180.7276		
remale	21.42381096	20.70932961	0.714481	-6.00426	64.18937		
	19.81147766	19.71036148	0.101116	-6.61763	98.19841		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relative	e standard er	ror of mean	40.52562		

Table S.2.10: *Cyp2c13* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2c13</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	16.92094994	22.33282661	-5.41188	-0.48094	1.395653		
	14.83981609	20.00734138	-5.16753	-0.23659	1.178204		
D.AI	16.39010429	22.09959602	-5.70949	-0.77856	1.715413		
Male	15.9002533	20.5872097	-4.68696	0.24398	0.844413		
	15.28989124	18.93590355	-3.64601	1.284924	0.410392		
	15.3242588	20.28801346	-4.96375	-0.03282	1.023009		
	'	Mean ΔCt (control)	-4.93094				
			Fold gen	e expression	1.094514		
			Standa	ard deviation	0.451418		
		S	tandard error	of the mean	0.184291		
		Relati	ve standard e	rror of mean	16.83766		
	25.75004387	21.30310631	4.446938	9.377874	0.001503		
	25.67071533	20.04458618	5.626129	10.55707	0.000664		
Female	25.67986488	20.70932961	4.970535	9.901471	0.001046		
	25.45578194	19.17353249	6.282249	11.21319	0.000421		
	25.90918732	19.71036148	6.198826	11.12976	0.000446		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relati	ve standard e	rror of mean	25.1233		

Table S.2.11: *Cyp2c23* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	15.66025448	21.88467026	-6.22442	-0.34325	1.268612
	15.91801167	21.40517998	-5.48717	0.393997	0.761018
Male	15.85260105	21.32709694	-5.4745	0.406669	0.754363
iviale	16.08489227	22.04953957	-5.96465	-0.08348	1.059572
	15.95038223	21.73478508	-5.7844	0.096762	0.935129
	15.7673893	22.11924934	-6.35186	-0.4707	1.385777
		Mean ΔCt (control)	-5.88117		
			Fold gene	expression	1.027412
			Standa	rd deviation	0.261409
		St	andard error	of the mean	0.10672
		Relativ	e standard er	ror of mean	10.38726
	15.4875412	20.89143944	-5.4039	0.477267	0.718337
	15.31864643	20.72109985	-5.40245	0.478712	0.717618
Female	15.85948563	20.0019474	-4.14246	1.738703	0.299639
remaie	15.6661129	21.08501244	-5.4189	0.462265	0.725846
	15.53650284	20.66811371	-5.13161	0.749554	0.594787
	14.84587955	20.28797722	-5.4421	0.439067	0.737611
			Fold gene	expression	0.632306
	Standard deviation				
		St	andard error	of the mean	0.069903
		Relativ	e standard er	ror of mean	11.05526

Table S.2.12: Cyp2d2 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2d2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	23.96702957	21.47598076	2.491049	0.438255	0.738027	
	22.2126503	20.26247215	1.950178	-0.10262	1.073719	
DA-I-	23.28487587	20.97448921	2.310387	0.257593	0.836483	
Male	23.06319427	20.92868996	2.134504	0.08171	0.944937	
	23.00608063	20.9538002	2.05228	-0.00051	1.000356	
	22.05823326	20.67986679	1.378366	-0.67443	1.595964	
	<u>'</u>	Mean ΔCt (control)	2.052794			
	Fold gene expression					
			Standar	d deviation	0.301006	
		Sta	ndard error o	f the mean	0.122885	
		Relative	e standard err	or of mean	11.91231	
	20.70548439	19.73490524	0.970579	-1.08221	2.117284	
	21.9311657	20.25279045	1.678375	-0.37442	1.296317	
Formula				-0.10262 0.257593 0.08171 -0.00051 -0.67443 e expression rd deviation of the mean rror of mean -1.08221 -0.37442 -0.41162 -1.87265 -1.72758 e expression rd deviation of the mean		
Female	22.03750229	20.39632988	1.641172	-0.41162	1.33018	
	20.27261925	20.09247208	0.180147	-1.87265	3.662039	
	20.80827522	20.48305702	0.325218	-1.72758	3.311709	
			Fold gene	expression	2.343506	
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.2.13: *Cyp2d3* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp2d3)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	21.05870438	21.47598076	-0.41728	-0.23108	1.17371		
	20.64422607	20.26247215	0.381754	0.567954	0.674573		
Mala	20.85568237	20.97448921	-0.11881	0.067394	0.954361		
iviale	20.79410553	20.92868996	-0.13458	0.051616	0.964855		
	20.4638958	20.9538002	-0.4899	-0.3037	1.234309		
	20.34148216	20.67986679	-0.33838	-0.15218	1.111251		
	Mean ΔCt (control) -0.1862						
			Fold gene	expression	1.018843		
			Standa	rd deviation	0.202259		
		Sta	ndard error	of the mean	0.082572		
		Relative	standard er	ror of mean	8.104475		
	20.32961273	19.73490524	0.594707	0.780908	0.582		
	Male	0.679545					
Female	20.78073311	18.72257614	2.058157	2.244357	0.211048		
remaie	21.78439331	20.39632988	1.388063	41728 -0.23108 881754 0.567954 11881 0.067394 13458 0.051616 9.4899 -0.3037 33838 -0.15218 9.1862 cold gene expression Standard deviation and error of the mean andard error of mean 9.4707 0.780908 9.71159 0.557359 9.58157 2.244357 9.88063 1.574264 9.72847 0.659047 9.56545 0.642745 9.610 gene expression Standard deviation Standard deviation and error of the mean	0.335814		
	20.56531906	20.09247208	0.472847	0.659047	0.633296		
	20.9396019	20.48305702	0.456545	0.642745	0.640493		
			Fold gene	expression	0.5137		
	Standard deviation						
		Sta	ndard error	of the mean	0.078696		
		Relative	standard er	ror of mean	15.31942		

Table S.2.14: Cyp2d4 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2d4)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	28.76899529	21.47598076	7.293015	0.113128	0.924581	
	27.19094658	20.26247215	6.928474	-0.25141	1.190372	
Mala	28.8094101	20.97448921	7.834921	0.655034	0.63506	
iviale	28.35194397	20.92868996	6.928474 -0.25141 7.834921 0.655034 7.423254 0.243367 6.371946 -0.80794 7.227711 0.047824 7.179887 Fold gene expression Standard deviation tandard error of the mean ve standard error of mean 7.432837 0.25295 7.760088 0.580201 8.028717 0.84883 7.712959 0.533072 6.910984 -0.2689	0.243367	0.844771	
	27.32574654	20.9538002	6.371946	-0.80794	1.75071	
	27.90757751	20.67986679	598076 7.293015 0.113128 247215 6.928474 -0.25141 448921 7.834921 0.655034 868996 7.423254 0.243367 538002 6.371946 -0.80794 986679 7.227711 0.047824 ACT (control) 7.179887 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 490524 7.432837 0.25295 279045 7.760088 0.580201 257614 8.028717 0.84883 632988 7.712959 0.533072 247208 6.910984 -0.2689 305702 6.63217 -0.54772 Fold gene expression Standard deviation	0.047824	0.967394	
		Mean ΔCt (control)	7.179887			
			Fold gen	e expression	1.052148	
			Standa	ard deviation	0.386657	
			Standard error	of the mean	0.157852	
		Relat	tive standard e	rror of mean	15.00284	
	27.16774178	19.73490524	7.432837	0.25295	0.839179	
	28.01287842	20.25279045	7.760088	0.580201	0.668871	
Formula	26.75129318	18.72257614	8.028717	0.84883	0.555235	
Female	28.10928917	20.39632988	7.712959	0.533072	0.691081	
	27.00345612	20.09247208	6.910984	-0.2689	1.204891	
	Male 28.76899529 21.47598076 7.293015 0.113128 27.19094658 20.26247215 6.928474 -0.25141 28.8094101 20.97448921 7.834921 0.655034 28.35194397 20.92868996 7.423254 0.243367 27.32574654 20.9538002 6.371946 -0.80794 27.90757751 20.67986679 7.227711 0.047824 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean Relative standard error of mean 27.16774178 19.73490524 7.432837 0.25295 28.01287842 20.25279045 7.760088 0.580201 26.75129318 18.72257614 8.028717 0.84883 28.10928917 20.39632988 7.712959 0.533072 27.00345612 20.09247208 6.910984 -0.2689 27.11522675 20.48305702 6.63217 -0.54772 Fold gene expression <td colspa<="" td=""><td>1.461771</td></td>	<td>1.461771</td>	1.461771			
			Fold gen	e expression	0.903505	
	Standard deviation					
			Standard error	of the mean	0.144695	
		Relat	tive standard e	rror of mean	16.01489	

Table S.2.15: Cyp2e1 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2e1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	17.13682938	21.88467026	-4.74784	0.770882	0.586059		
	16.604105	21.40517998	-4.80107	0.717648	0.608088		
Mala	15.93671608	21.32709694	-5.39038	0.128342	0.914882		
iviale	16.03761482	22.04953957	-6.01192	-0.4932	1.407565		
	15.62899399	21.73478508	-6.10579	-0.58707	1.502191		
	16.06392288	22.11924934	-6.05533	-0.5366	1.450553		
	Mean ΔCt (control) -5.51872						
	Fold gene expression						
			Standa	rd deviation	0.4282		
		Sta	ndard error	of the mean	0.174812		
		Relative	e standard er	ror of mean	16.21297		
	14.77024651	20.89143944	-6.12119	-0.60247	1.518314		
	Male 16.03761482 22.04953957 -6.01192 -0.4932 15.62899399 21.73478508 -6.10579 -0.58707 16.06392288 22.11924934 -6.05533 -0.5366 Mean ΔCt (control) -5.51872 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 14.77024651 20.89143944 -6.12119 -0.60247 15.0603714 20.72109985 -5.66073 -0.14201 14.62727261 20.0019474 -5.37467 0.144048 15.01053524 21.08501244 -6.07448 -0.55575	1.103438					
Familia	14.62727261	20.0019474	-5.37467	0.770882 0.717648 0.128342 0.4932 0.58707 0.5366 0.0000000000000000000000000000000000	0.904976		
remaie	15.01053524	21.08501244	-6.07448		1.469937		
	Male16.60410521.40517998-4.801070.71764815.9367160821.32709694-5.390380.12834216.0376148222.04953957-6.01192-0.493215.6289939921.73478508-6.10579-0.5870716.0639228822.11924934-6.05533-0.5366Fold gene expressionStandard deviationStandard error of the meanRelative standard error of meanRelative standard error of mean14.7702465120.89143944-6.12119-0.6024715.060371420.72109985-5.66073-0.1420114.6272726120.0019474-5.374670.14404815.0105352421.08501244-6.07448-0.5557514.8683586120.66811371-5.79976-0.2810314.3444652620.28797722-5.94351-0.42479Fold gene expression	1.215064					
	14.34446526	20.28797722	-5.94351	-0.42479	1.342376		
	Fold gene expression						
	Standard deviation						
		Sta	ndard error	of the mean	0.094895		
Relative standard error of mean							

Table S.2.16: *Cyp2j3* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	21.25211906	21.88467026	-0.63255	0.0593	0.95973		
	21.33333778	21.40517998	-0.07184	0.620009	0.650667		
Male	20.96491814	21.32709694	-0.36218	0.329672	0.795717		
iviale	20.95210457	22.04953957	-1.09743	-0.40558	1.324625		
	20.83299446	21.73478508	-0.90179	-0.20994	1.15664		
	21.03394127	22.11924934	-1.08531	-0.39346	1.313537		
	Mean ΔCt (control) -0.69185						
	Fold gene expression						
			Standar	d deviation	0.278166		
		Sta	ndard error o	f the mean	0.113561		
		Relative	e standard err	or of mean	10.98813		
	20.60764885	20.89143944	-0.28379	0.40806	0.753636		
	21.56151772	20.72109985	0.840418	1.532269	0.345733		
Female	21.03738594	21.08501244	-0.04763	0.644224	0.639837		
	21.83067703	20.66811371	1.162563	1.854414	0.276545		
	21.32190323	20.28797722	1.033926	1.725777	0.302336		
			Fold gene	expression	0.463617		
	Standard deviation						
		Sta	ndard error o	f the mean	0.097486		
		Relative	standard err	or of mean	21.02716		

Table S.2.17: *Cyp2j4* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	21.84089279	21.47598076	0.364912	-0.35876	1.282325		
	20.95622063	20.26247215	0.693748	-0.02993	1.020959		
Male	21.94081879	20.97448921	0.96633	0.242656	0.845188		
iviale	21.9086647	20.92868996	0.979975	0.256301	0.837232		
	21.56383514	20.9538002	0.610035	-0.11364	1.081954		
	21.40690994	20.67986679	0.727043	0.003369	0.997667		
	Mean ΔCt (control) 0.723674						
			Fold gene	expression	1.010888		
			Standard	d deviation	0.165298		
		Sta	ndard error o	f the mean	0.067483		
		Relative	standard err	or of mean	6.67559		
	21.12786674	19.73490524	1.392962	0.669288	0.628817		
	21.44385529	20.25279045	1.191065	0.467391	0.723271		
Female	21.31331444	18.72257614	6 0.364912 -0.35876 5 0.693748 -0.02993 1 0.96633 0.242656 6 0.979975 0.256301 2 0.610035 -0.11364 9 0.727043 0.003369 entrol) 0.723674 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 4 1.392962 0.669288 5 1.191065 0.467391 4 2.590738 1.867064 8 1.305544 0.58187 8 1.222208 0.498534 2 0.6628 -0.06087 Fold gene expression Standard deviation Standard deviation	1.867064	0.274131		
remale	21.70187378	20.39632988		0.58187	0.668097		
	21.3146801	20.09247208	1.222208	0.498534	0.707826		
	21.14585686	20.48305702	0.6628	-0.06087	1.043097		
Fold gene expression							
	Standard deviation						
		Sta	ndard error o	f the mean	0.100296		
Relative standard error of mean							

Table S.2.18: *Cyp2j10* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)			
	31.16314888	21.10664368	10.05651	-0.64912	1.568211			
	30.35653496	18.86980438	11.48673	0.781106	0.581921			
B. A I.	31.36725807	21.02325821	10.344	-0.36162	1.284872			
Male	30.54577827	20.01047897	10.5353	-0.17033	1.125312			
	31.51574898	20.85785294	94 10.6579 -0.04773 86 11.15332 0.447692 ntrol) 10.70562 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean	-0.04773	1.033636			
	30.63983536	19.48651886	11.15332	0.447692	0.733215			
	Mean ΔCt (control) 10.70562							
	Fold gene expression							
			Standa	rd deviation	0.360271			
		S	tandard error	of the mean	0.14708			
		Relati	ve standard er	ror of mean	13.94747			
	31.54032135	19.73880577	11.80152	1.095891	0.467847			
	31.89519119	20.37353134	11.52166	0.816035	0.568001			
Female	30.68645477	19.03637123	11.65008	0.944459	0.519624			
Female	30.30920219	19.04911041	11.26009	0.554467	0.680908			
	30.94412613	18.86980438 11.48673 0.781106 21.02325821 10.344 -0.36162 20.01047897 10.5353 -0.17033 20.85785294 10.6579 -0.04773 19.48651886 11.15332 0.447692 Mean ΔCt (control) 10.70562 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 19.73880577 11.80152 1.095891 20.37353134 11.52166 0.816035 19.03637123 11.65008 0.944459 19.04911041 11.26009 0.554467 17.97408104 12.97005 2.264421 18.18707848 12.2209 1.515275 Fold gene expression Standard deviation	0.208133					
	30.40797806	18.18707848	12.2209	1.515275	0.34983			
			Fold gene	e expression	0.465724			
			Standa	rd deviation	0.167008			
		S	tandard error	of the mean	0.068181			
	Relative standard error of mean							

Table S.2.19: *Cyp3a1* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	17.78360367	21.47598076	-3.69238	0.083422	0.943816	
	16.49839973	20.26247215	-3.76407	0.011727	0.991905	
Mala	17.98199463	20.97448921	-2.99249	0.783305	0.581034	
Male	17.20392227	20.92868996	-3.72477	0.051031	0.965246	
	17.08751869	20.9538002	-3.86628	-0.09048	1.064726	
	16.06506538	20.67986679	-4.6148	-0.839	1.788813	
Mean ΔCt (control) -3.7758						
			Fold gene	expression	1.055923	
			Standar	d deviation	0.396891	
		Sta	ndard error o	f the mean	0.16203	
		Relative	standard err	or of mean	15.34487	
	16.05394554	19.73490524	-3.68096	0.094839	0.936376	
	17.18310738	20.25279045	-3.06968	0.706116	0.612968	
Familia	17.4712429	18.72257614	-1.25133	138 0.083422 107 0.011727 149 0.783305 177 0.051031 128 -0.09048 148 -0.839 158 158 158 158 159 169 0.094839 169 0.094839 160 0.706116 161 0.33 2.524466 161 0.14232 161 0.14232 162 0.432078 163 0.432078 164 0.432078 165 0.432078 166 0.432078 166 0.432078 166 0.432078 167 0.432078 168 0.706116	0.173804	
Female	16.47821426	20.39632988	-3.91812		1.103676	
	17.50841522	20.09247208	-2.58406	1.191742	0.437774	
	17.13933563	20.48305702	-3.34372	0.432078	0.741194	
			Fold gene	expression	0.667632	
	Standard deviation					
		Sta	ndard error o	f the mean	0.137533	
	Relative standard error of mean					

Table S.2.20: Cyp3a2 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp3a2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	19.74202156	20.94281006	-1.20079	-0.1165	1.084099	
	16.85701942	18.45859718	-1.60158	-0.51729	1.43126	
Mala	19.55093384	20.57956696	-1.02863	0.055659	0.962155	
iviale	19.21705246	19.27609253	-0.05904	1.025252	0.491324	
	18.96092796	20.4620533	-1.50113	-0.41683	1.334994	
	17.89879799	19.01338577	-1.11459	-0.0303	1.021221	
	Mean ΔCt (control) -1.08429					
	Fold gene expression					
			Standard	d deviation	0.33115	
		Sta	ndard error o	f the mean	0.135191	
		Relative	standard err	or of mean	12.82438	
	20.46822929	19.9620285	0.506201	1.590493	0.332058	
Familia	Male 19.21705246 19.27609253 -0.05904 1.025252 18.96092796 20.4620533 -1.50113 -0.41683 17.89879799 19.01338577 -1.11459 -0.0303 Mean ΔCt (control) -1.08429 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 20.46822929 19.9620285 0.506201 1.590493 20.12581253 18.85101891 1.274794 2.359086	0.194915				
Female	17.68200111	18.57664299	-1.20079 -0.1165 -1.60158 -0.51729 -1.02863 0.055659 -0.05904 1.025252 -1.50113 -0.41683 -1.11459 -0.0303 -1.08429 Fold gene expression Standard deviation Standard error of the mean tive standard error of mean 1.274794 2.359086 -0.89464 0.18965 1.195944 2.280236 -0.63286 0.45143 Fold gene expression Standard deviation Standard deviation	0.18965	0.876818	
	19.21388435	18.01794052	1.195944	2.280236	0.205864	
	17.55674362	18.18960571	-0.63286	0.45143	0.731318	
			Fold gene	expression	0.468195	
	Standard deviation					
		Sta	ndard error o	f the mean	0.14111	
	Relative standard error of mean					

Table S.2.21: *Cyp3a9* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp3a9</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)			
	26.52981758	21.10664368	5.423174	0.613039	0.653818			
	24.6495533	18.86980438	5.779749	0.969614	0.510643			
Male	26.36856842	21.02325821	5.34531	0.535176	0.690075			
iviale								
	24.46473312	20.85785294	3.60688	-1.20325	2.302585			
	23.38207817	19.48651886	3.895559	-0.91458	1.885014			
	Mean ΔCt (control) 4.810135							
	Fold gene expression							
			Standa	ard deviation	0.824338			
		S	tandard error	of the mean	0.368655			
		Relati	ive standard e	rror of mean	30.50705			
	21.68917656	19.73880577	1.950371	-2.85976	7.258964			
				0.613039 0.969614 0.535176 -1.20325 -0.91458 ene expression dard deviation or of the mean error of mean -2.85976 -2.03833 -3.38114 -4.24487 -3.43581 ene expression				
Female	21.80817795	19.03637123	2.771807	-2.03833	4.107691			
remale	20.47810936	19.04911041	1.428999	-3.38114	10.41893			
	18.53935051	17.97408104	0.565269	-4.24487	18.95971			
	19.56140137	18.18707848	1.374323	-3.43581	10.82137			
			Fold gen	e expression	10.31333			
	Standard deviation							
		S	tandard error	of the mean	2.478591			
		Relati	ive standard e	rror of mean	24.03288			

