

4 Sex Differences in Drug Metabolism

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4.1 SUMMARY

Sex differences in drug metabolism are recognized as a major determinant of sex differences in pharmacokinetics and an important contributor to interindividual differences in drug metabolism and, in some cases, drug action. Historically, sex differences in drug metabolism were widely studied in rat and mouse models in the 1960s and 1970s. Sex differences in human drug metabolism have also been observed with a variety of drugs; however, the quantitative differences in rates of metabolism between men and women are typically much smaller than those seen in rats. The biochemical basis of sex differences in hepatic drug metabolism was largely unknown until the 1980s when it was discovered that expression of some of the drug-metabolizing enzymes, including those in the cytochrome P450 (CYP) family, is sex-dependent in mouse and rat liver. Thus, the expression of certain rat CYP enzymes is male specific (e.g., CYP2C11), whereas others are female specific (e.g., CYP2C12) or female predominant (e.g., CYP2A1). The major gonadal hormones, testosterone and estradiol, regulate the sex-dependent expression of rat hepatic CYP enzymes indirectly through their effects on the hypothalamus and its regulation of the sexually dimorphic, ultradian

rhythm of pituitary growth hormone (GH) secretion, which ultimately dictates the sex-dependent patterns of expression of select CYPs and certain other drug-metabolizing enzymes. In male rats, the intermittent, pulsatile pattern of GH secretion stimulates the expression of male-specific liver enzymes, whereas the more continuous pattern of GH release in females suppresses the expression of male-specific enzymes while inducing female-specific enzymes. Studies conducted in the past decade have identified essential molecular mediators of the effects of GH on sex-dependent liver CYP enzymes and their genes, in particular, the transcription factors—signal transducer and activator of transcription (STAT) 5b and hepatocyte nuclear factors (i.e., HNF4 α). The current view is that STAT5b and HNF4 α coordinately regulate the action of GH and its resultant effect on the sex-dependent expression of hepatic *CYP* genes and thus provide a molecular basis for sex differences in drug metabolism.

4.2 INTRODUCTION

Individual variability in drug metabolism and drug response is well documented in the biomedical literature [1–3]. This variability is manifested as qualitative and/or quantitative differences in drug response within a group of patients (i.e., interindividual variability) or in the same patient (i.e., intraindividual variability). Within a population, patient-specific characteristics such as sex [4], age [5], and genetics [6,7] have been identified as determinants of interindividual variability in drug action. Intraindividual differences in drug action can be induced by administration of a second drug leading to drug–drug interactions, may be caused by deterioration of organ function, or may be by progression in the severity of a disease state, among others [8]. The pharmacological basis for variability in drug response can be attributed to interindividual differences in pharmacokinetics (i.e., drug absorption, distribution, metabolism, and excretion) and/or pharmacodynamics (i.e., the physiological, biochemical, cellular, and molecular actions of a drug). Indeed, patient-specific characteristics are known to influence the biological processes that govern drug absorption, distribution, metabolism, excretion, and functionality of therapeutic targets (e.g., receptors, enzymes) and their associated signal transduction pathways.

Sex differences in drug action were first observed in the 1930s, where adult female rats were found to require only half the dosage of amobarbital (Amytal), as compared to adult male rats, to elicit an anesthetic effect [9], and the duration of action of certain barbiturates (e.g., amobarbital and pentobarbital) was found to be longer in female than in male rats [10]. Subsequent animal and human studies have shown sex differences in the pharmacokinetics and pharmacodynamics of various other drugs [11–15]. Considerably more is known about sex differences in pharmacokinetics, especially sex differences in drug metabolism in the liver than about sex differences in pharmacodynamics. The overall aim of this chapter is to provide an overview of our current understanding of sex differences in hepatic drug metabolism.