Table S.2.22: *Cyp3a18* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a18</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	31.24169731	20.8724823	10.36922	0.285637	0.82038		
	30.88352966	20.7446537	10.13888	0.055297	0.962396		
Male	31.58528328	21.34989548	10.23539	0.151809	0.900121		
iviale	31.07600021	21.55722809	9.518772	-0.56481	1.479189		
	31.27173042	21.11608887	10.15564	0.072063	0.951277		
	Mean ΔCt (control) 10.08358						
	Fold gene expression						
			Standa	ard deviation	0.261285		
		S	tandard error	of the mean	0.11685		
		Relati	ve standard e	rror of mean	11.42596		
	35.04505157	21.04141045	14.00364	3.920063	0.066061		
	36.68312073	20.96590233	15.71722	5.63364	0.020142		
Female	33.97431946	19.11339188	14.86093	4.777349	0.036465		
remale	34.89445114	20.8316555	14.0628	3.979217	0.063407		
	35.3510704	20.29788017	15.05319	4.969612	0.031915		
			Fold gen	e expression	0.043598		
	Standard deviation						
		S	tandard error	of the mean	0.00904		
		Relati	ve standard e	rror of mean	20.73543		

Table S.2.23: *Cyp3a23* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.23514175	21.10664368	-2.8715	0.226627	0.85463		
	16.25152397	18.86980438	-2.61828	0.479849	0.717053		
Mala	18.53732872	21.02325821	-2.48593	0.6122	0.654198		
Male	16.53012276	20.01047897	-3.48036	-0.38223	1.303352		
	17.75306892	20.85785294	-3.10478	-0.00665	1.004623		
	15.45859528	19.48651886	-4.02792	-0.92979	1.905004		
		Mean ΔCt (control)	-3.09813				
			Fold gene	expression	1.073144		
			Standar	d deviation	0.468988		
		St	andard error o	f the mean	0.191463		
		Relativ	e standard err	or of mean	17.84136		
	16.46657181	19.73880577	-3.27223	-0.1741	1.128264		
	17.5465374	20.37353134	-2.82699	0.271135	0.828667		
Female	17.74831581	19.03637123	-1.28806	1.810074	0.285176		
Female	15.51676464	19.04911041	-3.53235	-0.43422	1.351177		
	16.39876938	17.97408104	-1.57531	1.522818	0.348006		
	15.97563839	18.18707848	-2.21144	0.886689	0.540854		
	Fold gene expression						
	Standard deviation						
		St	andard error o	f the mean	0.176261		
		Relativ	e standard err	or of mean	23.59506		

Table S.2.24: *Cyp4a1* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp4a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)			
	21.02525902	21.88467026	-0.85941	-0.02779	1.019446			
	21.30592918	21.40517998	-0.09925	0.732375	0.601912			
20.1	21.27701378	21.32709694	-0.05008	0.781542	0.581744			
Male	21.09216309	22.04953957	-0.95738	-0.12575	1.091075			
	20.35990143	21.73478508	-1.37488	-0.54326	1.45726			
	20.47050095	22.11924934	-1.64875	-0.81712	1.761889			
		Mean ΔCt (control)	-0.83163					
	Fold gene expression							
			Standa	rd deviation	0.466418			
		St	andard error	of the mean	0.190414			
		Relativ	e standard er	ror of mean	17.54074			
	19.23550034	20.89143944	-1.65594	-0.82431	1.770692			
	18.66377068	20.72109985	-2.05733	-1.2257	2.338695			
Female	20.0784111	20.0019474	0.076464	0.908089	0.53289			
remaie	20.26852798	21.08501244	-0.81648	0.015141	0.98956			
	19.99027252	20.66811371	-0.67784	0.153784	0.898889			
	19.10308075	20.28797722	-1.1849	-0.35327	1.277454			
	Fold gene expression							
	Standard deviation							
	Standard error of the mean							
		Relativ	e standard er	ror of mean	20.55521			

Table S.2.25: *Cyp4a2* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.02228355	20.8724823	-2.8502	0.591914	0.663462		
	17.504879	20.7446537	-3.23977	0.202338	0.869141		
Mole	18.159132	21.34989548	-3.19076	0.251349	0.84011		
iviale	17.97693634	21.55722809	-3.58029	-0.13818	1.100515		
	17.72946548	22.07900047	-4.34953	-0.90742	1.875691		
	Mean ΔCt (control) -3.44211						
	Fold gene expression						
			Standa	rd deviation	0.476586		
		St	andard error	of the mean	0.213136		
		Relativ	e standard er	ror of mean	19.92324		
	28.64541245	20.96590233	7.67951	11.12162	0.000449		
Familia	27.9602375	19.11339188	8.846846	12.28896	0.0002		
remaie	28.15448189	20.8316555	7.322826	10.76494	0.000575		
	18.02228355 17.504879 18.159132 17.97693634 17.72946548 28.64541245 27.9602375	20.86181259	8.521231	11.96334	0.00025		
	28.95898056	20.29788017	8.6611	12.10321	0.000227		
Fold gene expression							
Standard deviation							
	Standard error of the mean						
	Relative standard error of mean						

Table S.2.26: *Cyp4a3* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.83239746	20.8724823	-2.04008	0.033601	0.976979		
	18.80909157	20.7446537	-1.93556	0.138123	0.9087		
Male	19.2974205	21.34989548	-2.05247	0.02121	0.985406		
	19.02155495	21.55722809	-2.53567	-0.46199	1.377438		
	19.31145668	21.11608887	-1.80463	0.269053	0.829864		
		Mean ΔCt (control)	-2.07369				
			Fold gene	expression	1.015677		
			Standa	rd deviation	0.211693		
		S	tandard error	of the mean	0.094672		
		Relati	ve standard er	ror of mean	9.321066		
	18.9805069	21.04141045	-2.0609	0.012782	0.991179		
	18.61083412	20.96590233	-2.35507	-0.28138	1.215359		
Female	18.51231956	19.11339188	-0.60107	1.472613	0.360329		
remale	19.00722885	20.8316555	-1.82443	0.249259	0.841329		
	19.07697487	20.86181259	-1.78484	0.288848	0.818556		
	18.78233337	20.29788017	-1.51555	0.558139	0.679178		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relati	ve standard er	ror of mean	14.4219		

Table S.2.27: *Cyp4a8* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a8</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	35.93387222	20.8724823	15.06139	0.522375	0.696225		
	35.91472626	20.7446537	15.17007	0.631058	0.645703		
Mala	35.8786087	21.34989548	14.52871	-0.0103	1.007166		
Male	35.93473434	21.55722809	14.37751	-0.16151	1.118456		
	35.74394608	21.11608887	14.62786	0.088842	0.940277		
	35.5475502	22.07900047	13.46855	-1.07047	2.10011		
	Mean ΔCt (control) 14.53901						
	Fold gene expression						
			Standar	d deviation	0.52966		
		Star	ndard error o	f the mean	0.216233		
		Relative	standard err	or of mean	19.93561		
	35.97494888	21.04141045	14.93354	0.394524	0.760741		
Formula	35.08459473	19.11339188	15.9712	1.432188	0.370568		
Female	36.28350067	20.8316555	15.45185	0.91283	0.531142		
	37.85520172	20.86181259	16.99339	2.454374	0.182457		
	35.62527466	20.29788017	15.32739	0.78838	0.578994		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.2.28: *Cyp4f1* mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	16.8526535	20.8724823	-4.01983	0.343711	0.788012		
	16.50281715	20.7446537	-4.24184	0.121703	0.919102		
Mala	16.89003944	21.34989548	-4.45986	-0.09632	1.069041		
Male	17.26995087	21.55722809	-4.28728	0.076262	0.948512		
	16.77445984	21.11608887	-4.34163	0.02191	0.984928		
	17.24819183	22.07900047	-4.83081	-0.46727	1.38249		
		Mean ΔCt (control)	-4.36354				
			Fold gen	e expression	1.015347		
			Standa	ard deviation	0.201997		
		St	andard error	of the mean	0.082465		
		Relativ	e standard e	rror of mean	8.121839		
	17.07052994	21.04141045	-3.97088	0.392659	0.761724		
	16.98641396	20.96590233	-3.97949	0.384051	0.766283		
Female	17.18163109	19.11339188	-1.93176	2.431779	0.185337		
remale	17.26861572	20.8316555	-3.56304	0.8005	0.57415		
	17.37354469	20.86181259	-3.48827	0.875271	0.545151		
	16.61227226	20.29788017	-3.68561	0.677931	0.625061		
	Fold gene expression						
	Standard deviation						
		St	andard error	of the mean	0.086925		
		Relativ	e standard e	rror of mean	15.08376		

Table S.2.29: Cyp4f4 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	19.58791351	20.8724823	-1.28457	0.080768	0.945554	
	19.55678749	20.7446537	-1.18787	0.177471	0.884252	
DA-I-	20.3129673	21.34989548	-1.03693	0.328409	0.796415	
Male	19.98579597	21.55722809	-1.57143	-0.2061	1.153562	
	20.00549126	21.11608887	-1.1106	0.254739	0.838139	
	20.07837296	22.07900047	-2.00063	-0.63529	1.553251	
		Mean ΔCt (control)	-1.36534			
			Fold gene	expression	1.028529	
			Standard	d deviation	0.285934	
		St	andard error o	f the mean	0.116732	
		Relativ	e standard err	or of mean	11.34944	
	19.95122147	21.04141045	-1.09019	0.275148	0.826366	
	19.77783394	20.96590233	-1.18807	0.177268	0.884376	
Familia	19.5415535	19.11339188	0.428162	157 0.080768 787 0.177471 793 0.328409 743 -0.2061 706 0.254739 706 0.63529 707 0.63529 708 108 108 108 108 108 108 108 108 108 1	0.288472	
Female	19.99121475	20.8316555	-0.84044	0.524896	0.695009	
	19.86571503	20.86181259	-0.9961	0.369239	0.774191	
	19.12087059	20.29788017	-1.17701	0.188327	0.877623	
Fold gene expression						
Standard deviation						
	Standard error of the mean					
	Relative standard error of mean					

Table S.2.30: Cyp4f5 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	24.76859283	20.8724823	3.896111	-0.34808	1.272867		
	24.69163322	20.7446537	3.94698	-0.29721	1.228768		
Mala	26.22362328	21.34989548	4.873728	0.629536	0.646384		
iviale	25.45830536	21.55722809	3.901077	-0.34311	1.268492		
	25.93483162	21.11608887	4.818743	0.574551	0.671495		
	26.10751534	22.07900047	4.028515	-0.21568	1.161249		
	Mean ΔCt (control) 4.244192						
			Fold gene	expression	1.041543		
			Standar	d deviation	0.299161		
		St	andard error o	f the mean	0.122132		
		Relativ	e standard err	or of mean	11.72605		
	26.00374031	21.04141045	4.96233	0.718138	0.607882		
	25.39862633	20.96590233	4.432724	0.188532	0.877498		
Familia	25.12063026	19.11339188	6.007238	1.763046	0.294625		
Female	Male 24.69163322 20.7446537 3.9469 26.22362328 21.34989548 4.87372 25.45830536 21.55722809 3.90107 25.93483162 21.11608887 4.81874 26.10751534 22.07900047 4.02851 Mean ΔCt (control) 4.24419 Start Standard err Relative standard Relative standard 25.39862633 20.96590233 4.43272 25.39862633 20.96590233 4.43272 25.12063026 19.11339188 6.00723 24.80643654 20.8316555 3.97478 25.6361084 20.86181259 4.77429 24.71218109 20.29788017 4.41436	3.974781	-0.26941	1.205316			
	25.6361084	20.86181259	4.774296	0.530104	0.692505		
	24.71218109	20.29788017	4.414301	0.170109	0.888776		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.2.31: Cyp4f6 mRNA expression in the liver of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	23.78008461	21.88467026	1.895414	-0.17112	1.125932		
	24.18930626	21.40517998	2.784126	0.717592	0.608112		
Male	23.3608532	21.32709694	2.033756	-0.03278	1.02298		
IVIdle	24.5755558	22.04953957	2.526016	0.459482	0.727248		
	23.54052353	21.73478508	1.805738	-0.2608	1.19814		
	23.47340584	22.11924934	1.354156	-0.71238	1.638503		
	Mean ΔCt (control) 2.066535						
	Fold gene expression						
			Standa	rd deviation	0.367189		
		Sta	ndard error	of the mean	0.149904		
		Relative	e standard er	ror of mean	14.22935		
	22.72532654	20.89143944	1.833887	-0.23265	1.174989		
	22.93718719	20.72109985	2.216087	0.149553	0.90153		
Female	23.35550499	20.0019474	3.353558	1.287023	0.409796		
remaie	23.8314743	21.08501244	2.746462	0.679927	0.624197		
	23.29389572	20.66811371	2.625782	0.559247	0.678656		
	22.76893044	20.28797722	2.480953	0.414419	0.750322		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

S.3 Calculation of the fold change in the kidney level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta CT$ method:

Table S.3.1: *Cyp1a1* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp1a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	27.0172863	21.44843102	5.568855	0.010524	0.992732		
	26.90682793	20.36410713	6.542721	0.98439	0.505439		
0.0-1-	27.00440216	20.92403412	6.080368	0.522037	0.696388		
Male							
	25.0068264	20.36148262	4.645344	-0.91299	1.88294		
	26.14905357	21.19468689	4.954367	-0.60396	1.519887		
		Mean ΔCt (control)	5.558331				
			Fold gene	expression	1.119477		
			Standar	d deviation	0.573436		
		St	andard error o	f the mean	0.256448		
		Relativ	e standard err	or of mean	22.90787		
	20.73973083	20.15422821	0.585503	-4.97283	31.40295		
	23.42601204	21.36679268	2.059219	-3.49911	11.30674		
Female							
remale	20.66752052	20.15905762	0.508463	-5.04987	33.12545		
	23.18659592	21.71059608	1.476	-4.08233	16.93964		
	21.16689491	21.34286499	-0.17597	-5.7343	53.23492		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relativ	e standard err	or of mean	25.0291		

Table S.3.2: *Cyp1a2* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp1a2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	29.06984901	20.36410713	8.705742	-0.04391	1.030907		
Mala	29.76661682	20.92403412	8.842583	0.092927	0.937618		
iviale	30.68917084	21.1464119	9.542759	0.793103	0.577101		
	28.79200363	20.36148262	8.430521	-0.31913	1.247582		
	29.42136002	21.19468689	8.226673	-0.52298	1.436923		
	Mean ΔCt (control) 8.749656						
			Fold gene	expression	1.046026		
			Standa	rd deviation	0.326074		
		St	andard error	of the mean	0.145825		
		Relativ	e standard er	ror of mean	13.94081		
	29.45443153	20.15422821	9.300203	0.550548	0.682761		
Male 29.76661682 20.92403412 30.68917084 21.1464119 28.79200363 20.36148262 29.42136002 21.19468689 Mean ΔCt (control) Sta Relative 29.45443153 20.15422821 29.37855339 21.36679268 29.56826591 20.80569077 28.72387314 20.15905762 29.58572006 21.34286499	8.011761	-0.73789	1.66774				
	29.56826591	20.80569077	8.762575	0.01292	0.991085		
Female	28.72387314	20.15905762	8.564816	-0.18484	1.136691		
	29.58572006	21.34286499	8.242855	-0.5068	1.420896		
	Fold gene expression						
	Standard deviation						
		St	andard error	of the mean	0.170401		
	Relative standard error of mean						

Table S.3.3: *Cyp1b1* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp1b1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	26.20146561	20.6704731	5.530993	-0.08746	1.062496		
	25.3973484	19.91502762	5.482321	-0.13613	1.098952		
na-l-	25.60379028	20.17363167	5.430159	-0.18829	1.139413		
Male	26.28948021	20.20077324	6.088707	0.470257	0.721836		
	25.80596733	20.43040276	5.375565	-0.24288	1.183357		
	26.35974121	20.55678749	5.802954	0.184504	0.879951		
		Mean ΔCt (control)	5.61845				
	Fold gene expression						
			Standar	d deviation	0.177374		
		St	andard error o	f the mean	0.072413		
		Relativ	e standard err	or of mean	7.13892		
	26.32404518	19.731493	6.592552	0.974103	0.509056		
	26.9568615	20.44017982	6.516682	0.898232	0.536544		
Famala	27.86956787	19.90690994	5.530993 -0.08746 5.482321 -0.13613 5.430159 -0.18829 6.088707 0.470257 5.375565 -0.24288 5.802954 0.184504 5.61845 Fold gene expression Standard deviation tandard error of the mean ve standard error of mean 6.592552 0.974103 6.516682 0.898232 7.962658 2.344208 7.281013 1.662564 5.725563 0.107114 7.755882 2.137433 Fold gene expression Standard deviation	0.196935			
Female	25.98503113	18.70401764	7.281013	1.662564	0.315877		
	26.5469799	20.82141685	5.725563	0.107114	0.928444		
	26.00157547	18.24569321	7.755882	2.137433	0.227284		
	Fold gene expression						
Standard deviation							
Standard error of the mean							
		Relativ	e standard err	or of mean	24.59968		

Table S.3.4: Cyp2a1 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	30.78098869	20.33044052	10.45055	-0.64262	1.561165		
Mala	33.46582413	21.64178658	11.82404	0.730866	0.602542		
Male	32.90471649	21.69516945	11.20955	0.116375	0.922502		
	29.86737442	18.9788208	10.88855	-0.20462	1.152381		
		Mean ΔCt (control)	11.09317				
	Fold gene expression						
			Standa	rd deviation	0.403267		
		St	andard error	of the mean	0.201634		
		Relativ	e standard er	ror of mean	19.02837		
	36.00477982	20.74038124	15.2644	4.171227	0.055505		
	37.58100128	23.71830177	13.8627	2.769528	0.146652		
Familia	34.07191467	22.45113373	11.62078	0.527609	0.693703		
Female							
	33.37465286	21.98809242	11.38656	0.293389	0.815983		
	33.24389648	20.23757744	13.00632	1.913147	0.265513		
			Fold gene	expression	0.395471		
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean							

Table S.3.5: Cyp2b1 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2b1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	30.08128357	21.44843102	8.632853	-0.05513	1.038951		
	29.82489204	20.36410713	9.460785	0.772805	0.585279		
Male							
iviale	31.02060699	21.1464119	9.874195	1.186215	0.439454		
	28.02316475	20.36148262	7.661682	-1.0263	2.036791		
	29.00507355	21.19468689	7.810387	-0.87759	1.837308		
	Mean ΔCt (control) 8.68798						
			Fold gene	expression	1.187557		
			Standa	rd deviation	0.722471		
		St	andard error	of the mean	0.323099		
		Relativ	e standard er	ror of mean	27.20702		
	30.06482315	20.15422821	9.910595	1.222615	0.428505		
	29.56221008	21.36679268	8.195417	-0.49256	1.406942		
Female	29.68480873	20.80569077	8.879118	0.191138	0.875915		
remaie	28.9195137	20.15905762	8.760456	0.072476	0.951005		
	28.98249245	21.34286499	7.639627	-1.04835	2.068167		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.3.6: Cyp2b2 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2b2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	27.84736252	22.29681969	5.550543	-0.92456	1.898103		
	27.30856323	21.2739296	6.034634	-0.44047	1.357044		
Male	24 02070720	22.47000224	0.50005	2.002004	0.224240		
	31.03978729 26.5728817	22.47080231 21.93065071	8.568985 4.642231	2.093884 -1.83287	0.234249 3.562451		
	29.60146141	22.0223484	7.579113	1.104012	0.465221		
	23.00110111	Mean ΔCt (control)	6.475101	1.10 1012	0.103221		
	Fold gene expression						
			Standa	ard deviation	1.332803		
		St	andard error	of the mean	0.596048		
		Relativ	e standard e	rror of mean	39.6463		
	21.98049355	21.44104385	0.53945	-5.93565	61.20813		
	28.75837898	24.35465431	4.403725	-2.07138	4.202875		
Female							
	00.00044407		0.674.770	0.000.00	10.00000		
	26.60241127	23.930933	2.671478	-3.80362	13.96383		
	22.37498474	22.16269875	0.212286	-6.26282	76.78833 39.04079		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.3.7: Cyp2c6 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	31.19462585	21.2739296	9.920696	-0.40614	1.325137		
Mala	33.10131073	22.35676956	10.74454	0.417703	0.748616		
Male	32.9033165	22.47080231	10.43251	0.105676	0.929369		
	33.62343597	21.93065071	11.69279	1.365947	0.38798		
	31.19462585 21.2739296 9.920696 -0.40614 33.10131073 22.35676956 10.74454 0.417703 32.9033165 22.47080231 10.43251 0.105676 33.62343597 21.93065071 11.69279 1.365947 30.86600304 22.0223484 8.843655 -1.48318 Mean ΔCt (control) 10.32684 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 32.25588226 21.44104385 10.81484 0.488 32.7257843 24.35465431 8.37113 -1.95571 32.53884125 24.16374207 8.375099 -1.95174	2.79565					
		Mean ΔCt (control)	10.32684				
			Standa	rd deviation	0.934231		
			Standard error	of the mean	0.417801		
		Rela	ative standard er	ror of mean	33.76575		
	32.25588226	21.44104385	10.81484	0.488	0.713013		
	32.7257843	24.35465431	8.37113	-1.95571	3.879063		
Family	32.53884125	24.16374207	8.375099	-1.95174	3.868406		
Female	30.6821804	21.61458778	9.067593	-1.25925	2.393706		
	33.75783157	23.930933	9.826899	-0.49994	1.414154		
	32.98618317	21.2739296 9.920696 -0.40614 22.35676956 10.74454 0.417703 22.47080231 10.43251 0.105676 21.93065071 11.69279 1.365947 22.0223484 8.843655 -1.48318 Mean ΔCt (control) 10.32684 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 21.44104385 10.81484 0.488 24.35465431 8.37113 -1.95571 24.16374207 8.375099 -1.95174 21.61458778 9.067593 -1.25925 23.930933 9.826899 -0.49994 22.16269875 10.82348 0.496646 Fold gene expression Standard deviation Standard deviation	0.708753				
			Fold gene	expression	2.162849		
	Standard deviation						
			Standard error	of the mean	0.596785		
		Rela	ative standard er	ror of mean	27.59255		

Table S.3.8: Cyp2c11 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c11</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	23.4070015	20.84217834	2.564823	-1.30236	2.466327	
	26.0033474	20.97036934	5.032978	1.165791	0.44572	
Mala	26.1450634	21.33662796	4.808435	0.941248	0.520782	
iviale	25.41716003	22.14450264	3.272657	-0.59453	1.509981	
	24.84078979	21.18374634	3.657043	-0.21014	1.156804	
	23.4070015 20.84217834 2.564823 -1.30236 26.0033474 20.97036934 5.032978 1.165791 26.1450634 21.33662796 4.808435 0.941248 25.41716003 22.14450264 3.272657 -0.59453 24.84078979 21.18374634 3.657043 -0.21014					
		Mean ΔCt (control)	3.867188			
Fold gene expression						
			Standa	rd deviation	0.82611	
			Standard error	of the mean	0.369448	
		Relat	tive standard er	ror of mean	30.2845	
	29.71196556	20.82183647	8.890129	5.022942	0.030757	
	30.10930252	23.80994415	6.299358	2.432171	0.185286	
Fomala	29.63360596	23.01904488	6.614561	2.747374	0.148922	
remale	28.98827362	20.93088341	8.05739	4.190203	0.05478	
	29.81073952	21.30577278	8.504967	4.637779	0.040169	
			Fold gene	expression	0.091983	
	Standard deviation					
	Standard error of the mean					
		Relat	tive standard er	ror of mean	34.17637	

Table S.3.9: Cyp2c23 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.03046227	19.53005981	-1.4996	0.490877	0.711592		
	18.16113281	19.37013245	-1.209	0.781475	0.581772		
Male	18.17616272	19.68417549	-1.50801	0.482462	0.715755		
iviale	17.42055702	20.35167122	-2.93111	-0.94064	1.919378		
	17.90238953	20.21253204	-2.31014	-0.31967	1.248043		
	17.92873955	20.41372299	-2.48498	-0.49451	1.408841		
	Mean ΔCt (control) -1.99048						
	Fold gene expression						
			Standa	ard deviation	0.520713		
		St	andard error	of the mean	0.21258		
		Relativ	e standard e	rror of mean	19.36837		
	17.5458107	20.22820854	-2.6824	-0.69192	1.615435		
	18.24902916	21.51779556	-3.26877	-1.27829	2.425515		
Familia	17.90238953 20.21253204 -2.31014 17.92873955 20.41372299 -2.48498 Mean ΔCt (control) -1.99048 Fold gene Standard Standard error of Relative standard error of Relative standard error of 18.24902916 21.51779556 -3.26877 17.60991669 21.09210587 -3.48219 17.00697899 20.30070305 -3.29372	-1.49171	2.812229				
Female	17.00697899	20.30070305	-3.29372	0.490877 0.781475 1 0.482462 1 -0.94064 4 -0.31967 8 -0.49451 8 gene expression indard deviation iror of the mean 4 -0.69192 7 -1.27829 9 -1.49171 2 -1.30325 3 -1.37056 5 -1.11287 gene expression indard deviation iror of the mean	2.46784		
	18.69623375	22.05726624	-3.36103	-1.37056	2.585705		
	16.89379501	19.99714088	-3.10335	-1.11287	2.162756		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
Relative standard error of mean							

Table S.3.10: *Cyp2d2* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2d2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	26.99505615	21.44843102	5.546625	-0.98519	1.979571		
	26.5629406	20.36410713	6.198833	-0.33298	1.259612		
Mala	28.65539932	20.92403412	7.731365	1.199553	0.43541		
Male	27.98278046	21.1464119	6.836369	0.304556	0.809691		
	26.58566856	20.36148262	6.224186	-0.30763	1.23767		
	27.84818459	21.19468689	6.653498	0.121685	0.919114		
	Mean ΔCt (control) 6.531813						
			Fold gen	e expression	1.106845		
			Standa	ard deviation	0.524685		
		S	tandard error	of the mean	0.214202		
		Relati	ive standard e	rror of mean	19.35246		
	27.44242477	20.15422821	7.288197	0.756384	0.591978		
	30.07309914	Society Soc	0.22152				
Female	28.77518654	20.80569077	7.969496	1.437683	0.36916		
remaie	27.63694572	20.15905762	7.477888	-0.98519 -0.33298 1.199553 0.304556 -0.30763 0.121685 ne expression dard deviation or of the mean 0.756384 2.174494 1.437683 0.946075 1.251892 0.742731 ne expression dard deviation or of the mean	0.519042		
	29.49430084	21.71059608	7.783705	1.251892	0.419897		
	28.61740875	21.34286499	7.274544	0.742731	0.597607		
	Fold gene expression						
Standard deviation							
		S	tandard error	of the mean	0.059525		
		Relati	ive standard e	rror of mean	13.13429		

Table S.3.11: *Cyp2d3* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2d3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	32.96843719	20.36410713	12.60433	-0.97505	1.965714	
Male	34.16418839	20.92403412	13.24015	-0.33923	1.265081	
iviale	35.26021194	21.1464119	14.1138	0.534416	0.690438	
	34.72073364	20.36148262	14.35925	0.779867	0.58242	
	Mean ΔCt (control) 13.57938					
	Fold gene expression					
			Standa	rd deviation	0.634995	
			Standard error	of the mean	0.317498	
		Relat	tive standard er	ror of mean	28.19911	
	37.07509995	20.15422821	16.92087	3.657394	0.079253	
	35.50923157	21.36679268	14.14244	0.878961	0.543759	
Female	35.92437744	20.15905762	15.76532	2.501842	0.176551	
	37.17380142	21.34286499	15.83094	2.567458	0.168701	
			Fold gene	e expression	0.242066	
	Standard deviation					
	Standard error of the mean					
		Relat	tive standard er	ror of mean	42.53255	

Table S.3.12: *Cyp2d4* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2d4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	27.71310043	19.53005981	8.183041	0.585552	0.666394		
	26.88541222	19.37013245	7.51528	-0.08221	1.058637		
Mala	27.67696953	19.68417549	7.992794	0.395306	0.760328		
iviale	27.85234642	20.35167122	7.500675	-0.09681	1.069409		
	27.44627571	20.21253204	7.233744	-0.36374	1.286762		
	27.57312012	20.41372299	7.159397	-0.43809	1.354811		
		Mean ΔCt (control)	7.597488				
			Fold gene	expression	1.032723		
			Standar	d deviation	0.275205		
		S	tandard error o	f the mean	0.112352		
		Relati	ve standard err	or of mean	10.87919		
	28.45509529	20.22820854	8.226887	0.629398	0.646446		
	31.38321114	21.51779556	9.865416	2.267927	0.207628		
Family	30.03599167	21.09210587	8.943886	1.346397	0.393273		
remaie	28.57620811	20.30070305	8.275505	0.678017	0.625024		
	Male 27.71310043 19.53005981 8.183041 0.585552 26.88541222 19.37013245 7.51528 -0.08221 27.67696953 19.68417549 7.992794 0.395306 27.85234642 20.35167122 7.500675 -0.09681 27.44627571 20.21253204 7.233744 -0.36374 27.57312012 20.41372299 7.159397 -0.43809 Mean ΔCt (control) 7.597488 Fold gene expression Standard error of the mean Relative standard error of mean Relative standard error of mean 31.38321114 21.51779556 9.865416 2.267927 30.03599167 21.09210587 8.943886 1.346397 28.57620811 20.30070305 8.275505 0.678017 30.57167816 22.05726624 8.514412 0.916924 29.08075142 19.99714088 9.083611 1.486122 Fold gene expression Standard deviation Standard deviation	0.916924	0.529637				
	29.08075142	19.99714088	9.083611	1.486122	0.356971		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean							

Table S.3.13: Cyp2e1 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2e1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.97686958	19.53005981	-0.55319	0.515709	0.699449		
	18.88059616	19.37013245	-0.48954	0.579363	0.669259		
Male	18.84788704	19.68417549	-0.83629	0.23261	0.851094		
iviale	18.32757378	20.35167122	-2.0241	-0.9552	1.938847		
	19.27022934	20.21253204	-0.9423	0.126596	0.91599		
	18.84574509	20.41372299	-1.56798	-0.49908	1.413311		
	Mean ΔCt (control) -1.0689						
			Fold gene	expression	1.081325		
			Standar	d deviation	0.498303		
		Sta	ndard error o	f the mean	0.203431		
		Relative	standard err	or of mean	18.81314		
	17.8197155	20.22820854	-2.40849	-1.33959	2.530801		
	18.68653679	21.51779556	-2.83126	-1.76236	3.392526		
Familia	17.32378769	21.09210587	-3.76832	-2.69942	6.495404		
Female	17.24805641	20.30070305	-3.05265	-1.98375	3.955192		
	18.87420464	22.05726624	-3.18306	-2.11416	4.329387		
	17.29385567	19.99714088	-2.70329	-1.63439	3.104555		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean					14.30133		

Table S.3.14: *Cyp2j3* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.78109169	19.53005981	5.251032	0.51916	0.697778	
	24.33361626	19.37013245	4.963484	0.231612	0.851683	
Male	24.43241119	19.68417549	4.748236	0.016363	0.988722	
iviale	25.01156425	20.35167122	4.659893	-0.07198	1.051158	
	24.53747749	20.21253204	4.324945	-0.40693	1.325858	
	24.85736656	20.41372299	4.443644	-0.28823	1.22114	
Mean ΔCt (control) 4.731872						
Fold gene expression						
			Standa	rd deviation	0.231546	
		Sta	ndard error	of the mean	0.094528	
		Relative	e standard er	ror of mean	9.24282	
	24.15125465	20.22820854	3.923046	-0.80883	1.751785	
	25.74804497	21.51779556	4.230249	-0.50162	1.415805	
Female	25.7058506	21.09210587	4.613745	-0.11813	1.085325	
	25.60566711	20.30070305	5.304964	0.573092	0.672175	
	25.86975861	22.05726624	3.812492	-0.91938	1.891302	
			Fold gene	expression	1.363279	
	Standard deviation					
	Standard error of the mean					
		Relative	e standard er	ror of mean	16.29838	

Table S.3.15: *Cyp2j4* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	25.71083069	21.44843102	4.2624	0.138124	0.9087		
	24.42203903	20.36410713	4.057932	-0.06634	1.04706		
Mala	24.92321587	20.92403412	3.999182	-0.12509	1.090579		
Male	25.47284126	21.1464119	4.326429	0.202153	0.869252		
	25.20201683	20.36148262	4.840534	0.716258	0.608674		
	24.45386696	21.19468689	3.25918	-0.8651	1.821461		
	Mean ΔCt (control) 4.124276						
Fold gene expression							
			Standa	rd deviation	0.410819		
		S	tandard error	of the mean	0.167716		
		Relati	ve standard er	ror of mean	15.85788		
	25.36391258	20.15422821	5.209684	1.085408	0.471259		
	26.14167976	21.36679268	4.774887	0.650611	0.637011		
Famala	25.95587158	20.80569077	5.150181	1.025905	0.491102		
Female	25.59830093	20.15905762	5.439243	1.314967	0.401935		
	26.42259979	21.71059608	4.712004	0.587728	0.66539		
			Fold gene	expression	0.533339		
	Standard deviation						
		S	tandard error	of the mean	0.050542		
Relative standard error of mean							

Table S.3.16: Cyp2j10 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.65837097	21.44843102	13.20994	-0.63978	1.558089	
	33.91353989	20.36410713	13.54943	-0.30029	1.231388	
Mala	35.96679306	20.92403412	15.04276	1.193041	0.43738	
iviale	34.63116074	21.1464119	13.48475	-0.36497	1.287854	
	34.45147324	20.36148262	14.08999	0.240273	0.846585	
	34.91612244	21.19468689	13.72144	-0.12828	1.092992	
		Mean ΔCt (control)	13.84972			
			Fold gene	expression	1.075715	
			Standa	rd deviation	0.390434	
		S	tandard error	of the mean	0.159394	
		Relati	ive standard er	ror of mean	14.81749	
	34.55791092	20.15422821	14.40368	0.553965	0.681146	
	33.5796051	21.36679268	12.21281	-1.63691	3.10998	
Famala	Male 34.65837097 21.44843102 13.20994 -0.63978 33.91353989 20.36410713 13.54943 -0.30029 35.96679306 20.92403412 15.04276 1.193041 34.63116074 21.1464119 13.48475 -0.36497 34.45147324 20.36148262 14.08999 0.240273 34.91612244 21.19468689 13.72144 -0.12828 Mean ΔCt (control) 13.84972 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 34.55791092 20.15422821 14.40368 0.553965	1.312829				
remale	34.64805603	20.15905762	14.489	0.639281	0.642033	
	34.03968811	21.34286499	12.69682	-1.15289	2.223596	
			Fold gene	expression	1.593917	
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					29.80308	

Table S.3.17: *Cyp3a1* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp3a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.30351257	18.7430172	15.5605	-0.79485	1.734902	
	38.3971138	21.22470284	17.17241	0.817062	0.567597	
Male	37.59463882	21.17545128	16.41919	0.063838	0.956715	
	36.89120865	19.68737793	17.20383	0.848481	0.555369	
	34.52952957	19.10870743	15.42082	-0.93453	1.911264	
		Mean ΔCt (control)	16.35535			
			Fold ge	ne expression	1.145169	
			Stand	lard deviation	0.642584	
			Standard erro	r of the mean	0.287372	
		Rela	tive standard	error of mean	25.09429	
	37.00455475	20.61572647	16.38883	0.033479	0.977061	
	34.30351257 18.7430172 15.5605 -0.79485 38.3971138 21.22470284 17.17241 0.817062 37.59463882 21.17545128 16.41919 0.063838 36.89120865 19.68737793 17.20383 0.848481 34.52952957 19.10870743 15.42082 -0.93453 Mean ΔCt (control) 16.35535 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 37.00455475 20.61572647 16.38883 0.033479					
Female	37.08632278	3.30351257 18.7430172 15.5605 -0.79485 1 3.3971138 21.22470284 17.17241 0.817062 0 5.59463882 21.17545128 16.41919 0.063838 0 8.89120865 19.68737793 17.20383 0.848481 0 5.52952957 19.10870743 15.42082 -0.93453 1 Mean ΔCt (control) 16.35535 Fold gene expression 1 Standard deviation 0 Standard error of the mean 2 Relative standard error of mean 2 0.00455475 20.61572647 16.38883 0.033479 0 0.08632278 21.51637077 15.56995 -0.7854 1 0.9864006 19.91755867 16.06884 -0.28651 1 0.33658638 22.14803505 16.21783 -0.13752 1 0.46936035 21.31115723 16.1582 -0.19715 1 0.54006 Standard deviation 0 0.55006 Standard deviation 0 0.55006 Standard deviation 0 0.55006 Standard error of the mean 0 0.55006 Standard deviation 0 0.55006 Standard error of the mean 0	1.723567			
remale	35.9864006	19.91755867	16.06884	-0.28651	1.219684	
	38.3658638	22.14803505	16.21783	-0.13752	1.100013	
	37.46936035	21.31115723	16.1582	-0.19715	1.146428	
			Fold ge	ne expression	1.233351	
	Standard deviation					
			Standard erro	r of the mean	0.128747	
Relative standard error of mean						

Table S.3.18: *Cyp3a2* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp3a2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	34.46009445	18.7430172	15.71708	1.488748	0.356322		
Male	36.56611633	21.22470284	15.34141	1.113084	0.462305		
iviale	34.27514267	21.17545128	13.09969	-1.12864	2.186522		
	32.79494476	19.68737793	13.10757	-1.12076	2.174618		
	32.98460388	19.10870743	13.8759	-0.35243	1.276712		
	Mean ΔCt (control) 14.22833						
			Fold gene	expression	1.291296		
			Standard	d deviation	0.886468		
		Star	ndard error o	f the mean	0.396441		
		Relative	standard err	or of mean	30.701		
	33.75891113	22.99111748	10.76779	-3.46054	11.00842		
	33.00808334	21.51637077	11.49171	-2.73662	6.665054		
Female	33.25848007	19.91755867	13.34092	-0.88741	1.849849		
	33.80696487	21.31115723	12.49581	-1.73252	3.323081		
			Fold gene	expression	5.711601		
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean					35.58936		

Table S.3.19: Cyp3a9 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a9</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.0511322	19.86431313	14.18682	0.35606	0.781295	
Mala	34.71797943	20.96670532	13.75127	-0.07948	1.05664	
Male	35.27788544	21.0174675	14.26042	0.429659	0.742437	
	33.58153915	20.51798058	13.06356	-0.7672	1.701963	
	33.46962357	19.57789993	13.89172	0.060965	0.958623	
		Mean ΔCt (control)	13.83076			
			Fold gene	expression	1.048192	
			Standa	rd deviation	0.387373	
		St	andard error	of the mean	0.173238	
		Relativ	e standard er	ror of mean	16.52736	
	35.39659882	20.31739998	15.0792	1.24844	0.420903	
200	34.45935822	21.91688538	12.54247	-1.28829	2.442377	
Female	31.73126602	19.66669464	12.06457	-1.76619	3.401538	
	37.22182846	22.88987923	14.33195	0.501191	0.706523	
	34.31126785	20.18359756	14.12767	0.296912	0.813993	
			Fold gene	expression	1.557067	
			Standa	rd deviation	1.299259	
		St	andard error	of the mean	0.581046	
	Relative standard error of mean					

Table S.3.20: *Cyp3a23* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	30.25981331	19.86431313	10.3955	-0.28118	1.215184	
Male	33.15713882	20.96670532	12.19043	1.513758	0.350198	
iviale	32.32994461	21.0174675	11.31248	0.635802	0.643583	
	30.07069588	20.51798058	9.552715	-1.12396	2.179444	
	29.51015091	19.57789993	9.932251	-0.74442	1.675306	
		Mean ΔCt (control)	10.67668			
			Fold gene	expression	1.212743	
			Standard	d deviation	0.74438	
		Sta	ndard error o	f the mean	0.332897	
		Relative	standard err	or of mean	27.44992	
	30.75619888	20.31739998	10.4388	-0.23788	1.179256	
Formula	31.16948891	21.91688538	9.252604	-1.42407	2.683418	
Female	29.66583443	19.66669464	9.99914	-0.67754	1.599405	
	30.44386101	20.18359756	10.26026	-0.41641	1.334604	
			Fold gene	expression	1.699171	
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					19.97162	