4.3 SEX DIFFERENCES IN HEPATIC DRUG METABOLISM

4.3.1 Rat Hepatic Drug Metabolism

Sex differences in hepatic drug metabolism were widely studied in the 1960s and 1970s in experiments involving the incubation of rat hepatic microsomes (liver membrane

TABLE 4.1 Metabolism of Drugs and Other Substrates Catalyzed by Hepatic Microsomes Isolated from Male and Female Rats

Drug/ Substrate	Reaction	Strain of Rat	Sex Difference (Male/Female Ratio)	References
5 α -androstane- 3 α ,17 β -diol-3, 17-disulfate	15 β -Hydroxylation	Long-Evans	<0.05	21
Aminopyrine	N-demethylation	Sprague-Dawley ^a	3.5 4.0	22 23
Aniline	Hydroxylation	Sprague-Dawley ^a	1.1	22
Diazepam	3-Hydroxylation	Wistar ^a Wistar ^b BD ^a	5.2 0.7 10	16 16 16
	N-desmethylation	Wistar ^a Wistar ^b BD ^a	0.9 0.9 2.6	16 16 16
Diphenhydramine	N-demethylation	Sprague-Dawley ^a	2.4	23
Hexobarbital	Hydroxylation	Sprague-Dawley ^a	4.2 3.0	22 23
Imipramine	N-demethylation	Sprague-Dawley ^a	2.5	24
	2-Hydroxylation	Sprague-Dawley ^a	0.7	24
	N-oxidation	Sprague-Dawley ^a	2.9	24
Indinavir	Oxidation ^c	Sprague-Dawley ^a	3.3	25
Lidocaine	N-demethylation	Sprague-Dawley ^a	8.5	24
	3-Hydroxylation	Sprague-Dawley ^a	0.7	24
Meperidine	N-demethylation	Sprague-Dawley ^a	2.4	23
3,4-Methylenedioxy- methamphetamine	N-demethylation	Sprague-Dawley ^a	3.3	18
Morphine	N-demethylation	Sprague-Dawley ^a	2.6	23
Pentobarbital	Oxidation	Sprague-Dawley ^a	2.6	23

^aPostpubertal rats.^bNeonatal rats.^cTotal oxidation products.

preparations enriched in endoplasmic reticulum) with drugs, other foreign chemicals, or endogenous lipophilic substrates (e.g., endogenous steroids and fatty acids). As illustrated in Table 4.1, drugs such as diazepam, lidocaine, and indinavir are metabolized more rapidly by liver microsomes isolated from postpubertal male rats than by those from postpubertal female rats. In contrast, other substrates (e.g., the steroid 5 α -androstane-3 α ,17 β -diol-3,17-disulfate) are more rapidly metabolized by hepatic microsomes from female rats. However, not all substrates (e.g., aniline) exhibit sex difference in hepatic metabolism. Table 4.1 also shows that the sex dependence of drug metabolism can be pathway specific; for example, lidocaine N-demethylation, but not lidocaine 3-hydroxylation, is preferentially catalyzed by hepatic microsomes from male rats. With some drugs, sex differences in hepatic metabolism are strain specific; for example, liver microsomes prepared from BD rats but not those from Wistar rats exhibit sex differences in diazepam N-desmethylation. Sex differences in hepatic drug metabolism may also vary as a function of age (Table 4.1), typically being

manifested during the pubertal and adult periods but not prepubertally [16] and present at a reduced level during senescence [17]. Sex differences in hepatic drug bioactivation can lead to deleterious consequences, as exemplified by the finding that the enhanced N-demethylation of 3,4-methylenedioxymethamphetamine (MDMA, ecstasy) by hepatic microsomes from male rats is associated with an increased susceptibility to the acute toxicity of MDMA, when compared to female rats [18]. In addition to phase I oxidative drug metabolism, sex differences in phase II metabolism affecting glucuronidation, sulfation, and glutathione conjugation, have also been reported in animal studies [19,20].