Table S.3.21: *Cyp4a1* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	19.91963768	20.89357758	-0.97394	-0.74817	1.679659	
	19.9793663	19.86431313	0.115053	0.340825	0.78959	
na.t.	19.96125603	20.96670532	-1.00545	-0.77968	1.716747	
Male	21.54629517	21.0174675	0.528828	0.754599	0.592711	
	20.2638607	20.51798058	-0.25412	-0.02835	1.019844	
	19.81289864	19.57789993	0.234999	0.46077	0.726598	
		Mean ΔCt (control)	-0.22577			
			Fold gene	expression	1.087525	
			Standa	rd deviation	0.49295	
		St	andard error	of the mean	0.201246	
		Relativ	e standard er	ror of mean	18.50496	
	20.83655357	20.31739998	0.519154	0.744925	0.596699	
	22.4444294	22.59498405	-0.15055	0.075217	0.949199	
Formula	20.9527607	21.91688538	-0.96412	-0.73835	1.66827	
Female	20.85179329	19.66669464	1.185099	1.41087	0.376085	
	22.2225914	22.88987923	-0.66729	-0.44152	1.358031	
	20.75555611	20.18359756	0.571959	0.79773	0.575254	
			Fold gene	expression	0.92059	
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.3.22: *Cyp4a2* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	18.50077629	20.89357758	-2.3928	-0.35147	1.275862		
	17.91036987	19.86431313	-1.95394	0.087385	0.941227		
Mala	18.37908554	20.96670532	-2.58762	-0.54629	1.460327		
Male	19.3446846	21.0174675	-1.67278	0.368546	0.774563		
	18.34843636	20.51798058	-2.16954	-0.12822	1.092941		
	18.10661888	19.57789993	-1.47128	0.570048	0.673595		
	Mean ΔCt (control) -2.04133						
			Fold gene	expression	1.036419		
			Standa	rd deviation	0.299832		
		Sta	ndard error	of the mean	0.122406		
		Relative	standard er	ror of mean	11.81048		
	24.86208725	20.31739998	4.544687	6.586016	0.010409		
	26.35299301	22.59498405	3.758009	5.799338	0.017957		
Famala							
Female	24.8263092	19.66669464	5.159615	7.200943	0.006797		
	26.93806648	22.88987923	4.048187	6.089516	0.014685		
	23.79533195	20.18359756	3.611734	5.653063	0.019873		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
Relative standard error of mean					17.2251		

Table S.3.23: *Cyp4a3* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	18.78629303	20.582304	-1.79601	0.139024	0.908133	
	18.39201546	19.87765503	-1.48564	0.449396	0.73235	
Male	18.79459953	20.79586792	-2.00127	-0.06623	1.046979	
ividie	19.45201492	21.4719944	-2.01998	-0.08494	1.060647	
	18.725914	21.09819221	-2.37228	-0.43724	1.354014	
		Mean ΔCt (control)	-1.93504			
			Fold gene	expression	1.020425	
			Standar	d deviation	0.228693	
		Stan	dard error o	f the mean	0.102275	
		Relative	standard err	or of mean	10.02274	
	19.34219742	20.18075752	-0.83856	1.096475	0.467658	
Female	20.04523468	22.27972412	-2.23449	-0.29945	1.230679	
remale	18.8917942	21.04664612	-2.15485	-0.21982	1.164586	
	20.49616814	23.84099388	-3.34483	-1.40979	2.656986	
	19.84767151	20.96439171	-1.11672	0.818315	0.567104	
			Fold gene	expression	1.217402	
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.3.24: *Cyp4a8* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a8</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	33.2581749	20.582304	12.67587	-1.64706	3.131948	
	35.19928741	19.87765503	15.32163	0.998701	0.50045	
Male	34.73405838	20.79586792	13.93819	-0.38474	1.305625	
iviale	38.2901268	21.4719944	16.81813	2.495201	0.177366	
	33.95902252	21.09819221	12.86083	-1.4621	2.755093	
		Mean ΔCt (control)	14.32293			
			Fold gene	expression	1.574097	
			Standar	d deviation	1.322621	
		Sta	ndard error o	f the mean	0.591494	
		Relative	standard err	or of mean	37.57673	
	34.92519379	24.4500885	10.47511	-3.84783	14.39829	
Female	32.1306572	22.27972412	9.850933	-4.472	22.19247	
Female	33.94862366	21.04664612	12.90198	-1.42095	2.677625	
	33.98335266	23.84099388	10.14236	-4.18057	18.13334	
	33.77131271	20.96439171	12.80692	-1.51601	2.85999	
			Fold gene	expression	12.05234	
	Standard deviation					
Standard error of the mean						
Relative standard error of mean						

Table S.3.25: *Cyp4f1* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	19.92054939	20.582304	-0.66175	-0.03506	1.024601		
	19.26861954	19.87765503	-0.60904	0.017657	0.987836		
Mala	21.16665077	20.79586792	0.370783	0.997475	0.500876		
ividie	20.55848312	21.4719944	-0.91351	-0.28682	1.219947		
	19.98826408	20.58220482	-0.59394	0.032752	0.977554		
	19.74549675	21.09819221	-1.3527	-0.726	1.65405		
		Mean ΔCt (control)	-0.62669				
			Fold gene	expression	1.060811		
			Standa	rd deviation	0.375327		
		St	andard error	of the mean	0.153227		
		Relativ	e standard er	ror of mean	14.4443		
	19.63010979	20.18075752	-0.55065	0.076045	0.948655		
	22.1161232	24.4500885	-2.33397	-1.70727	3.26543		
Famala	Male	2.359471					
Female	19.15427589	21.04664612	-0.66175 -0.03506 -0.60904 0.017657 0.370783 0.997475 -0.91351 -0.28682 -0.59394 0.032752 -1.3527 -0.726 Fold gene expression Standard deviation Standard error of the mean tive standard error of mean -0.55065 0.076045 -2.33397 -1.70727 -1.86516 -1.23846 -1.89237 -1.26568 -1.88016 -1.25347 -1.66775 -1.04106 Fold gene expression Standard deviation Standard error of the mean	-1.26568	2.404401		
	21.9608326	23.84099388	-1.88016	-1.25347	2.38414		
	19.2966404	20.96439171	-1.66775	-1.04106	2.057737		
	Fold gene expression						
Standard deviation					0.750241		
	Standard error of the mean						
Relative standard error of mean					13.69397		

Table S.3.26: Cyp4f4 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	29.40390396	20.84189224	8.562012	-1.03432	2.048142	
	31.44765091	20.75020409	10.69745	1.101119	0.466155	
Mala						
iviale	32.02010345	21.57505989	10.44504	0.848716	0.555279	
	29.77568245	20.87166214	8.90402	-0.69231	1.615866	
	30.52752113	21.15440559	9.373116	-0.22321	1.16733	
		Mean ΔCt (control)	9.596328			
			Fold gen	e expression	1.170554	
			Standa	ard deviation	0.678825	
		;	Standard error	of the mean	0.30358	
		Relat	tive standard e	rror of mean	25.93472	
	30.72869301	23.19133377	7.537359	-2.05897	4.166882	
Famala	Male 31.44765091 20.75020409 10.69745 1.101119 32.02010345 21.57505989 10.44504 0.848716 29.77568245 20.87166214 8.90402 -0.69231 30.52752113 21.15440559 9.373116 -0.22321 Mean ΔCt (control) 9.596328 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean	-2.0538	4.151969			
remale	29.30051613	20.96542549	8.335091	-1.26124	2.397012	
	28.56533051	20.82875443	7.736576	-1.85975	3.629451	
			Fold gen	e expression	3.586329	
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean						

Table S.3.27: *Cyp4f5* mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.50182152	22.29681969	2.205002	-0.45799	1.373626	
	24.18658447	21.2739296	2.912655	0.249664	0.841092	
0.0.1.	24.66133308	22.35676956	2.304564	-0.35843	1.282028	
Male	24.7171669	22.47080231	2.246365	-0.41663	1.334803	
	24.33512306	21.93065071	2.404472	-0.25852	1.19625	
	25.92723846	22.0223484	3.90489	1.241899	0.422816	
		Mean ΔCt (control)	2.662991			
			Fold gene	expression	1.075103	
			Standa	rd deviation	0.372638	
		St	andard error	of the mean	0.152129	
		Relativ	e standard er	ror of mean	14.15016	
	24.25989723	21.44104385	2.818853	0.155862	0.897596	
	25.4653492	24.35465431	1.110695	-1.5523	2.932836	
Female	24.92349815	24.16374207	0.759756	-1.90324	3.74051	
remaie						
	24.83056068	23.930933	0.899628	-1.76336	3.394887	
	24.45025826	22.16269875	2.28756	-0.37543	1.297228	
			Fold gene	expression	2.452611	
	Standard deviation					
		St	andard error	of the mean	0.571409	
Relative standard error of mean					23.29799	

Table S.3.28: Cyp4f6 mRNA expression in the kidney of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	23.98557663	22.29681969	1.688757	-0.41763	1.335733	
	23.92938042	21.2739296	2.655451	0.549062	0.683464	
Mala	25.51048088	22.35676956	3.153711	1.047323	0.483865	
iviale	23.90467453	22.47080231	1.433872	-0.67252	1.59385	
	23.51361084	21.93065071	1.58296	-0.52343	1.437367	
	24.14592743	22.0223484	2.123579	0.017191	0.988155	
		Mean ΔCt (control)	2.106388			
			Fold gene	expression	1.087072	
			Standard	d deviation	0.442245	
		Sta	ndard error o	f the mean	0.180546	
		Relative	standard err	or of mean	16.60846	
	23.30430603	21.44104385	1.863262	-0.24313	1.183555	
	24.89033699	24.35465431	0.535683	-1.57071	2.9705	
Famala	Male 23.92938042 21.2739296 2.655451 0.549062 (25.51048088 22.35676956 3.153711 1.047323 (23.90467453 22.47080231 1.433872 -0.67252 (23.51361084 21.93065071 1.58296 -0.52343 (24.14592743 22.0223484 2.123579 0.017191 (Mean ΔCt (control) 2.106388 Fold gene expression (Standard deviation (Standard error of the mean (Relative standard error of mean (Relative standard error of mean (23.30430603 21.44104385 1.863262 -0.24313 (24.89033699 24.35465431 0.535683 -1.57071 (24.3901062 24.16374207 0.226364 -1.88002 (23.67495537 21.61458778 2.060368 -0.04602 (25.06542778 23.930933 1.134495 -0.97189 (23.30242538 22.16269875 1.139727 -0.96666 (Fold gene expression (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (Standard deviation (3.680813				
remale	23.67495537	21.61458778	2.060368	-0.04602	1.032413	
	25.06542778	23.930933	1.134495	-0.97189	1.961413	
	23.30242538	22.16269875	1.139727	-0.96666	1.954313	
Fold gene expression						
Standard deviation					1.026762	
	Standard error of the mean					
		Relative	standard err	or of mean	19.67489	

S.4 Calculation of the fold change in the lung level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta CT$ method:

Table S.4.1: *Cyp1a1* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp1a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	22.92127228	17.87739372	5.043879	0.43672	0.738813	
Male	23.39588547	18.78137016	4.614515	0.007356	0.994914	
iviale	24.70204353	19.61618996	5.085854	0.478695	0.717627	
	22.8150177	18.69833565	4.116682	-0.49048	1.404909	
	23.2163372	19.04147148	4.174866	-0.43229	1.349377	
		Mean ΔCt (control)	4.607159			
			Fold gen	e expression	1.041128	
			Standa	ard deviation	0.326166	
		S	tandard error	of the mean	0.145866	
		Relati	ive standard e	rror of mean	14.01038	
	22.48679352	18.92680931	3.559984	-1.04717	2.066479	
	24.05761147	18.64676285	5.410849	0.80369	0.572882	
Female						
remale	21.62118721	18.72934532	2.891842	-1.71532	3.283688	
	23.25413704	18.08415794	5.169979	0.56282	0.676978	
	22.0242157	19.15018082	2.874035	-1.73312	3.32447	
			Fold gen	e expression	1.984899	
	Standard deviation					
	Standard error of the mean					
		Relati	ive standard e	rror of mean	30.21229	

Table S.4.2: Cyp1a2 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	26.82986832	17.87739372	8.952475	-0.14126	1.102865
Male	27.35612488	18.78137016	8.574755	-0.51898	1.432937
iviale	28.65088081	19.61618996	9.034691	-0.05904	1.041772
	27.89149475	18.69833565	9.193159	0.099429	0.933402
	28.75504303	19.04147148	9.713572	0.619841	0.650742
		Mean ΔCt (control)	9.09373		
			Fold gen	e expression	1.032344
			Standa	ard deviation	0.283261
		S	tandard error	of the mean	0.126678
		Relati	ive standard e	rror of mean	12.27091
Familia	26.82986832 17.87739372 8.952475 -0.14126 1 27.35612488 18.78137016 8.574755 -0.51898 1 28.65088081 19.61618996 9.034691 -0.05904 1 27.89149475 18.69833565 9.193159 0.099429 0 28.75504303 19.04147148 9.713572 0.619841 0 Mean ΔCt (control) 9.09373 Fold gene expression 1 Standard deviation Standard error of the mean 1 Relative standard error of mean 1 27.893013 18.55441666 9.338596 0.244866 0 29.1862278 18.72934532 10.45688 1.363152 0 28.41287041 18.08415794 10.32871 1.234982 0 28.88457108 19.15018082 9.73439 0.64066 0 Fold gene expression 0 Standard deviation 0 Standard deviation 0	0.843894			
Female	29.1862278	18.72934532	10.45688	1.363152	0.388732
	28.41287041	18.08415794	10.32871	1.234982	0.424848
	28.88457108	19.15018082	9.73439	0.64066	0.641419
			Fold gen	e expression	0.574723
Standard deviation					
		S	tandard error	of the mean	0.105655
		Relati	ive standard e	rror of mean	18.3837

Table S.4.3: Cyp1b1 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1b1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	22.38321114	17.87739372	4.505817	0.496854	0.70865		
	22.62660599	18.3462429	4.280363	0.271399	0.828515		
Male	23.49168205	18.78137016	4.710312	0.701348	0.614997		
iviale	23.34890366	19.61618996	3.732714	-0.27625	1.211043		
	22.3320179	18.69833565	3.633682	-0.37528	1.297092		
	22.23236465	19.04147148	3.190893	-0.81807	1.763046		
		Mean ΔCt (control)	4.008964				
			Fold gene	expression	1.070557		
			Standa	rd deviation	0.43536		
		St	andard error	of the mean	0.177735		
		Relativ	e standard er	ror of mean	16.60209		
	23.1882515	18.92680931	4.261442	0.252479	0.839453		
	22.27731323	18.64676285	3.63055	-0.37841	1.299911		
Female	22.54185867	18.55441666	3.987442	-0.02152	1.015029		
remale	23.95024681	18.72934532	5.220901	1.211938	0.431688		
	22.37326431	18.08415794	4.289106	0.280143	0.82351		
	22.13584709	19.15018082	2.985666	-1.0233	2.032559		
	Fold gene expression						
	Standard deviation						
		St	andard error	of the mean	0.22388		
		Relativ	e standard er	ror of mean	20.85144		

Table S.4.4: Cyp2b1 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2b1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	19.55262947	18.99743843	0.555191	0.571249	0.673034		
	19.74834633	19.06721306	0.681133	0.697191	0.616772		
Male	18.61517525	19.44244194	-0.82727	-0.81121	1.754681		
iviale	19.68612289	20.28934479	-0.60322	-0.58716	1.502291		
	19.668993	19.13036537	0.538628	0.554686	0.680805		
	18.90013123	19.34094238	-0.44081	-0.42475	1.342343		
		Mean ΔCt (control)	-0.01606				
			Fold gene	expression	1.094988		
			Standar	d deviation	0.498108		
		Sta	ndard error o	f the mean	0.203352		
		Relative	standard err	or of mean	18.57115		
	18.47485161	19.32472038	-0.84987	-0.83381	1.782387		
	17.61610985	19.25806427	-1.64195	-1.6259	3.086339		
Female	19.55262947 18.99743843 0.555191 0.571249 0.571249 1.574834633 19.06721306 0.681133 0.697191 0.571249 1.574834633 19.06721306 0.681133 0.697191 0.571249 1.574834633 19.06721306 0.681133 0.697191 0.571249 1.574834633 19.44244194 -0.82727 -0.81121 1.574834639 19.44244194 -0.82727 -0.60322 -0.58716 1.57483479	1.859925					
Female	17.70389557	19.70411301	-2.00022	-1.98416	3.956321		
	18.18362617	19.19081879	-1.00719	-0.99113	1.987748		
	18.32147217	19.74034691	-1.41887	-1.40282	2.644173		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relative	standard err	or of mean	13.67445		

Table S.4.5: Cyp2b2 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2b2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	29.08254433	18.99743843	10.08511	-0.55304	1.46717		
	30.22611427	19.06721306	11.1589	0.520759	0.697005		
D.AI	31.10868835	19.44244194	11.66625	1.028104	0.490354		
Male	30.29436302	20.28934479	10.00502	-0.63312	1.550919		
	29.23172569	19.13036537	10.10136	-0.53678	1.450733		
	30.153162	19.34094238	10.81222	0.174078	0.886334		
	-	Mean ΔCt (control)	10.63814				
			Fold gene	expression	1.090419		
			Standa	rd deviation	0.456143		
		St	andard error	of the mean	0.18622		
		Relativ	e standard er	ror of mean	17.07779		
	29.84101105	19.32472038	10.51629	-0.12185	1.08813		
	28.76152992	19.25806427	9.503466	-1.13468	2.195693		
Female	29.84632683	19.08437729	10.76195	11 -0.55304 29 0.520759 25 1.028104 02 -0.63312 36 -0.53678 22 0.174078 14 29 ene expression 29 -0.12185 36 -1.13468 39 0.123808 78 -0.36036 39 0.918844 74 -1.05847 gene expression andard deviation aror of the mean	0.917762		
Female	29.98189545	19.70411301	10.27778	-0.36036	1.283746		
	30.74780464	19.19081879	11.55699	0.918844	0.528933		
	29.32002068	19.74034691	9.579674	-1.05847	2.082719		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relativ	e standard er	ror of mean	20.00296		

Table S.4.6: Cyp2c6 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	32.79861832	19.06721306	13.73141	0.733075	0.60162		
Mala	33.62251663	19.44244194	14.18007	1.181745	0.440818		
Male	32.11355591	20.28934479	11.82421	-1.17412	2.25655		
	32.25492859	19.13036537	13.12456	0.126233	0.91622		
	31.47233772	19.34094238	12.1314	-0.86693	1.823784		
		Mean ΔCt (control)	12.99833				
			Fold gene	expression	1.207798		
			Standard	d deviation	0.793733		
		Star	ndard error o	f the mean	0.354968		
		Relative	standard err	or of mean	29.38968		
	32.89268875	19.32472038	13.56797	0.569638	0.673786		
	33.08142853	19.08437729	13.99705	0.998721	0.500443		
Female	32.06086349	19.70411301	12.35675	-0.64158	1.560036		
	32.49713135	19.19081879	13.30631	0.307983	0.807771		
	31.85897255	19.74034691	12.11863	-0.8797	1.839998		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean					24.42954		

Table S.4.7: Cyp2c11 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c11</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	29.29161263	18.99743843	10.29417	-0.71718	1.643963		
	30.40301704	19.06721306	11.3358	0.324452	0.798601		
Mala	31.34239197	19.44244194	11.89995	0.888598	0.540139		
iviale	30.64749336	20.28934479	10.35815	-0.6532	1.572656		
	30.34089088	19.13036537	11.21053	0.199174	0.871049		
	30.3104496	19.34094238	10.96951	-0.04184	1.029429		
		Mean ΔCt (control)	11.01135				
			Fold gene	expression	1.075973		
			Standa	rd deviation	0.442158		
		St	andard error	of the mean	0.18051		
		Relativ	e standard er	ror of mean	16.77648		
	30.63634872	19.32472038	11.31163	0.300277	0.812097		
	30.25811386	19.25806427	11.00005	-0.0113	1.007865		
Famala	29.29161263 18.99743843 10.29417 -0.71718 30.40301704 19.06721306 11.3358 0.324452 31.34239197 19.44244194 11.89995 0.888598 30.64749336 20.28934479 10.35815 -0.6532 30.34089088 19.13036537 11.21053 0.199174 30.3104496 19.34094238 10.96951 -0.04184 Mean ΔCt (control) 11.01135 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 30.63634872 19.32472038 11.31163 0.300277 30.25811386 19.25806427 11.00005 -0.0113 30.10378265 19.08437729 11.01941 0.008054 30.08021355 19.70411301 10.3761 -0.63525 30.51275253 19.19081879 11.32193 0.310582 Fold gene expression Standard deviation Standard error of the mean Standard e	0.994433					
remale	30.08021355	19.70411301	10.3761	-0.63525	1.553208		
	30.51275253	19.19081879	11.32193	0.310582	0.806316		
	Fold gene expression						
Standard deviation							
Standard error of the mean							
		Relativ	e standard er	ror of mean	13.19574		

Table S.4.8: Cyp2c23 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	22.77787781	18.38732338	4.390554	0.014815	0.989784		
	22.6652565	18.35128784	4.313969	-0.06177	1.043746		
Mala	22.53532028	18.76119232	3.774128	-0.60161	1.517411		
Ividie							
	23.10972023	18.19241142	4.917309	0.541569	0.687023		
	22.97971916	18.49698067	4.482738	0.106999	0.928518		
		Mean ΔCt (control)	4.37574				
			Fold gene	expression	1.033296		
			Standa	rd deviation	0.303007		
		Sta	ndard error	of the mean	0.135509		
		Relative	e standard er	ror of mean	13.11423		
	22.82199287	18.6565609	4.165432	-0.21031	1.156935		
	22.31262016	18.44511795	3.867502	-0.50824	1.422311		
Fomolo	Male	1.275769					
remaie	22.8443737	18.43808937	4.406284	0.030545	0.979051		
	22.54894447	17.74411392	4.804831	0.429091	0.74273		
	22.94019127	18.40876198	4.531429	0.15569	0.897703		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.4.9: Cyp2d4 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2d4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	26.43734169	18.38732338	8.050018	-0.62095	1.537885		
	26.43478775	18.35128784	8.0835	-0.58747	1.502606		
Male	27.03805542	18.76119232	8.276863	-0.3941	1.314126		
iviale	29.17103577	19.52798462	9.643051	0.972085	0.509769		
	27.7983799	18.49698067	9.301399	0.630433	0.645983		
		Mean ΔCt (control)	8.670966				
			Fold gene	expression	1.102074		
			Standa	rd deviation	0.488408		
		St	andard error	of the mean	0.218423		
		Relativ	e standard er	ror of mean	19.81923		
	27.12688637	18.6565609	8.470325	-0.20064	1.149209		
	26.79421997	18.44511795	8.349102	-0.32186	1.249945		
Female							
remale	30.28576851	18.43808937	11.84768	3.176713	0.11059		
	29.29802132	18.40876198	10.88926	2.218293	0.214895		
	Fold gene expression						
Standard deviation							
Standard error of the mean							
		Relativ	e standard er	ror of mean	44.15533		

Table S.4.10: Cyp2e1 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2e1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	21.81419182	18.57818794	3.236004	-0.24256	1.183093		
	21.84495544	18.38136482	3.463591	-0.01498	1.010435		
8.6.1.	21.95317459	18.97838593	2.974789	-0.50378	1.417923		
Male	22.49595833	19.63236046	2.863598	-0.61497	1.531526		
	22.26360512	18.45039368	3.813211	0.334644	0.79298		
	23.04731178	18.52709961	4.520212	1.041645	0.485773		
	'	Mean ΔCt (control)	3.478567				
			Fold gene	expression	1.070288		
			Standa	rd deviation	0.392034		
		St	andard error	of the mean	0.160047		
		Relativ	e standard er	ror of mean	14.95367		
	21.80897331	18.50517273	3.303801	-0.17477	1.128782		
	21.83387566	18.7165184	3.117357	-0.36121	1.284503		
Female							
remaie	21.72060013	18.3507061	3.369894	-0.10867	1.078236		
	21.21471214	17.98646927	3.228243	-0.25032	1.189475		
	22.30658913	18.20016098	4.106428	0.627861	0.647135		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relativ	e standard er	ror of mean	10.33382		

Table S.4.11: *Cyp2j3* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	24.35410118	16.40213203	7.951969	-0.57328	1.487906		
	24.3428669	15.77114773	8.571719	0.046467	0.968305		
Mala	24.99538231	16.00558853	8.989794	0.464542	0.724701		
iviale	25.91841888	17.14575195	8.772667	0.247415	0.842405		
	25.24705696	16.45843124	8.788626	0.263374	0.833137		
	24.42102814	16.34428978	8.076738	-0.44851	1.364634		
		Mean ΔCt (control)	8.525252				
			Fold gen	e expression	1.036848		
			Standa	ard deviation	0.313795		
		S	tandard error	of the mean	0.128106		
		Relati	ive standard e	rror of mean	12.35536		
	24.85449982	16.40930367	8.445196	-0.08006	1.057059		
24.3428669 15.77 24.99538231 16.00 25.91841888 17.14 25.24705696 16.45 24.42102814 16.34 Mean Δ Female 24.85449982 16.40 24.52044868 16.24 24.24489975 16.33 24.82716179 16.53 23.9208622 15.90	16.24884605	8.271603	-0.25365	1.192219			
Famala	24.24489975	16.33191109	7.912989	-0.61226	1.528656		
remale	24.82716179	16.53628731	8.290874	-0.23438	1.176399		
	23.9208622	15.90371037	8.017152	-0.5081	1.422176		
	24.68640137	16.67467499	8.011726	-0.51353	1.427535		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
		Relati	ive standard e	rror of mean	5.777114		

Table S.4.12: *Cyp2j4* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.37016296	18.57818794	5.791975	-0.41707	1.335216	
	24.35871696	18.38136482	5.977352	-0.2317	1.174214	
Mala	25.20401001	18.97838593	6.225624	0.016576	0.988576	
Male	25.64874458	19.63236046	6.016384	-0.19266	1.142872	
	25.43470383	18.45039368	6.98431	0.775262	0.584282	
	24.78574181	18.52709961	6.258642	0.049594	0.966208	
	•	Mean ΔCt (control)	6.209048			
			Fold gene	expression	1.031895	
			Standa	rd deviation	0.257341	
		St	andard error	of the mean	0.105059	
		Relativ	e standard er	ror of mean	10.18118	
	25.07962608	18.50517273	6.574453	0.365405	0.776251	
	24.46627235	18.7165184	5.749754	-0.45929	1.374869	
F	24.27939606	18.7470932	5.532303	-0.67675	1.598529	
Female	25.50915337	18.3507061	7.158447	0.949399	0.517848	
	23.99829292	17.98646927	6.011824	-0.19722	1.14649	
	24.61742401	18.20016098	6.417263	0.208215	0.865608	
Fold gene expression						
Standard deviation						
	Standard error of the mean					
Relative standard error of mean						

Table S.4.13: *Cyp2j10* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	30.84857368	16.40213203	14.44644	-0.12772	1.092568		
	30.40653038	15.77114773	14.63538	0.061218	0.958454		
Mala	30.69090843	16.00558853	14.68532	0.111156	0.925846		
Male	31.18015289	17.14575195	14.0344	-0.53976	1.453734		
	31.91260147	16.45843124	15.45417	0.880006	0.543365		
	30.5335598	16.34428978	14.18927	-0.38489	1.305764		
	Mean ΔCt (control) 14.57416						
			Fold gene	expression	1.046622		
			Standa	rd deviation	0.319573		
		St	andard error	of the mean	0.130465		
		Relativ	e standard er	ror of mean	12.46534		
	30.49056244	16.40930367	14.08126	-0.49291	1.407276		
	30.18668175	16.24884605	13.93784	-0.63633	1.554368		
Female	30.4241333	16.33191109	14.09222	-0.48194	1.396622		
Female	31.6965847	16.53628731	15.1603	0.586133	0.666126		
	30.37435722	15.90371037	14.47065	-0.10352	1.07439		
	32.01665878	16.67467499	15.34198	0.76782	0.587304		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.4.14: Cyp3a2 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	33.79129791	18.69672966	15.09457	1.065265	0.477885		
Male	34.06178284	19.07109451	14.99069	0.961385	0.513564		
iviale	33.04243469	20.00292778	13.03951	-0.9898	1.985904		
	31.88895798	18.61770439	13.27125	-0.75805	1.691202		
	32.4364624	18.68596458	13.7505	-0.27881	1.21319		
	Mean ΔCt (control) 14.0293						
			Fold ger	e expression	1.176349		
			Standa	ard deviation	0.67988		
		S	tandard error	of the mean	0.304052		
		Relati	ve standard e	rror of mean	25.84706		
	34.78320313	18.77750015	16.0057	1.9764	0.254123		
	32.60792542	18.58622551	14.0217	-0.0076	1.005284		
200	33.42826462	18.83141136	14.59685	0.56755	0.674762		
Female	32.79399109	18.85993767	13.93405	-0.09525	1.06825		
	34.10342026	18.42444038	15.67898	1.649677	0.318712		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean							

Table S.4.15: Cyp3a9 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a9</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	26.31387329	18.62302208	7.690851	-0.19149	1.14194		
	26.75233841	18.69672966	8.055609	0.173271	0.88683		
Male	26.97164726	19.07109451	7.900553	0.018215	0.987454		
iviale							
		Mean ΔCt (control)	7.882338				
			Fold gene	eexpression	1.005408		
			Standa	rd deviation	0.128499		
		S	tandard error	of the mean	0.074189		
		Relati	ve standard er	ror of mean	7.379001		
	28.16468811	18.77750015	9.387188	1.50485	0.352367		
	27.47090149	18.58622551	8.884676	1.002338	0.49919		
Female	26.27395821	18.83141136	7.442547	-0.43979	1.356408		
Terriale							
	26.59047699	18.42444038	8.166037	0.283699	0.821482		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relati	ve standard er	ror of mean	29.36757		

Table S.4.16: Cyp3a23 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	29.34032059	16.40213203	12.93819	-0.78537	1.723532	
	29.72708893	15.77114773	13.95594	0.232384	0.851227	
D.AI	31.24424934	16.00558853	15.23866	1.515104	0.349871	
Male	29.9582901	17.14575195	12.81254	-0.91102	1.880373	
	30.01674461	16.34428978	13.67245	-0.0511	1.036056	
		Mean ΔCt (control)	13.72356			
			Fold gen	e expression	1.168212	
			Standa	ard deviation	0.633084	
		S	tandard error	of the mean	0.283124	
		Relati	ve standard e	rror of mean	24.23564	
	30.63632965	16.40930367	14.22703	0.503469	0.705408	
Female	30.72545242	16.33191109	14.39354	0.669985	0.628513	
remale	30.19988823	16.53628731	13.6636	-0.05996	1.042434	
	30.33937454	15.90371037	14.43566	0.712107	0.610428	
Fold gene expression						
Standard deviation						
Standard error of the mean						
Relative standard error of mean						

Table S.4.17: *Cyp4a1* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)			
	25.17637634	18.62302208	6.553354	0.026118	0.982059			
	24.97893333	18.69672966	6.282204	-0.24503	1.185119			
D.AI	25.91881752	19.07109451	6.847723	0.320487	0.800799			
Male								
	25.07326508	18.61770439	6.455561	-0.07168	1.050936			
	25.18330193	18.68596458	6.497337	-0.0299	1.02094			
	Mean ΔCt (control) 6.527236							
			Fold ge	ne expression	1.007971			
			Stand	lard deviation	0.138733			
			Standard erro	r of the mean	0.062043			
		Rela	tive standard	error of mean	6.155258			
	24.39503098	18.58622551	5.808805	-0.71843	1.645391			
Female	25.06937599	18.83141136	6.237965	-0.28927	1.222023			
remale	24.81588936	18.85993767	5.955952	-0.57128	1.485845			
	24.60747147	18.42444038	6.183031	-0.3442	1.269451			
	25.71463203	19.72647858	5.988153	-0.53908	1.453048			
	Fold gene expression							
	Standard deviation							
	Standard error of the mean							
Relative standard error of mean								

Table S.4.18: *Cyp4a3* mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	30.68878555	17.78814507	12.90064	-0.53887	1.452839		
	30.86901855	17.76878548	13.10023	-0.33928	1.265127		
Male	31.07027245	18.14568329	12.92459	-0.51493	1.428921		
iviale	33.37727356	19.40160942	13.97566	0.536149	0.689609		
	32.9477005	18.65125275	14.29645	0.856933	0.552125		
		Mean ΔCt (control)	13.43951				
			Fold gen	e expression	1.077724		
			Standa	ard deviation	0.426045		
		S	tandard error	of the mean	0.190533		
		Relati	ive standard e	rror of mean	17.67921		
	31.9852562	18.4491787	13.53608	0.096563	0.935259		
	30.68878555 17.78814507 12.90064 -0.53887 1 30.86901855 17.76878548 13.10023 -0.33928 1 31.07027245 18.14568329 12.92459 -0.51493 1 33.37727356 19.40160942 13.97566 0.536149 0						
Female	31.20719719	18.34965897	12.85754	-0.58198	1.496899		
remale	32.93345261	18.3086586	14.62479	1.185279	0.439739		
	31.0255127	17.64413452	13.38138	-0.05814	1.04112		
	32.1522522	18.43346977	13.71878	0.279268	0.824009		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						

Table S.4.19: Cyp4f1 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	31.08237648	17.78814507	13.29423	0.20195	0.869375		
	31.69880104	17.76878548	13.93002	0.837734	0.559522		
Male	31.53038216	18.14568329	13.3847	0.292418	0.816533		
iviale	31.50394821	19.40160942	12.10234	-0.98994	1.986106		
	31.7729454	18.4796257	13.29332	0.201038	0.869924		
	31.20033646	18.65125275	12.54908	-0.5432	1.457199		
		Mean ΔCt (control)	13.09228				
			Fold gene	expression	1.09311		
			Standa	rd deviation	0.527462		
			Standard error	of the mean	0.215336		
		Relat	tive standard er	ror of mean	19.69936		
	31.43117523	18.4491787	12.982	-0.11028	1.079441		
Female	32.04748154	18.34965897	13.69782	0.605541	0.657225		
remale	32.74836731	18.3086586	14.43971	1.347427	0.392992		
	32.17800522	17.64413452	14.53387	1.441589	0.368161		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
		Relat	tive standard er	ror of mean	26.44923		

Table S.4.20: Cyp4f4 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	31.61248589	17.78814507	13.82434	0.85487	0.552915		
	31.15543175	17.76878548	13.38665	0.417176	0.748889		
Mala	32.20957947	18.14568329	14.0639	1.094426	0.468322		
Male	32.01199341	19.40160942	12.61038	-0.35909	1.282613		
	30.17363548	18.4796257	11.69401	-1.27546	2.420761		
	30.88879776	18.65125275	12.23755	-0.73193	1.660854		
	Mean ΔCt (control) 12.96947						
			Fold gene	expression	1.189059		
			Standar	d deviation	0.757142		
		St	andard error o	f the mean	0.309102		
		Relativ	e standard err	or of mean	25.99552		
	31.85640335	18.4491787	13.40722	0.437754	0.738283		
	30.42624092	18.31050491	12.11574	-0.85373	1.807173		
Female	31.10124969	18.34965897	12.75159	-0.21788	1.163023		
Female	31.51767921	18.3086586	13.20902	0.23955	0.847009		
	31.5107708	17.64413452	13.86664	0.897166	0.53694		
	31.20701981	18.43346977	12.77355	-0.19592	1.145455		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.4.21: Cyp4f5 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	22.49903679	17.78814507	4.710892	0.469703	0.722113	
	22.18119049	17.76878548	4.412405	0.171216	0.888094	
Mala	22.31451988	18.14568329	4.168837	-0.07235	1.05143	
Male	22.78550911	19.40160942	3.3839	-0.85729	1.811631	
	22.40570831	18.4796257	3.926083	-0.31511	1.244103	
	23.49627113	18.65125275	4.845018	0.603829	0.658005	
		Mean ΔCt (control)	4.241189			
			Fold gene	expression	1.062563	
			Standard	d deviation	0.425252	
		Sta	ndard error o	f the mean	0.173608	
		Relative	standard err	or of mean	16.33865	
	22.37131882	18.4491787	3.92214	-0.31905	1.247508	
	22.63340569	18.31050491	4.322901	0.081712	0.944936	
Female	22.42640305	18.34965897	4.076744	-0.16444	1.120735	
Female	22.50321579	18.3086586	4.194557	-0.04663	1.032851	
	21.97297478	17.64413452	4.32884	0.087651	0.941054	
	22.58729744	18.43346977	4.153828	-0.08736	1.062425	
Fold gene expression						
Standard deviation					0.115724	
Standard error of the mean					0.047244	
Relative standard error of mean					4.464368	

Table S.4.22: Cyp4f6 mRNA expression in the lung of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	24.28021622	17.78814507	6.492071	0.192998	0.874786		
	24.16589165	17.76878548	6.397106	0.098033	0.934306		
D.AI	24.22339821	18.14568329	6.077715	-0.22136	1.16583		
Male	25.06870079	19.40160942	5.667091	-0.63198	1.549692		
	25.10493088	18.4796257	6.625305	0.326232	0.797617		
	25.18640137	18.65125275	6.535149	0.236076	0.849052		
		Mean ΔCt (control)	6.299073				
			Fold gen	e expression	1.028547		
			Standa	ard deviation	0.285836		
		S	tandard error	of the mean	0.116692		
		Relati	ve standard e	rror of mean	11.34532		
	24.25156212	18.4491787	5.802383	-0.49669	1.410972		
	24.60692596	18.31050491	6.296421	-0.00265	1.00184		
Female	24.59147263	18.34965897	6.241814	-0.05726	1.040487		
remale	25.28375435	18.3086586	6.975096	0.676023	0.625888		
	24.31830597	18.45649719	5.861809	-0.43726	1.354034		
	25.18746758	18.43346977	6.753998	0.454925	0.729548		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
		Relati	ve standard e	rror of mean	12.62627		

S.5 Calculation of the fold change in the brain level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta CT$ method:

Table S.5.1: *Cyp1a1* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (Cyp1a1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^- (ΔΔCt)		
	29.85753632	21.95253754	7.904999	-0.30201	1.232859		
	30.37665939	22.01350594	8.363153	0.156147	0.897419		
Male	29.41322899	22.00631142	7.406918	-0.80009	1.741209		
iviale	30.44920921	21.12924385	9.319965	1.112959	0.462345		
	29.52134705	21.48134804	8.039999	-0.16701	1.122728		
		Mean ΔCt (control)	8.207007				
			Fold ge	ne expression	1.091312		
			Stand	lard deviation	0.46813		
			Standard erro	or of the mean	0.209354		
		Rela	tive standard	error of mean	19.18371		
	30.63818169	21.29172325	9.346458	1.139452	0.453932		
	30.29972458	21.2209301	9.078794	0.871788	0.546469		
Female	30.91126251	21.59479332	9.316469	1.109462	0.463467		
remale	29.87859535	21.26463318	8.613962	0.406955	0.754213		
	30.99826622	21.23768806	9.760578	1.553571	0.340666		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
		Rela	tive standard	error of mean	13.46318		

Table S.5.2: Cyp1a2 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp1a2</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	27.48363686	21.95253754	5.531099	-0.79458	1.734572		
	29.45805359	22.01350594	7.444548	1.118869	0.460455		
Mala	28.02119827	22.00631142	6.014887	-0.31079	1.240389		
iviale							
	28.63006783	21.48134804	7.14872	0.823041	0.565249		
	27.00094032	21.51179886	5.489141	-0.83654	1.785759		
		Mean ΔCt (control)	6.325679				
			Fold gene	e expression	1.157285		
			Standa	rd deviation	0.626743		
		S	tandard error	of the mean	0.280288		
		Relati	ive standard er	ror of mean	24.21943		
	27.65212059	21.29172325	6.360397	0.034718	0.976222		
	27.48363686 21.95253754 5.531099 -0.79458 29.45805359 22.01350594 7.444548 1.118869 28.02119827 22.00631142 6.014887 -0.31079 28.63006783 21.48134804 7.14872 0.823041 -0.83654 27.00094032 21.51179886 5.489141 -0.83654 27.00094032 21.51179886 Standard deviation Standard deviation Standard error of the mean Relative standard error of mean Relative standard error of mean 28.81288338 21.2209301 7.591953 1.266274 29.3457756 21.59479332 7.750982 1.425303 27.80597305 21.26463318 6.54134 0.215661 28.88311195 21.23768806 7.645424 1.319745 27.90197372 21.12900352 6.77297 0.447291 Fold gene expression Standard deviation	0.415732					
Fomala	29.3457756	21.59479332	7.750982	-0.79458 1.118869 -0.31079 0.823041 -0.83654 De expression and deviation of the mean one of the mean of the	0.372341		
remale	27.80597305	21.26463318	6.54134	0.215661	0.861152		
	28.88311195	21.23768806	7.645424	1.319745	0.400606		
	27.90197372	21.12900352	6.77297	0.447291	0.733419		
	Fold gene expression						
	Standard deviation						
		S	tandard error	of the mean	0.107834		
	Relative standard error of mean						