4.3.2 Human Hepatic Drug Metabolism

Sex differences in human hepatic drug metabolism have been investigated in clinical pharmacokinetic studies that compare the clearance of a drug in men and women. Sex differences characterize the clearance of certain drugs, including cyclosporine [26], erythromycin [27], nifedipine [28], triazolam [29], midazolam [30,31], alprazolam [32], and verapamil [33]. However, in some cases, conflicting data exists with regard to the effect of sex on drug clearance [34–36]. Possible reasons for the conflicting data include (i) the small sample size employed in some of the studies resulting in a lack of statistical power to detect a sex-based difference and (ii) the role of confounding variables, such as age [12,35], ethnicity [37], and other biological determinants of drug clearance, which can include the level of expression of uptake and efflux transporters, although it is debatable whether human hepatic P-glycoprotein expression differs between males and females [38], given the conflicting data in the literature [39,40]. With the increased availability of human hepatic microsomes for research, the *in vitro* metabolism of drugs and other substrates can readily be assessed using microsomes isolated from liver tissue samples obtained from male and female donors. As shown in Table 4.2, in general, sex differences are not apparent or are relatively modest in human hepatic microsomal drug metabolism. An exception is *S*-mephenytoin N-demethylation, whose metabolism is substantively greater in hepatic microsomes from females than in those from males, but this occurs only in samples from Hispanic donors [37]. Overall, sex differences occur in hepatic phase I and phase II drug metabolism, and the differences are generally larger and hence more apparent in rats than in humans.

4.4 SEX-DEPENDENT EXPRESSION OF HEPATIC DRUG-METABOLIZING ENZYMES

The underlying biochemical basis for sex differences in drug metabolism came to light when it was discovered in the 1980s that many of the rat hepatic CYP enzymes (Table 4.3) are expressed in a sex-dependent manner [47]. Since then, sex-dependent expression of other rodent drug-metabolizing enzymes, including UDP-glucuronosyltransferases [48–50], sulfotransferases [49,51,52], and glutathione *S*-transferases [49,53,54], has also been reported. The available experimental evidence suggests that certain human hepatic CYP enzymes may also be expressed differentially in men and women. Overall, among the drug-metabolizing enzymes, the hepatic CYPs are the best characterized with respect to their sex-dependent expression. Thus, the

TABLE 4.2 Effect of Sex on Metabolism of Drugs and Other Substrates Catalyzed by Human Hepatic Microsomes

Drug	Reaction	Effect	References
Aminopyrine	N-demethylation	M = F	41
Chlorzoxazone	6-Hydroxylation	M = F	42
Coumarin	7-Hydroxylation	M = F	42
Dextromethorphan	O-demethylation	M = F	42
Diclofenac	4'-Hydroxylation	M = F	42
Erythromycin	N-demethylation	F > M (1.2-fold)	43
7-Ethoxyresorufin	O-demethylation	M = F	42
Ifosfamide	N-dechloroethylation	F > M (2.2-fold)	44
Lauric acid	12-Hydroxylation	M = F	42
S-mephenytoin	N-demethylation	F > M (7.8-fold) ^a	37
		M = F ^{b,c}	37
		M = F ^d	42
	4'-Hydroxylation	M = F	42
Nicotine	C-oxidation	M = F	45
S-oxazepam	Glucuronidation	M > F (1.7-fold)	46
Paclitaxel	6 α -Hydroxylation	M = F	42
Testosterone	6 β -Hydroxylation	M = F	42

^aHispanic.^bAfrican-American.^cCaucasian.^dSamples from Hispanics, African-Americans, and Caucasians.

remainder of this chapter focuses on hepatic CYP enzymes and the various factors that regulate their sex-dependent expression.

4.4.1 Rat Cytochrome P450s

As summarized in Table 4.3, multiple CYP enzymes are expressed constitutively in a sex-dependent manner in rat liver, and their expression is developmentally regulated. Interestingly, a catalytic function of many of these enzymes is the oxidative metabolism of sex steroid hormones such as testosterone. The prototype male-specific P450 enzyme of rat liver is CYP2C11 [55–57, 75]. CYP2C11 is essentially undetectable in both male and female rat liver before puberty, but its expression increases rapidly at puberty (at approximately day 35 of age) in males only and reaches a maximum during early adulthood [58,59]. A similar pattern of postnatal developmental expression has been described for several other male-specific rat hepatic CYP enzymes, that is, CYP2A2 [60,61], CYP2C13 [62], and CYP3A18 [63,64]. In contrast, hepatic expression of the male-specific CYP3A2 is detectable during the prepubertal period in both sexes, whereas at the onset of puberty, the level of CYP3A2 is increased in male liver, but at about the same time it decreases to barely quantifiable levels in females [58]. Another sex-dependent rat hepatic CYP is the female-specific CYP2C12 (Table 4.3). Although CYP2C12 is expressed at low levels in the livers of both sexes prepubertally, its expression is increased at puberty in female rat liver but is suppressed in male rats [58,65]. The female-predominant enzymes CYP2A1, CYP2A7, and CYP3A9 (Table 4.3) comprise a third group of sex-dependent liver P450s. Expression is either similar (CYP2A1

TABLE 4.3 Sex-Dependent Hepatic Expression of Rat Cytochrome P450 Enzymes^a

Enzyme	Catalytic Reaction	Developmental Expression	
		Prepuberty	Puberty
<i>Male Specific</i>			
CYP2A2	Testosterone 15 α -hydroxylation	None	↑ in male
CYP2C11	Testosterone 2 α -hydroxylation	None	↑ in male
CYP2C13	Testosterone 15 α -hydroxylation	Minimal	↑ in male
CYP3A2	Testosterone 6 β -hydroxylation	Male ~ female	↑ in male, ↓ in female
CYP3A18	Testosterone 6 β -hydroxylation	Minimal	↑ in male
CYP4A2	Lauric acid ω -hydroxylation	None ^b	↑ in male ^b
<i>Female Specific</i>			
CYP2C12	5 α -Androstane-3 α ,17 β -diol-3,17-disulfate 15 β -hydroxylation	Minimal/moderate	↑ in female None in male
<i>Female Predominant</i>			
CYP2A1	Testosterone 7 α -hydroxylation	Male ~ female	Female > male
CYP2C7	Retinoic acid 4-hydroxylation	Male ~ female	Female > male
CYP3A9	Testosterone 6 β -hydroxylation	↓	Female > male

^aBased on Refs 21,55–73.^bBased on data for renal CYP4A2 [74].

and CYP2C7) or absent (CYP3A9) in both sexes during prepuberty, and the levels of these enzymes are greater in postpubertal female rats than in postpubertal male rats [58,62,64].

4.4.2 Human Cytochrome P450s

In contrast to the rat CYP enzymes, relatively little is known about the sex-dependent expression of human CYPs. With the available information in the literature, many of the conclusions have been drawn from *in vivo* pharmacokinetic studies that measured the plasma clearance of an enzyme-selective substrate (e.g., midazolam for CYP3A enzymes [31]) or urinary formation of a biomarker (e.g., 6 β -hydroxycortisol for CYP3A enzymes [76]). In other cases, conclusions regarding the sex dependence of human enzyme levels have been drawn from *in vitro* metabolism studies (Table 4.2) based on the conversion of an enzyme-selective substrate to its metabolite (e.g., microsomal formation of 6 β -hydroxytestosterone as a catalytic marker for CYP3A [42]). More limited information is available on the immunoreactive protein levels of human CYP enzymes in livers from both sexes. In a panel of 94 liver samples (48 from males and 46 from females), the median level of CYP3A4 was approximately twofold greater in liver samples from females than in samples from males [40]. By comparison, CYP2B6 protein levels were found to be approximately twofold higher in livers from Caucasian women ($N = 15$) than from Caucasian men ($N = 28$), whereas it was approximately fourfold higher in liver samples from Hispanic women ($N = 3$) than in those from Hispanic men ($N = 4$) [37]. In another study of human livers from donors of unknown ethnic origin, multiple linear regression analysis indicated that sex was a

predictor of hepatic CYP2B6 protein levels, but only accounted for 14% of the variance [77]. Overall, it appears that sex differences occur in hepatic expression of human CYP, namely, CYP3A4 and CYP2B6, although the effects are rather modest and considerably less pronounced when compared to the sex differences in the expression of rat hepatic CYP enzymes (Table 4.3).