Table S.5.3: Cyp1b1 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp1b1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.50772476	21.95253754	2.555187	-0.57608	1.490795	
	25.18506813	22.01350594	3.171562	0.040293	0.972457	
0.0-1-	25.61454964	22.00631142	3.608238	0.476969	0.718485	
Male	25.10695457	21.12924385	3.977711	0.846442	0.556155	
	24.31092453	21.48134804	2.829576	-0.30169	1.23259	
	24.15713882	21.51179886	2.64534	-0.48593	1.400488	
		Mean ΔCt (control)	3.131269			
			Fold gene	expression	1.061828	
			Standa	rd deviation	0.37668	
		St	andard error	of the mean	0.153779	
		Relativ	e standard er	ror of mean	14.48247	
	24.23845863	21.29172325	2.946735	-0.18453	1.13645	
	24.765728	21.2209301	2.5555187 -0.57608 2.53754 2.555187 -0.57608 2.50594 3.171562 0.040293 2.31142 3.608238 0.476969 2.24385 3.977711 0.846442 2.34804 2.829576 -0.30169 2.79886 2.64534 -0.48593 Ct (control) 3.131269 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean Relative standard error of mean 7.2325 2.946735 -0.18453 2.9301 3.544798 0.413529 2.49103 -0.64024 2.63318 4.416653 1.285384 2.68806 3.665916 0.534647	0.750785		
Female	24.08582306	21.59479332	2.49103	-0.57608 0.040293 0.476969 0.846442 -0.30169 -0.48593 e expression rd deviation of the mean ror of mean -0.18453 0.413529 -0.64024 1.285384 0.534647 -0.45651 e expression rd deviation of the mean	1.558588	
remale	24.50772476 21.95253754 2.555187 -0.57608 25.18506813 22.01350594 3.171562 0.040293 25.61454964 22.00631142 3.608238 0.476969 25.10695457 21.12924385 3.977711 0.846442 24.31092453 21.48134804 2.829576 -0.30169 24.15713882 21.51179886 2.64534 -0.48593 Mean ΔCt (control) 3.131269 Fold gene expression Standard error of the mean Relative standard error of mean Relative standard error of mean 24.23845863 21.29172325 2.946735 -0.18453 24.765728 21.2209301 3.544798 0.413529 24.08582306 21.59479332 2.49103 -0.64024 25.68128586 21.26463318 4.416653 1.285384 24.90360451 21.23768806 3.665916 0.534647 23.80375862 21.12900352 2.674755 -0.45651 Fold gene expression Standard deviation Standard error of the mean	0.410262				
	24.90360451	21.23768806	3.665916	0.534647	0.690327	
	23.80375862	21.12900352	2.674755	-0.45651	1.372222	
			Fold gene	expression	0.986439	
	Standard deviation					
		St	andard error	of the mean	0.180174	
	Relative standard error of mean					

Table S.5.4: *Cyp2a1* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	34.17876434	21.95253754	12.22623	-1.68856	3.22334		
	35.20835114	22.01350594	13.19485	-0.71994	1.647111		
Mala	36.75863266	22.00631142	14.75232	0.837538	0.559598		
Male							
	36.33463669	21.48134804	14.85329	0.938506	0.521773		
	36.05903244	21.51179886	14.54723	0.63245	0.64508		
		Mean ΔCt (control)	13.91478				
			Fold gene	expression	1.31938		
			Standar	d deviation	1.161959		
		Sta	ndard error o	f the mean	0.519644		
		Relative	standard err	or of mean	39.38546		
	28.20470047	21.29172325	6.912977	-7.00181	128.1603		
	33.10462952	21.2209301	11.8837	-2.03108	4.087117		
Female							
remaie	27.43479919	21.26463318	6.170166	-7.74462	214.4678		
	31.88129616	21.23768806	10.64361	-3.27118	9.654322		
	33.82952118	21.12900352	12.70052	-1.21427	2.320226		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
	Relative standard error of mean						

Table S.5.5: Cyp2b1 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2b1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	27.04485893	20.6919651	6.352894	-1.26488	2.40308	
	28.96102715	20.69056702	8.27046	0.652682	0.636097	
Mala	28.21276855	20.91047478	7.302294	-0.31548	1.24443	
Male	28.14666557	20.16667366	7.979992	0.362214	0.77797	
	28.91030884	20.7270565	8.183252	0.565474	0.675733	
		Mean ΔCt (control)	7.617778			
			Fold gene	expression	1.147462	
			Standard	d deviation	0.742719	
		Stan	dard error o	f the mean	0.332154	
		Relative	standard err	or of mean	28.94684	
	29.1765976	20.48005295	8.696545	1.078766	0.473434	
	28.16078949	20.2576561	7.903133	0.285355	0.82054	
Female	30.07255173	20.42548943	9.647062	2.029284	0.244977	
remale	28.07061768	20.45305443	7.617563	-0.00022	1.000149	
	29.40126228	20.31982613	9.081436	1.463658	0.362573	
			Fold gene	expression	0.580334	
	Standard deviation					
	Standard error of the mean					
		Relative	standard err	or of mean	24.52852	

Table S.5.6: Cyp2b2 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2b2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	28.43515968	21.95253754	6.482622	-0.87884	1.838898	
	29.67820549	22.01350594	7.6647	0.303236	0.810433	
Male	29.23870087	22.00631142	7.232389	-0.12907	1.093592	
iviale	28.97444916	21.12924385	7.845205	0.483741	0.715121	
	29.06375122	21.48134804	7.582403	0.220939	0.858007	
		Mean ΔCt (control)	7.361464			
			Fold gene	expression	1.06321	
			Standa	rd deviation	0.455463	
		Star	ndard error	of the mean	0.203689	
		Relative	standard er	ror of mean	19.15795	
	29.36781693	21.29172325	8.076094	0.71463	0.609361	
	28.95804024	21.2209301	7.73711	0.375646	0.77076	
Female	30.14954185	21.59479332	8.554749	1.193285	0.437306	
remale	28.69236946	21.26463318	7.427736	0.066272	0.955103	
	29.77027702	21.23768806	8.532589	1.171125	0.444075	
			Fold gene	expression	0.643321	
	Standard deviation					
		Star	ndard error	of the mean	0.099182	
Relative standard error of mean						

Table S.5.7: Cyp2c6 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	34.24165726	21.95253754	12.28912	-0.05387	1.038042		
	34.65101624	22.01350594	12.63751	0.294525	0.815341		
Mala	34.79759979	22.00631142	12.79129	0.448303	0.732904		
iviale	34.06654358	21.12924385	12.9373	0.594315	0.662359		
	32.99552917	21.48134804	11.51418	-0.8288	1.776212		
	33.40031052	21.51179886	11.88851	-0.45447	1.370283		
		Mean ΔCt (control)	12.34299				
			Fold gene	expression	1.065857		
			Standa	rd deviation	0.432315		
		Sta	ndard error	of the mean	0.176492		
		Relative	e standard er	ror of mean	16.55867		
	33.55820084	21.2209301	12.33727	-0.00571	1.003969		
Fomala	Male 34.06654358 21.12924385 12.9373 0.594315 0 32.99552917 21.48134804 11.51418 -0.8288 1 33.40031052 21.51179886 11.88851 -0.45447 1 Mean ΔCt (control) 12.34299 Fold gene expression 1 Standard deviation 0 Relative standard error of the mean 1 33.55820084 21.2209301 12.33727 -0.00571 1	1.884648					
remale							
	33.76731491	21.23768806	12.52963	0.186642	0.878649		
	32.69213867	21.12900352	11.56314	-0.77985	1.716952		
	Fold gene expression						
Standard deviation					0.503534		
		Sta	ndard error	of the mean	0.251767		
Relative standard error of mean							

Table S.5.8: Cyp2c11 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c11</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	28.23395538	21.95253754	6.281418	-0.47869	1.393476		
	29.25510025	22.01350594	7.241594	0.481489	0.716238		
Mala	28.05605125	22.00631142	6.04974	-0.71037	1.636219		
Iviale	28.623312	21.12924385	7.494068	0.733963	0.60125		
	28.21505547	21.48134804	6.733707	-0.0264	1.018466		
		Mean ΔCt (control)	6.760106				
			Fold gene	expression	1.07313		
			Standar	d deviation	0.439523		
		Sta	ndard error o	f the mean	0.196561		
		Relative	standard err	or of mean	18.31657		
	28.96500587	21.29172325	7.673283	0.913177	0.531014		
	28.39972687	21.2209301	7.178797	0.418691	0.748103		
Fomolo	Male 28.23395538 21.95253754 6.281418 -0.47869 29.25510025 22.01350594 7.241594 0.481489 28.05605125 22.00631142 6.04974 -0.71037 28.623312 21.12924385 7.494068 0.733963 28.21505547 21.48134804 6.733707 -0.0264 Mean ΔCt (control) 6.760106 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean Relative standard error of mean 28.96500587 21.29172325 7.673283 0.913177 28.39972687 21.2209301 7.178797 0.418691 28.98079872 21.59479332 7.386005 0.6259 28.64322853 21.26463318 7.378595 0.61849 29.60293961 21.23768806 8.365252 1.605146 Fold gene expression Standard deviation Standard deviation	0.648015					
remale	28.64322853	21.26463318	7.378595	0.61849	0.651352		
	29.60293961	21.23768806	8.365252	1.605146	0.328702		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
		Relative	standard err	or of mean	12.37328		

Table S.5.9: *Cyp2c13* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2c13</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.77687073	20.91020012	13.86667	-0.99182	1.988686	
	36.55079651	21.23644829	15.31435	0.455862	0.729074	
Malo	35.66214752	21.20062065	14.46153	-0.39696	1.316729	
ividie						
	37.54246902	21.14937782	16.39309	1.534605	0.345174	
	35.31303406	21.05624199	14.25679	-0.60169	1.517497	
		Mean ΔCt (control)	14.85849			
			Fold gene	expression	1.179432	
			Standard	d deviation	0.649392	
		Star	dard error o	f the mean	0.290417	
		Relative	standard err	or of mean	24.62345	
	37.3074913	20.75029945	16.55719	1.698706	0.308062	
	36.66149902	20.74041748	15.92108	1.062596	0.47877	
Formula	Male 36.55079651 21.23644829 15.31435 0.455862 35.66214752 21.20062065 14.46153 -0.39696 37.54246902 21.14937782 16.39309 1.534605 35.31303406 21.05624199 14.25679 -0.60169 Mean ΔCt (control) 14.85849 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean Relative standard error of mean 37.3074913 20.75029945 16.55719 1.698706 36.66149902 20.74041748 15.92108 1.062596 36.75853729 20.70553017 16.05301 1.194521 37.45233536 20.68715286 16.76518 1.906697 38.07253647 20.51906967 17.55347 2.694981 37.68747711 20.89987946 16.7876 1.929112	0.436931				
remale	37.45233536	20.68715286	16.76518	1.906697	0.266703	
	38.07253647	20.51906967	17.55347	2.694981	0.154429	
	37.68747711	20.89987946	16.7876	1.929112	0.262591	
Fold gene expression						
Standard deviation					0.120455	
	Standard error of the mean					
Relative standard error of mean						

Table S.5.10: *Cyp2c23* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	28.88280869	20.91020012	7.972609	-0.56892	1.483411		
	29.67432976	21.23644829	8.437881	-0.10365	1.074485		
Male	30.19148636	21.20062065	8.990866	0.449339	0.732378		
iviale	30.34165001	20.83230591	9.509344	0.967818	0.511279		
	29.57940865	21.14937782	8.430031	-0.1115	1.080348		
	28.96467018	21.05624199	7.908428	-0.6331	1.550892		
		Mean ΔCt (control)	8.541526				
			Fold gene	expression	1.072132		
			Standard	d deviation	0.407044		
		Star	dard error o	f the mean	0.166175		
		Relative	standard err	or of mean	15.49948		
	29.56397247	20.74041748	8.823555	0.282029	0.822434		
Female	31.24544907	20.70553017	10.53992	1.998392	0.250279		
remale							
	28.6942215	20.51906967	8.175152	-0.36637	1.289109		
	31.86293793	20.89987946	10.96306	2.421532	0.186658		
	Fold gene expression						
	Standard deviation						
		Star	ndard error o	f the mean	0.260127		
Relative standard error of mean					40.8286		

Table S.5.11: Cyp2d4 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2d4)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	24.74249458	20.6919651	4.050529	-0.72336	1.651019	
	25.50859261	20.69056702	4.818026	0.04414	0.969868	
	26.12778282	20.91047478	5.217308	0.443422	0.735388	
Male	25.75165749	20.16667366	5.584984	0.811098	0.569948	
	25.15022469	20.7270565	4.423168	-0.35072	1.275195	
	25.16477966	20.61547852	4.549301	-0.22458	1.168441	
		Mean ΔCt (control)	4.773886			
			Fold gene	expression	1.061643	
			Standa	rd deviation	0.390129	
		St	andard error	of the mean	0.159269	
		Relativ	e standard er	ror of mean	15.00217	
	25.28174019	20.48005295	4.801687	0.027801	0.980914	
	25.64151955	20.2576561	5.383863	0.609977	0.655207	
Female	25.28036118	20.42548943	4.854872	0.080986	0.945411	
Female	25.73700142	20.6919651 4.050529 -0.72336 20.69056702 4.818026 0.04414 20.91047478 5.217308 0.443422 20.16667366 5.584984 0.811098 20.7270565 4.423168 -0.35072 20.61547852 4.549301 -0.22458 Mean ΔCt (control) 4.773886 Fold gene expression Standard deviation Standard error of the mean Relative standard error of mean 20.48005295 4.801687 0.027801 20.2576561 5.383863 0.609977	0.702193			
	25.56605911	20.31982613	5.246233	0.472347	0.720791	
			Fold gene	expression	0.800903	
		St	andard error	of the mean	0.067333	
		Relativ	e standard er	ror of mean	8.40718	

Table S.5.12: Cyp2e1 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp2e1)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	24.97688675	20.6919651	4.284922	-0.8663	1.822986		
	26.75543594	20.69056702	6.064869	0.913644	0.530843		
Male	26.70380592	20.91047478	5.793331	0.642106	0.640777		
ividie							
	25.93351936	20.7270565	5.206463	0.055238	0.962436		
	25.02202034	20.61547852	4.406542	-0.74468	1.675607		
		Mean ΔCt (control)	5.151225				
			Fold gene	expression	1.12653		
			Standa	rd deviation	0.592508		
		St	andard error	of the mean	0.264978		
		Relativ	e standard er	ror of mean	23.52158		
	25.61355209	20.48005295	5.133499	-0.01773	1.012363		
	26.62290001	20.2576561	6.365244	1.214019	0.431066		
Female	27.19693565	20.42548943	6.771446	1.620221	0.325286		
remale							
	26.87777138	20.31982613	6.557945	1.40672	0.377168		
	25.29464531	20.27284431	5.021801	-0.12942	1.093857		
	Fold gene expression						
Standard deviation							
	Standard error of the mean						
Relative standard error of mean							

Table S.5.13: *Cyp2j3* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β-actin housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	22.66023445	22.56298447	0.09725	-0.80681	1.74934
	23.89534569	22.7211895	1.174156	0.270095	0.829265
Mala	22.96236992	22.25282288	0.709547	-0.19451	1.144339
Male	23.90988922	22.0347538	1.875135	0.971074	0.510126
	22.5336113	21.94296837	0.590643	-0.31342	1.242648
	22.76453781	21.78690338	0.977634	0.073573	0.950281
		Mean ΔCt (control)	0.904061		
			Fold gene	expression	1.071
			Standard	d deviation	0.420137
		Sta	ndard error o	f the mean	0.17152
		Relative	standard err	or of mean	16.01496
	23.69215202	22.32851219	1.36364	0.459579	0.727199
	23.40186501	21.3197937	2.082071	1.17801	0.441961
Female	23.34770775	22.2055912	1.142117	0.238056	0.847887
remale					
	23.75100327	22.27744484	1.473558	0.569497	0.673851
	22.46575737	21.60790634	0.857851	-0.04621	1.032549
			Fold gene	expression	0.744689
	Standard deviation				
		Sta	ndard error o	f the mean	0.097591
		Relative	standard err	or of mean	13.10498

Table S.5.14: Cyp2j4 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	27.88459206	22.56298447	5.321608	0.037323	0.974462
	27.99481392	22.7211895	5.273624	-0.01066	1.007417
Male	27.2287178	22.25282288	4.975895	-0.30839	1.238325
iviale	27.78040886	22.0347538	5.745655	0.46137	0.726296
	27.04761124	21.94296837	5.104643	-0.17964	1.132603
		Mean ΔCt (control)	5.284285		
			Fold gene	expression	1.015821
			Standar	d deviation	0.192842
		Star	ndard error o	f the mean	0.086241
		Relative	standard err	or of mean	8.489836
	27.90842247	22.32851219	5.57991	0.295625	0.814719
	26.9987545	21.3197937	5.678961	0.394676	0.76066
Female	27.68693542	22.2055912	5.481344	0.197059	0.872327
Female	27.68802834	22.48288155	5.205147	-0.07914	1.056387
	27.64679146	22.27744484	5.369347	0.085062	0.942744
	26.6769352	21.60790634	5.069029	-0.21526	1.16091
			Fold gene	expression	0.934625
	Standard deviation				
	Standard error of the mean				
		Relative	standard err	or of mean	6.613485

Table S.5.15: Cyp2j10 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	28.47204971	22.56298447	5.909065	-0.40297	1.322225
	28.58550072	22.7211895	5.864311	-0.44772	1.363885
Male	28.43003654	22.25282288	6.177214	-0.13482	1.097956
iviale	28.61799431	22.0347538	6.583241	0.271207	0.828626
	28.42879868	21.94296837	6.48583	0.173797	0.886506
	28.63944244	21.78690338	6.852539	0.540506	0.68753
		Mean ΔCt (control)	6.312033		
			Fold gene	expression	1.031121
			Standar	d deviation	0.275649
		Star	ndard error o	f the mean	0.112533
		Relative	standard err	or of mean	10.91369
	28.45299149	22.32851219	6.124479	-0.18755	1.138831
	28.14215088	21.3197937	6.822357	0.510324	0.702065
Female	28.322052	22.2055912	6.116461	-0.19557	1.145179
remale	29.18748093	22.48288155	6.704599	0.392566	0.761773
	28.20519638	22.27744484	5.927752	-0.38428	1.30521
	28.0281868	21.60790634	6.42028	0.108247	0.927715
Fold gene expression					
Standard deviation					0.238405
		Star	ndard error o	f the mean	0.097328
		Relative	standard err	or of mean	9.764135

Table S.5.16: Cyp3a1 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.74755478	20.6919651	14.05559	-0.28978	1.222457	
	34.7414093	20.69056702	14.05084	-0.29453	1.226487	
Mala	35.03034592	20.91047478	14.11987	-0.2255	1.169185	
Male	36.12369919	20.16667366	15.95703	1.611652	0.327224	
	34.15901947	20.61547852	13.54354	-0.80183	1.743315	
		Mean ΔCt (control)	14.34537			
			Fold gene	expression	1.137733	
			Standar	d deviation	0.509828	
		Sta	ndard error o	f the mean	0.228002	
		Relative	standard err	or of mean	20.04003	
	35.18642426	20.2576561	14.92877	0.583394	0.667392	
Female	35.30002213	20.42548943	14.87453	0.529159	0.692959	
remale						
	34.78128052	20.31982613	14.46145	0.11608	0.922691	
	34.71193314	20.27284431	14.43909	0.093715	0.937107	
	Fold gene expression					
	Standard deviation					
		Sta	ndard error o	f the mean	0.072338	
		Relative	standard err	or of mean	8.985615	

Table S.5.17: Cyp3a2 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a</i> 2)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	28.35564804	20.91020012	7.445448	-0.4318	1.348911
	29.4664669	21.23644829	8.230019	0.352775	0.783076
Male	29.22295761	21.20062065	8.022337	0.145094	0.904321
iviale	28.77317429	20.83230591	7.940868	0.063625	0.956857
	28.89692307	21.14937782	7.747545	-0.1297	1.094065
		Mean ΔCt (control)	7.877243		
			Fold gene	expression	1.017446
			Standar	d deviation	0.216297
		Star	ndard error o	f the mean	0.096731
		Relative	standard err	or of mean	9.507224
	29.07541847	20.75029945	8.325119	0.447876	0.733122
	28.70752335	20.74041748	7.967106	0.089862	0.939612
Female	30.04026222	20.70553017	9.334732	1.457489	0.364126
remale	28.63785744	20.68715286	7.950705	0.073461	0.950355
	29.50494766	20.51906967	8.985878	1.108635	0.463733
			Fold gene	expression	0.69019
	Standard deviation				
		Star	ndard error o	f the mean	0.120279
		Relative	standard err	or of mean	17.42695