4.5 HORMONAL DETERMINANTS OF SEX-DEPENDENT EXPRESSION OF HEPATIC CYTOCHROME P450s

A hormonal basis for sex differences in drug metabolism was apparent from early studies showing that (i) castration of postpubertal male rats decreased hepatic microsomal CYP-mediated metabolism of aminopyrine and hexobarbital and (ii) exogenous administration of an androgen (methyltestosterone) fully reversed the suppressive effects of castration [22,78]. These observations paved the way for subsequent studies by various research groups elucidating the hormonal factors that determine the sex-dependent expression of CYP enzymes in rat liver. Collectively, the results of those studies indicate that in addition to gonadal hormones, other hormones also regulate the sex-dependent expression of rat hepatic CYPs. Among these, GH is the most important and the best characterized [79].

4.5.1 Gonadal Hormones

4.5.1.1 Androgen. Androgen has distinct effects on the hepatic expression of the male-specific and female-specific rat CYP enzymes. In male rats, androgen is required for the pubertal expression of the male-specific forms, such as CYP2A2 [60], CYP2C11 [55,58,59,80], CYP2C13 [66], CYP3A2 [58,81], CYP3A18 [82], and CYP4A2 [67], as shown in experiments involving castration of neonatal male rats and exogenous administration of testosterone. Thus, exposure to androgen early in life (e.g., neonatal period) is sufficient to ensure enzyme expression later in adult life in a process known as *neonatal programming* [83]. Full adult expression levels of the male-specific enzymes are achieved in rats exposed to androgen during both the neonatal and peripubertal periods [47]. In contrast to the positive regulation of the male-specific rat hepatic CYP enzymes by androgen, this hormone negatively regulates the hepatic expression of the female-specific CYP2C12, as shown by the decreased enzyme levels in intact, adult female rats administered with exogenous androgen [65,84]. An important conclusion from the available experimental evidence [47,85] is that androgen controls hepatic expression of the sex-dependent rat hepatic CYP enzymes in an indirect manner through its action on the hypothalamic pituitary axis, which results in a sex-specific pattern of pituitary GH secretion (Section 4.5.2).

4.5.1.2 Estrogen. Estrogen also has distinct effects on the expression of sex-dependent rat liver CYP enzymes. In female rats, ovariectomy and exogenous estrogen replacement experiments show that estrogen is required for liver expression of the female-specific CYP2C12 [58,65] and the female-predominant CYP2C7 [80] and CYP3A9 [86]. However, estrogen is not absolutely required for hepatic expression of these enzymes, as these enzymes are still expressed, albeit at low levels, in ovariectomized rats. In intact and castrated male rats, estrogen suppresses the hepatic

expression of the male-specific CYP2C11 [59,80,84]. The effects of estrogen on these sex-dependent CYP enzymes also involves its effects on the sex-specific pattern of pituitary GH secretion [47,85].

4.5.2 Growth Hormone

GH is a 191-amino-acid protein hormone that is secreted by the anterior pituitary gland. Its secretion is rhythmic and episodic over a 24-h period in mammals, including rats [87] and humans [88]. The ultradian rhythm of pituitary GH secretion can be influenced by various factors, including age [89], prolonged food deprivation [90], and sex steroids, particularly, estrogen [91]. There is a distinct sexual dimorphic pattern of GH secretion, which becomes apparent at the onset of puberty [92]. In adult male rats, the plasma GH profile is characterized by well-defined peaks (up to 250 ng GH/mL) and troughs where the levels are very low or undetectable (Fig. 4.1). The period between peaks typically lasts 3–4 h. By comparison, in female rats, GH is secreted in a more continuous manner such as the plasma GH peaks are lower (75 ng/mL), but circulating hormone levels are nearly always detectable (Fig. 4.1).