Table S.5.18: Cyp3a9 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a9</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	27.48248291	21.86958313	5.6129	0.401224	0.757215		
	27.58653641	21.931036	5.6555	0.443825	0.735183		
Mala	26.97591972	22.0233326	4.952587	-0.25909	1.196722		
Male	26.75855064	21.61818314	5.140368	-0.07131	1.050669		
	27.02022743	21.90874672	5.111481	-0.10019	1.071918		
	26.72475052	21.9275341	4.797216	-0.41446	1.332799		
	Mean ΔCt (control) 5.211675						
			Fold gene	expression	1.024084		
			Standard	d deviation	0.237818		
		Stan	dard error o	f the mean	0.097089		
		Relative s	tandard err	or of mean	9.480536		
	26.95716286	21.78549576	5.171667	-0.04001	1.02812		
	26.97912979	21.55132675	5.427803	0.216128	0.860873		
Famala	26.61027718	21.27036667	5.339911	0.128235	0.91495		
Female	27.53744125	21.745224	5.792217	0.580542	0.668713		
	27.09966469	21.42292404	5.676741	0.465065	0.724438		
	26.75687599	21.66075897	5.096117	-0.11556	1.083394		
			Fold gene	expression	0.880081		
	Standard deviation						
		Stan	dard error o	f the mean	0.066768		
	Relative standard error of mean						

Table S.5.19: Cyp3a23 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	29.1442585	21.86958313	7.274675	-0.20085	1.149373	
	30.353899	21.931036	8.422863	0.94734	0.518588	
B.OI	29.05274391	22.0233326	7.029411	-0.44611	1.362363	
Male	28.98163414	21.61818314	7.363451	-0.11207	1.080779	
	29.195961	21.90874672	7.287214	-0.18831	1.139427	
		Mean ΔCt (control)	7.475523			
			Fold gene	expression	1.050106	
			Standa	rd deviation	0.315749	
		Sta	ndard error	of the mean	0.141207	
		Relative	e standard er	ror of mean	13.44695	
	28.05164146	21.78549576	6.266146	-1.20938	2.312378	
	28.60787392	21.55132675	7.056547	-0.41898	1.336978	
Female						
remaie	27.90856743	21.745224	6.163343	-1.31218	2.483164	
	29.43889427	21.42292404	8.01597	0.540447	0.687558	
	27.11496735	21.66075897	5.454208	-2.02131	4.059535	
	Fold gene expression					
	Standard deviation					
		Sta	ndard error	of the mean	0.573588	
	Relative standard error of mean					

Table S.5.20: *Cyp4a1* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	30.0105381	22.66277885	7.347759	-0.30606	1.236327	
	30.89823151	22.40570641	8.492525	0.838705	0.559145	
	29.91341782	22.53338432	7.380033	-0.27379	1.208977	
Male	29.78141975	22.07455635	7.706863	0.053043	0.963901	
	29.86969948	22.52778053	7.341919	-0.3119	1.241342	
		Mean ΔCt (control)	7.65382			
			Fold gene	expression	1.041939	
			Standard	d deviation	0.293526	
		Stan	dard error o	f the mean	0.131269	
		Relative	standard err	or of mean	12.59852	
	29.76352692	22.17542458	7.588102	-0.06572	1.046605	
	29.64139557	21.97904015	7.662355	0.008535	0.994101	
Female	30.65661621	21.90332222	8.753294	1.099474	0.466687	
remale	29.13052177	22.25868607	6.871836	-0.78198	1.719494	
	30.74164581	21.83511353	8.906532	1.252712	0.419659	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
	Relative standard error of mean					

Table S.5.21: *Cyp4a3* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (Cyp4a3)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	34.31351471	20.6919651	13.62155	-0.40172	1.321083	
	35.51944733	20.69056702	14.82888	0.80561	0.57212	
Mala	34.62796021	20.91047478	13.71749	-0.30579	1.236091	
Male						
	35.36398697	20.7270565	14.63693	0.61366	0.653537	
	33.92698669	20.61547852	13.31151	-0.71176	1.637804	
		Mean ΔCt (control)	14.02327			
			Fold gene	expression	1.084127	
			Standar	d deviation	0.456442	
		Stan	ndard error o	f the mean	0.204127	
		Relative	standard err	or of mean	18.82871	
	37.29369736	20.2576561	17.03604	3.01277	0.123898	
Female	38.43450165	20.42548943	18.00901	3.985741	0.063121	
remale						
	35.36367798	20.31982613	15.04385	1.020581	0.492918	
	37.94108582	20.27284431	17.66824	3.644971	0.079938	
			Fold gene	expression	0.189969	
	Standard deviation					
		Stan	dard error o	f the mean	0.101792	
		Relative	standard err	or of mean	53.5838	

Table S.5.22: *Cyp4a8* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4a8</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	38.1027565	22.66277885	15.43998	-0.29835	1.229738	
Male	39.00237274	22.53338432	16.46899	0.730659	0.602628	
Iviaic	38.14551926	22.07455635	16.07096	0.332634	0.794085	
	38.48109818	22.52778053	15.95332	0.214989	0.861553	
	37.11241531	22.35401726	14.7584	-0.97993	1.972371	
		Mean ΔCt (control)	15.73833			
			Fold gene	expression	1.092075	
			Standard	d deviation	0.542072	
		Stand	dard error o	f the mean	0.242422	
		Relative s	tandard err	or of mean	22.19829	
	38.01499557	22.17542458	15.83957	0.101242	0.93223	
	37.75903702	21.97904015	15.78	0.041668	0.971531	
Female	37.20704269	21.90332222	15.30372	-0.43461	1.351544	
remale	38.44969177	22.25868607	16.19101	0.452677	0.730686	
	38.71764374	21.98641777	16.73123	0.992897	0.502468	
	Fold gene expression					
	Standard deviation					
		Stan	dard error o	f the mean	0.14084	
		Relative s	tandard err	or of mean	15.68916	

Table S.5.23: *Cyp4f1* mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f1</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	26.03436661	22.66277885	3.371588	0.096731	0.93515	
	25.75484848	22.40570641	3.349142	0.074285	0.949813	
Male	26.02673912	22.53338432	3.493355	0.218498	0.85946	
iviale	25.07918549	22.07455635	3.004629	-0.27023	1.205998	
	25.9437542	22.52778053	3.415974	0.141116	0.906817	
	25.36847305	22.35401726	3.014456	-0.2604	1.197812	
		Mean ΔCt (control)	3.274857			
			Fold gene	expression	1.009175	
			Standard	d deviation	0.152459	
		Sta	ndard error o	f the mean	0.062241	
		Relative	standard err	or of mean	6.167524	
	25.0563488	22.17542458	2.880924	-0.39393	1.313971	
	25.15913391	21.97904015	3.180094	-0.09476	1.06789	
Female	25.02727318	21.90332222	3.123951	-0.15091	1.110267	
remale	25.40101814	22.25868607	3.142332	-0.13253	1.096211	
	25.12705994	21.83511353	3.291946	0.017089	0.988225	
	25.35582733	21.98641777	3.36941	0.094552	0.936563	
Fold gene expression						
Standard deviation					0.130293	
		Sta	ndard error o	f the mean	0.053192	
	Relative standard error of mean					

Table S.5.24: Cyp4f4 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)	
	29.64159012	22.40570641	7.235884	0.257257	0.836677	
Mala	30.55644226	22.53338432	8.023058	1.044431	0.484836	
Male	28.24415016	22.07455635	6.169594	-0.80903	1.752036	
	30.02590942	22.52778053	7.498129	0.519502	0.697612	
	28.32048607	22.35401726	5.966469	-1.01216	2.016926	
		Mean ΔCt (control)	6.978627			
			Fold gene	expression	1.157618	
			Standard	d deviation	0.681723	
		Stan	dard error o	f the mean	0.304876	
		Relative	standard err	or of mean	26.3365	
	27.9193821	22.17542458	5.743958	-1.23467	2.353274	
	27.87650681	21.97904015	5.897467	-1.08116	2.115737	
Female	27.45651054	21.90332222	5.553188	-1.42544	2.685961	
remale	28.70817947	22.25868607	6.449493	-0.52913	1.443062	
	27.96807289	21.83511353	6.132959	-0.84567	1.797096	
	28.11343384	21.98641777	6.127016	-0.85161	1.804514	
	Fold gene expression					
	Standard deviation					
		Stan	dard error o	f the mean	0.181811	
	Relative standard error of mean					

Table S.5.25: Cyp4f5 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)		
	25.70376587	22.66277885	3.040987	0.131496	0.912884		
	26.49518776	22.40570641	4.089481	1.179991	0.441354		
Male	25.23944473	22.53338432	2.70606	-0.20343	1.151433		
ividie	25.37679672	22.07455635	3.30224	0.39275	0.761676		
	24.63045311	22.52778053	2.102673	-0.80682	1.749349		
	24.56951904	22.35401726	2.215502	-0.69399	1.61775		
		Mean ΔCt (control)	2.909491				
			Fold gene	expression	1.105741		
			Standa	rd deviation	0.505066		
		Star	ndard error	of the mean	0.206192		
		Relative	standard er	ror of mean	18.64742		
	24.37293053	22.17542458	2.197506	-0.71198	1.638056		
	24.25177765	21.97904015	2.272738	-0.63675	1.554826		
Female	24.21597099	21.90332222	2.312649	-0.59684	1.512402		
remale	24.22727585	22.25868607	1.96859	-0.9409	1.919727		
	24.29445648	21.83511353	2.459343	-0.45015	1.36618		
	24.21575356	21.98641777	2.229336	-0.68015	1.602312		
	Fold gene expression						
	Standard deviation						
		Star	ndard error	of the mean	0.074846		
	Relative standard error of mean						

Table S.5.26: Cyp4f6 mRNA expression in the brain of male and female Sprague Dawley rats normalized to β -actin housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (B-ACTIN)	ΔCt	ΔΔCt	2^-(ΔΔCt)
	24.8198452	22.66277885	2.157066	0.125634	0.916601
	25.20314789	22.40570641	2.797441	0.766009	0.588042
Mala	24.31415749	22.53338432	1.780773	-0.25066	1.189751
Male	24.56652832	22.07455635	2.491972	0.460539	0.726714
	23.89409065	22.52778053	1.36631	-0.66512	1.585703
	23.949049	22.35401726	1.595032	-0.4364	1.353224
		Mean ΔCt (control)	2.031432		
			Fold gene	expression	1.060006
			Standar	d deviation	0.383032
		Star	ndard error o	f the mean	0.156372
		Relative	standard err	or of mean	14.75199
	24.13739204	22.17542458	1.961967	-0.06947	1.049327
	24.38249969	21.97904015	2.40346	0.372027	0.772696
Female	25.05745316	21.90332222	3.154131	1.122698	0.459234
remale	24.28108597	22.25868607	2.0224	-0.00903	1.006281
	25.26451874	21.83511353	3.429405	1.397973	0.379462
	23.79139709	21.98641777	1.804979	-0.22645	1.169955
Fold gene expression					
Standard deviation					0.327131
	Standard error of the mean				
Relative standard error of mean					16.5663

S.6 Calculation of the fold change in the small intestine level of target genes between female and male rats normalized to the housekeeping gene using the $\Delta\Delta$ CT method:

Table S.6.1: *Cyp1a1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp1a1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.42038918	15.0469017	6.373487	-0.07512	1.05345	
	22.09064102	15.63680172	6.453839	0.00523	0.996382	
Male	22.13542557	15.80290794	6.332518	-0.11609	1.083795	
iviale	22.58141327	15.58490181	6.996511	0.547902	0.684014	
	21.87853813	15.65109253	6.227446	-0.22116	1.165674	
	21.86164856	15.553792	6.307857	-0.14075	1.10248	
	Mean ΔCt (control) 6.44861					
			Fold gene	expression	1.014299	
			Standard	l deviation	0.17115	
		Stand	lard error of	the mean	0.069872	
		Relative s	tandard erro	or of mean	6.888673	
	21.78126526	15.26576138	6.515504	0.066894	0.954691	
	21.81917953	15.58676243	6.232417	-0.21619	1.161664	
Female	22.22576523	15.72859859	6.497167	0.048557	0.966903	
remale	22.1903038	15.84080791	6.349496	-0.09911	1.071115	
	21.86701012	15.89041996	5.97659	-0.47202	1.38705	
	22.72870445	16.63667107	6.092033	-0.35658	1.280384	
Fold gene expression						
Standard deviation					0.173322	
	Standard error of the mean					
Relative standard error of mean					6.223428	

Table S.6.2: *Cyp1a2* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp1a2</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	22.54761505	14.88868523	7.65893	-0.1963	1.145752	
	23.26819801	15.26900387	7.999194	0.143969	0.905026	
Male	23.51773071	15.34797287	8.169758	0.314533	0.804111	
iviale	22.91939163	15.24869919	7.670692	-0.18453	1.136449	
	23.09124374	15.07187366	8.01937	0.164145	0.892457	
	22.89465904	15.28125191	7.613407	-0.24182	1.182482	
	·	Mean ΔCt (control)	7.855225			
			Fold gene	expression	1.011046	
			Standard	deviation	0.162103	
		Stand	lard error o	f the mean	0.066178	
		Relative s	tandard err	or of mean	6.545535	
	22.90989685	15.29334736	7.616549	-0.23868	1.179909	
	22.90000343	15.16924477	7.730759	-0.12447	1.090105	
Female	23.16272354	15.29349518	7.869228	0.014003	0.990341	
remale	23.49699783	15.89090347	7.606094	-0.24913	1.188491	
	23.16763878	15.78082657	7.386812	-0.46841	1.383587	
	24.96151352	16.44277382	8.51874	0.663514	0.631338	
	Fold gene expression					
Standard deviation					0.254288	
	Standard error of the mean					
		Relative s	tandard erro	or of mean	9.636405	

Table S.6.3: *Cyp1b1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp1b1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)		
	22.43733978	15.0469017	7.390438	0.179531	0.88299		
	22.75009346	15.63680172	7.113292	-0.09761	1.070003		
Mala	23.14750099	15.80290794	7.344593	0.133686	0.911499		
Male	22.62650299	15.58490181	7.041601	-0.16931	1.124517		
	22.89332008	15.65109253	7.242228	0.031321	0.978524		
	22.68708038	15.553792	7.133288	-0.07762	1.055274		
		Mean ΔCt (control)	7.210907				
			Fold gene	expression	1.003801		
			Standard	deviation	0.095245		
		Stand	lard error of	f the mean	0.038884		
		Relative s	tandard erro	or of mean	3.873654		
	22.62806129	15.26576138	7.3623	0.151393	0.900381		
	22.79387283	15.58676243	7.20711	-0.0038	1.002635		
Famala	23.04993057	15.72859859	7.321332	0.110425	0.926315		
Female	23.18860626	15.84080791	7.347798	0.136892	0.909477		
	23.87608337	15.89041996	7.985663	0.774757	0.584487		
	24.67870331	16.63667107	8.042032	0.831126	0.562091		
	Fold gene expression						
	Standard deviation						
	Standard error of the mean						
Relative standard error of mean					9.537232		

Table S.6.4: *Cyp2a1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2a1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	34.42749405	14.88868523	19.53881	0.469621	0.722154	
	33.82606125	15.26900387	18.55706	-0.51213	1.426155	
Male	34.58669281	15.34797287	19.23872	0.169532	0.889131	
iviale	34.41130447	15.24869919	19.16261	0.093417	0.9373	
	34.11862183	15.07187366	19.04675	-0.02244	1.015676	
	34.15243912	15.28125191	18.87119	-0.198	1.147107	
		Mean ΔCt (control)	19.06919			
			Fold gene	expression	1.02292	
			Standard	deviation	0.24239	
		Stand	lard error of	f the mean	0.098955	
		Relative s	tandard erro	or of mean	9.673785	
	34.59760284	15.29334736	19.30426	0.235068	0.849645	
	34.2593956	15.16924477	19.09015	0.020963	0.985575	
Female	34.16900253	15.29349518	18.87551	-0.19368	1.143678	
remaie	34.72328186	15.89090347	18.83238	-0.23681	1.178384	
	34.96700287	15.78082657	19.18618	0.116988	0.92211	
	36.20910263	16.44277382	19.76633	0.697141	0.616793	
Fold gene expression						
Standard deviation					0.206305	
	Standard error of the mean					
		Relative s	tandard erro	or of mean	8.87159	

Table S.6.5: *Cyp2b1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (Cyp2b1)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	20.62836266	14.88868523	5.739677	-0.00457	1.003173	
	21.12841606	15.26900387	5.859412	0.115164	0.923277	
Male	21.22377396	15.34797287	5.875801	0.131553	0.912848	
iviale	20.8448925	15.24869919	5.596193	-0.14805	1.108074	
	20.75500679	15.07187366	5.683133	-0.06111	1.043272	
	20.99252319	15.28125191	5.711271	-0.03298	1.023121	
		Mean ΔCt (control)	5.744248			
			Fold gene	expression	1.002294	
			Standard	deviation	0.074226	
		Stand	lard error of	f the mean	0.030303	
		Relative s	tandard erro	or of mean	3.023321	
	20.77105522	15.29334736	5.477708	-0.26654	1.20292	
	20.8916378	15.16924477	5.722393	-0.02186	1.015264	
Female	21.0814209	15.29349518	5.787926	0.043678	0.970179	
remale	22.27885628	15.89090347	6.387953	0.643705	0.640067	
	21.37211418	15.78082657	5.591288	-0.15296	1.111849	
	21.22587776	16.44277382	4.783104	-0.96114	1.946853	
Fold gene expression						
Standard deviation					0.435827	
	Standard error of the mean					
		Relative s	tandard erro	or of mean	15.50069	

Table S.6.6: *Cyp2b2* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (Cyp2b2)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	20.54189491	15.0469017	5.494993	-0.04	1.028116	
	21.3386898	15.63680172	5.701888	0.166892	0.890759	
Male	21.4197731	15.80290794	5.616865	0.081869	0.944833	
iviale	21.00575256	15.58490181	5.420851	-0.11414	1.082333	
	21.22142601	15.65109253	5.570333	0.035338	0.975803	
	20.9588356	15.553792	5.405044	-0.12995	1.094257	
	·	Mean ΔCt (control)	5.534996			
			Fold gene	expression	1.002684	
			Standard	d deviation	0.079966	
		Stand	lard error of	f the mean	0.032646	
		Relative s	tandard erro	or of mean	3.255869	
	20.83866119	15.26576138	5.5729	0.037904	0.974069	
	20.98480415	15.58676243	5.398042	-0.13695	1.099581	
Female	21.42878342	15.72859859	5.700185	0.165189	0.891812	
remale	22.68405342	15.84080791	6.843246	1.30825	0.40381	
	21.53352928	15.89041996	5.643109	0.108114	0.9278	
	22.80368614	16.63667107	6.167015	0.632019	0.645273	
	Fold gene expression					
Standard deviation					0.253906	
	Standard error of the mean					
	Relative standard error of mean					

Table S.6.7: *Cyp2c6* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2c6</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)
	36.61317444	14.88868523	21.72449	1.267428	0.4154
	34.90011978	15.26900387	19.63112	-0.82595	1.772696
Male	34.67299652	15.34797287	19.32502	-1.13204	2.19168
iviale	35.95380783	15.24869919	20.70511	0.248048	0.842035
	36.1480217	15.07187366	21.07615	0.619087	0.651083
	35.56173325	15.28125191	20.28048	-0.17658	1.130201
		Mean ΔCt (control)	20.45706		
			Fold gene	expression	1.167183
			Standard	deviation	0.686275
		Stand	lard error of	f the mean	0.280171
		Relative s	tandard erro	or of mean	24.00401
	36.99253464	15.29334736	21.69919	1.242126	0.422749
	35.68371964	15.16924477	20.51447	0.057414	0.960985
EI-	36.69110489	15.29349518	21.39761	0.940549	0.521035
Female	35.78203964	15.89090347	19.89114	-0.56592	1.480336
	35.76646042	15.78082657	19.98563	-0.47143	1.38648
			Fold gene	expression	0.954317
	Standard deviation				
	Standard error of the mean				
	Relative standard error of mean				

Table S.6.8: Cyp2c11 mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2c11</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	22.2719326	14.88868523	7.383247	-0.16109	1.118134	
	22.98664284	15.26900387	7.717639	0.173299	0.886813	
Male	23.23209	15.34797287	7.884117	0.339777	0.790164	
iviale	22.58089638	15.24869919	7.332197	-0.21214	1.158408	
	22.65786552	15.07187366	7.585992	0.041651	0.971542	
	22.6441021	15.28125191	7.36285	-0.18149	1.134055	
		Mean ΔCt (control)	7.54434			
			Fold gene	expression	1.009853	
			Standard	deviation	0.151056	
		Stand	lard error of	f the mean	0.061668	
		Relative s	tandard erro	or of mean	6.106665	
	22.63096428	15.29334736	7.337617	-0.20672	1.154064	
	22.6428833	15.16924477	7.473639	-0.0707	1.050228	
Female	22.89701462	15.29349518	7.603519	0.059179	0.95981	
remale	23.30892372	15.89090347	7.41802	-0.12632	1.091506	
	24.46821022	15.78082657	8.687384	1.143043	0.452803	
	24.78164673	16.44277382	8.338873	0.794532	0.57653	
	Fold gene expression					
Standard deviation					0.293183	
	Standard error of the mean					
Relative standard error of mean					13.58856	

Table S.6.9: Cyp2c13 mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2c13</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.72068977	15.0469017	6.673788	0.018677	0.987137	
	22.19749451	15.63680172	6.560693	-0.09442	1.067635	
Male	22.58212471	15.80290794	6.779217	0.124106	0.917572	
iviale	22.21775627	15.58490181	6.632854	-0.02226	1.015546	
	22.33726501	15.65109253	6.686172	0.031062	0.9787	
	22.15173149	15.553792	6.597939	-0.05717	1.040424	
		Mean ΔCt (control)	6.655111			
			Fold gene	expression	1.001169	
			Standard	d deviation	0.052641	
		Stand	lard error of	f the mean	0.021491	
		Relative s	tandard erro	or of mean	2.146557	
	22.2999115	15.26576138	7.03415	0.379039	0.768949	
	22.45824814	15.58676243	6.871486	0.216375	0.860725	
Female	22.44109535	15.72859859	6.712497	0.057386	0.961004	
remale	22.56281853	15.84080791	6.722011	0.0669	0.954687	
	24.93540382	15.89041996	9.044984	2.389873	0.190799	
	24.06513596	16.63667107	7.428465	0.773354	0.585056	
	Fold gene expression					
Standard deviation					0.294491	
	Standard error of the mean					
Relative standard error of mean					16.69324	

Table S.6.10: Cyp2c23 mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2c23</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)		
	35.11530304	14.99317646	20.12213	-0.0195	1.01361		
	35.56125259	15.53057957	20.03067	-0.11096	1.079944		
Male	35.26271439	15.73916912	19.52355	-0.61808	1.534835		
iviale	35.98874283	15.41041851	20.57832	0.436695	0.738825		
	36.56649399	15.53924561	21.02725	0.885619	0.541255		
	34.84709549	15.27923965	19.56786	-0.57377	1.488411		
		Mean ΔCt (control)	20.14163				
			Fold gene	expression	1.066147		
			Standard	l deviation	0.395873		
		Stand	lard error of	the mean	0.161614		
		Relative s	tandard erro	or of mean	15.15875		
	32.34098816	15.25453281	17.08646	-3.05517	8.311873		
	33.9764595	15.24050331	18.73596	-1.40567	2.649413		
Female	37.15181351	15.51179123	21.64002	1.498393	0.353947		
remale	36.91116714	15.64538002	21.26579	1.124158	0.45877		
	Fold gene expression						
Standard deviation					3.732094		
	Standard error of the mean						
Relative standard error of mean					63.3955		

Table S.6.11: *Cyp2e1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2e1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.00891876	15.0469017	5.962017	0.034478	0.976385	
	21.61394882	15.63680172	5.977147	0.049608	0.966199	
Male	21.91149902	15.80290794	6.108591	0.181052	0.882059	
IVIdle	21.3314724	15.58490181	5.746571	-0.18097	1.133644	
	21.53865433	15.65109253	5.887562	-0.03998	1.028097	
	21.4371376	15.553792	5.883346	-0.04419	1.031106	
		Mean ΔCt (control)	5.927539			
			Fold gene	expression	1.002915	
			Standard	deviation	0.083877	
		Stand	lard error of	f the mean	0.034243	
		Relative s	tandard erro	or of mean	3.414328	
	21.18091965	15.26576138	5.915158	-0.01238	1.008619	
	22.1549778	15.58676243	6.568215	0.640676	0.641412	
Female	21.76470566	15.72859859	6.036107	0.108568	0.927508	
remale	21.67921638	15.84080791	5.838408	-0.08913	1.063729	
	21.79675484	15.89041996	5.906335	-0.0212	1.014806	
	23.31801414	16.63667107	6.681343	0.753804	0.593038	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					9.559831	

Table S.6.12: *Cyp2j3* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2j3</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.20621109	15.0469017	6.159309	-0.10035	1.072032	
	21.78096199	15.63680172	6.14416	-0.1155	1.083349	
Male	22.99167442	15.80290794	7.188766	0.929109	0.525183	
iviale	21.65578842	15.58490181	6.070887	-0.18877	1.139793	
	21.70969009	15.65109253	6.058598	-0.20106	1.149543	
	21.49001884	15.553792	5.936227	-0.32343	1.251303	
		Mean ΔCt (control)	6.259658			
			Fold gene	expression	1.036867	
			Standard	d deviation	0.258647	
		Stand	lard error of	f the mean	0.105592	
		Relative s	tandard erro	or of mean	10.18376	
	21.29408836	15.26576138	6.028327	-0.23133	1.173917	
	21.67180824	15.58676243	6.085046	-0.17461	1.128661	
Female	21.8919754	15.72859859	6.163377	-0.09628	1.069014	
remale	21.99642372	15.84080791	6.155616	-0.10404	1.074781	
	22.43022537	15.89041996	6.539805	0.280148	0.823507	
	23.48759651	16.63667107	6.850925	0.591268	0.663759	
	Fold gene expression					
	Standard deviation					
		Stand	lard error of	f the mean	0.081778	
Relative standard error of mean					8.269266	

Table S.6.13: *Cyp2j4* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2j4</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.63014984	14.90790844	6.722241	-0.24304	1.183481	
	22.43364143	15.53313255	6.900509	-0.06477	1.045917	
Mala	23.24939537	16.04916191	7.200233	0.234956	0.849711	
Male	23.12949944	15.88101864	7.248481	0.283203	0.821765	
	22.36303902	15.43636131	6.926678	-0.0386	1.027117	
	22.52326584	15.7297411	6.793525	-0.17175	1.126426	
		Mean ΔCt (control)	6.965278			
			Fold gene	expression	1.00907	
			Standard	deviation	0.145845	
		Stand	lard error o	f the mean	0.059541	
		Relative s	tandard err	or of mean	5.900578	
	22.15753555	15.31814289	6.839393	-0.12589	1.091177	
	22.18677902	15.23503971	6.951739	-0.01354	1.009428	
Female	22.87580109	15.87814617	6.997655	0.032377	0.977808	
remaie	23.33951378	16.2695446	7.069969	0.104691	0.930004	
	23.10550117	16.11947632	6.986025	0.020747	0.985722	
	22.6664238	16.50775146	6.158672	-0.80661	1.749091	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					11.29088	

Table S.6.14: *Cyp2j10* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp2j10</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	31.6193409	14.90790844	16.71143	1.209574	0.432396	
	30.89755058	15.53313255	15.36442	-0.13744	1.099952	
Mala	30.9431057	16.04916191	14.89394	-0.60791	1.524055	
iviale	30.86358452	15.88101864	14.98257	-0.51929	1.433252	
	30.954216	15.43636131	15.51785	0.015996	0.988974	
	31.27067757	15.7297411	15.54094	0.039078	0.973277	
		Mean ΔCt (control)	15.50186			
			Fold gene	expression	1.075318	
			Standard	deviation	0.390206	
		Stand	lard error of	f the mean	0.159301	
		Relative s	tandard erro	or of mean	14.81432	
	31.29132462	15.31814289	15.97318	0.471323	0.721303	
Male 30.86358452 15.88101864 14.9825 30.954216 15.43636131 15.5178 31.27067757 15.7297411 15.5409	16.28966	0.787802	0.579226			
Formula	31.27067757 15.7297411 15.54094 0.039078 0 Mean ΔCt (control) 15.50186 Fold gene expression 1 Standard deviation 0 Standard error of the mean 1 Relative standard error of mean 1 31.29132462 15.31814289 15.97318 0.471323 0 31.52470016 15.23503971 16.28966 0.787802 0 31.03298569 15.87814617 15.15484 -0.34702 15	1.27193				
Female	31.09254265	16.2695446	14.823	-0.67886	1.600875	
	31.11616707	16.11947632	14.99669	-0.50517	1.419288	
	31.57313728	16.50775146	15.06539	-0.43647	1.353292	
	Fold gene expression					
Standard deviation					0.410174	
Standard error of the mean					0.167453	
Relative standard error of mean					14.46485	

Table S.6.15: *Cyp3a1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp3a1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	35.7577095	15.53313255	20.22458	-0.62844	1.545896	
Male	37.41846466	16.04916191	21.3693	0.516283	0.699171	
iviale						
	37.07433701	15.43636131	21.63798	0.784956	0.58037	
	35.90996552	15.7297411	20.18022	-0.6728	1.594159	
		Mean ΔCt (control)	20.85302			
			Fold gene	expression	1.104899	
			Standard	d deviation	0.539629	
		Stand	dard error o	f the mean	0.269815	
		Relative s	tandard err	or of mean	24.41985	
	36.60176849	15.31814289	21.28363	0.430606	0.74195	
	36.75820923	15.23503971	21.52317	0.67015	0.628442	
EI-	36.77846909	15.87814617	20.90032	0.047303	0.967744	
Female						
	37.16514969	16.11947632	21.04567	0.192653	0.874995	
	36.38945007	16.50775146	19.8817	-0.97132	1.960635	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean				0.238547	
	Relative standard error of mean					

Table S.6.16: *Cyp3a2* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp3a2</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	25.19059181	14.90790844	10.28268	0.050117	0.965858	
	25.76757813	15.53313255	10.23445	0.001879	0.998698	
Mala	26.27242661	16.04916191	10.22326	-0.0093	1.006468	
Male	26.12583733	15.88101864	10.24482	0.012252	0.991543	
	25.74394035	15.43636131	10.30758	0.075013	0.949334	
	25.83234787	15.7297411	10.10261	-0.12996	1.094263	
		Mean ΔCt (control)	10.23257			
			Fold gene	expression	1.001027	
			Standard	d deviation	0.050452	
		Stand	lard error o	f the mean	0.020597	
		Relative s	tandard err	or of mean	2.057577	
	25.70596695	15.31814289	10.38782	0.155258	0.897972	
	25.78372765	15.23503971	10.54869	0.316122	0.803226	
Female	26.11230087	15.87814617	10.23415	0.001588	0.9989	
remale	26.5971489	16.2695446	10.3276	0.095038	0.936248	
	28.95790291	16.11947632	12.83843	2.60586	0.16427	
	28.3625946	16.50775146	11.85484	1.622277	0.324822	
	Fold gene expression					
Standard deviation					0.352659	
	Standard error of the mean					
Relative standard error of mean					20.93921	

Table S.6.17: *Cyp3a9* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp3a9</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)
	37.20720291	15.63680172	21.5704	1.199007	0.435575
Male	35.86162949	15.80290794	20.05872	-0.31267	1.242007
iviale					
	34.38830566	15.65109253	18.73721	-1.63418	3.104114
	36.67303467	15.553792	21.11924	0.747848	0.595491
		Mean ΔCt (control)	20.37139		
			Fold gene	expression	1.344297
			Standard	d deviation	1.223915
		Stand	dard error o	f the mean	0.611958
		Relative s	tandard err	or of mean	45.5225
	26.90587044	15.26576138	11.64011	-8.73129	424.9901
	31.63671494	15.58676243	16.04995	-4.32144	19.99326
Female	36.15457535	15.72859859	20.42598	0.054582	0.962873
Female	36.53848648	15.84080791	20.69768	0.326284	0.797588
	26.74207306	15.89041996	10.85165	-9.51974	734.0536
			Fold gene	expression	236.1595
	Standard deviation				
		Stand	dard error o	f the mean	148.4918
Relative standard error of mean					62.87774

Table S.6.18: *Cyp3a23* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp3a23</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.01748848	14.90790844	6.10958	-0.03687	1.025887	
	21.60243797	15.53313255	6.069305	-0.07715	1.054929	
Male	22.36790657	16.04916191	6.318745	0.172293	0.887431	
iviale	22.15455437	15.88101864	6.273536	0.127084	0.91568	
	21.68592644	15.43636131	6.249565	0.103113	0.931022	
	21.58772087	15.7297411	5.85798	-0.28847	1.221346	
		Mean ΔCt (control)	6.146452			
			Fold gene	expression	1.006049	
			Standard	deviation	0.124098	
		Stand	lard error of	f the mean	0.050663	
		Relative s	tandard erro	or of mean	5.035812	
	21.42850304	15.31814289	6.11036	-0.03609	1.025332	
	21.44103432	15.23503971	6.205995	0.059543	0.959568	
Female	21.84907722	15.87814617	5.970931	-0.17552	1.129372	
remaie	22.43259048	16.2695446	6.163046	0.016594	0.988564	
	22.30660629	16.11947632	6.18713	0.040678	0.972198	
	24.13162231	16.50775146	7.623871	1.477419	0.359131	
	Fold gene expression					
Standard deviation					0.274689	
	Standard error of the mean					
Relative standard error of mean					12.38179	

Table S.6.19: *Cyp4a1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (Cyp4a1)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	22.4458313	15.0469017	7.39893	0.039977	0.972671	
	23.17233849	15.63680172	7.535537	0.176584	0.884796	
Male	23.34290695	15.80290794	7.539999	0.181046	0.882063	
iviale	22.71615982	15.58490181	7.131258	-0.22769	1.170962	
	22.98096848	15.65109253	7.329876	-0.02908	1.020359	
	22.77190971	15.553792	7.218118	-0.14084	1.102543	
		Mean ΔCt (control)	7.358953			
			Fold gene	expression	1.005566	
			Standard	d deviation	0.116533	
		Stand	lard error of	f the mean	0.047574	
		Relative s	tandard erro	or of mean	4.731117	
	22.63911819	15.26576138	7.373357	0.014404	0.990066	
	22.82960129	15.58676243	7.242839	-0.11611	1.083812	
Female	23.07598114	15.72859859	7.347383	-0.01157	1.008052	
remaie	23.06615639	15.84080791	7.225348	-0.1336	1.097031	
	22.95615959	15.89041996	7.06574	-0.29321	1.225366	
	24.61741447	16.63667107	7.980743	0.621791	0.649864	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					7.876582	

Table S.6.20: *Cyp4a8* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp4a8</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	35.37493515	15.63680172	19.73813	-0.31859	1.247111	
Mala	36.12532043	15.80290794	20.32241	0.265689	0.831802	
Male	35.63024521	15.58490181	20.04534	-0.01138	1.007919	
	36.1544342	15.65109253	20.50334	0.446618	0.733761	
	35.22817993	15.553792	19.67439	-0.38234	1.303451	
	<u>'</u>	Mean ΔCt (control)	20.05672			
			Fold gene	expression	1.024809	
			Standard	deviation	0.249653	
		Stand	lard error of	f the mean	0.111648	
		Relative s	tandard erro	or of mean	10.89452	
	36.65815735	15.58676243	21.07139	1.014671	0.494941	
Family	35.93002319	15.72859859	20.20142	0.144701	0.904567	
Female	35.30750656	15.84080791	19.4667	-0.59003	1.505273	
	35.41714096	15.89041996	19.52672	-0.53	1.443932	
	36.95736313	16.63667107	20.32069	0.263968	0.832794	
	Fold gene expression					
	Standard deviation					
		Stand	lard error of	f the mean	0.192086	
Relative standard error of mean					18.53568	

Table S.6.21: *Cyp4f1* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp4f1</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	25.97480392	14.37320328	11.6016	-1.28546	2.437598	
Male	27.30628777	14.60536575	12.70092	-0.18614	1.137715	
	28.27339172	15.02254772	13.25084	0.363783	0.777124	
	28.93395424	14.93907642	13.99488	1.107817	0.463996	
		Mean ΔCt (control)	12.88706			
			Fold gene	expression	1.204108	
			Standard	d deviation	0.867177	
		Stand	dard error o	f the mean	0.433589	
		Relative s	tandard erre	or of mean	36.0091	
	24.17294121	14.99263287	9.180308	-3.70675	13.05701	
	26.8288784	14.84191036	11.98697	-0.90009	1.866186	
Female	28.83596992	15.01529312	13.82068	0.933616	0.523545	
remate						
	24.56740379	15.37995338	9.18745	-3.69961	12.99253	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean					
Relative standard error of mean					48.18695	

Table S.6.22: *Cyp4f4* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp4f4</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)
	22.38262939	14.37320328	8.009426	-0.32184	1.249928
	23.09357071	14.77818966	8.315381	-0.01589	1.011075
Male	22.99298477	14.57935047	8.413634	0.082363	0.944509
iviale	22.9574337	14.60536575	8.352068	0.020797	0.985688
	23.49173927	15.02254772	8.469192	0.137921	0.908828
	23.36700058	14.93907642	8.427924	0.096653	0.9352
		Mean ΔCt (control)	8.331271		
			Fold gene	expression	1.005871
			Standard	l deviation	0.12504
		Stand	lard error of	the mean	0.051047
		Relative s	tandard erro	or of mean	5.074945
	23.43267632	14.99263287	8.440043	0.108773	0.927377
	23.02902031	14.84191036	8.18711	-0.14416	1.105088
Female	23.29996872	15.01529312	8.284676	-0.0466	1.032825
remale	23.8758812	15.52533817	8.350543	0.019272	0.98673
	23.9055233	15.37995338	8.52557	0.194299	0.873997
	25.44578171	16.05649185	9.38929	1.058019	0.480291
			Fold gene	expression	0.901051
Standard deviation					0.221269
		Stand	lard error of	the mean	0.090333
		Relative s	tandard erro	or of mean	10.02526

Table S.6.23: *Cyp4f5* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp4f5</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)	
	21.47459793	15.0469017	6.427696	0.402737	0.756422	
	21.69128036	15.63680172	6.054479	0.029519	0.979747	
Male	22.37661552	15.80290794	6.573708	0.548748	0.683613	
ividle	21.22067833	15.58490181	5.635777	-0.38918	1.309652	
	21.40433693	15.65109253	5.753244	-0.27172	1.207242	
	21.25864601	15.553792	5.704854	-0.32011	1.248422	
		Mean ΔCt (control)	6.02496			
			Fold gene	expression	1.03085	
			Standard	d deviation	0.266337	
		Stand	lard error o	f the mean	0.108732	
		Relative s	tandard err	or of mean	10.54778	
	21.15799141	15.26576138	5.89223	-0.13273	1.096366	
	21.41794205	15.58676243	5.83118	-0.19378	1.143757	
Female	21.77904892	15.72859859	6.05045	0.025491	0.982486	
remale	21.6143589	15.84080791	5.773551	-0.25141	1.190369	
	21.25339699	15.89041996	5.362977	-0.66198	1.582255	
	21.94256592	16.63667107	5.305895	-0.71906	1.646115	
	Fold gene expression					
	Standard deviation					
	Standard error of the mean				0.111651	
Relative standard error of mean					8.766888	

Table S.6.24: *Cyp4f6* mRNA expression in the small intestine of male and female Sprague Dawley rats normalized to GAPDH housekeeping gene

Group	Ct (<i>Cyp4f6</i>)	Ct (GAPDH)	ΔCt	ΔΔCt	2^- (ΔΔCt)
Male	28.59749222	15.0469017	13.55059	-0.49322	1.407582
	29.52577782	15.58490181	13.94088	-0.10293	1.073955
	29.70816994	15.65109253	14.05708	0.013268	0.990845
	30.18048477	15.553792	14.62669	0.582884	0.667628
Mean ΔCt (control) 14.04381					
Fold gene expression					1.035002
Standard deviation					0.303999
Standard error of the mean					0.152
Relative standard error of mean					14.68591
Female	27.54456329	15.26576138	12.2788	-1.76501	3.398757
	29.60769844	15.58676243	14.02094	-0.02287	1.015981
	30.53573418	15.72859859	14.80714	0.763326	0.589136
	27.59592628	15.89041996	11.70551	-2.3383	5.057074
Fold gene expression					2.515237
Standard deviation					2.097552
Standard error of the mean					1.048776
Relative standard error of mean					41.69691