An initial clue that GH plays a role in regulating hepatic CYP expression was provided by the findings that GH administration decreased total CYP content and hepatic microsomal metabolism of CYP substrates [94]. It is now widely recognized that the sexual dimorphic pattern of GH secretion is the major hormonal determinant

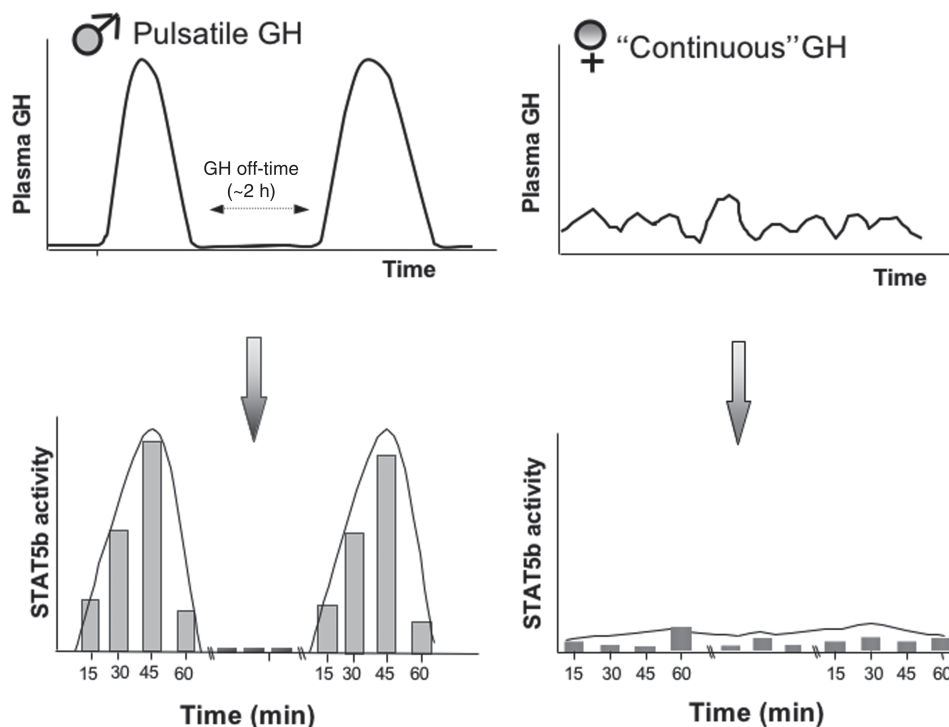


Figure 4.1 Sex dependence of the temporal pattern of plasma GH secretion and hepatic STAT5b activity in male and female rats [93]

TABLE 4.4 Responsiveness of Sex-Dependent Rodent Hepatic Cytochrome P450 Genes to Hypophysectomy and Exogenous Administration of GH^a

Rodent Hepatic Cytochrome P450 Genes	Hypophysectomy		Exogenous Administration of GH	
	Male	Female	Intermittent ^b	Continuous ^c
	Effect on Gene Expression		Effect on Gene Expression	
Rat, male-specific (class I) <i>CYP2C11</i>	↓	↑	↑	↓
Rat, male-specific (class II) <i>CYP2A2</i> , <i>CYP2C13</i> , and <i>CYP3A2</i>	No effect	↑	↓	↓
Rat, female-specific (class I) <i>CYP2C12</i>	↓	↓	↓	↑
Mouse, male-specific (class IA) <i>Cyp2d9</i>	↓	No effect	↑	↓
Mouse, male-specific (class IB) <i>Cyp7b1</i>	↓	↓	↑	↓
Mouse, male-specific (class IC) <i>Cyp4a12</i>	↓	↑	↑	↓
Mouse, female-specific (class II) <i>Cyp2a4</i> and <i>Cyp2b9</i>	↑	No effect	↓	↑

^aAs summarized in Ref. 79.

^bIntermittent GH administration to hypophysectomized male rats.

^cContinuous GH administration of intact male rats.

of the sex-dependent expression of rat hepatic CYPs [47]. On the basis of the available evidence, it appears that the interpulse period devoid of GH is a key signal in the masculinization of CYP expression in rat liver [95,96]. As shown in Table 4.4, the GH-regulated CYP enzymes can be categorized into different classes, based on their response to hypophysectomy [79,127,128]. The expression of a male class I gene (i.e., *CYP2C11*) is decreased in hypophysectomized male rats, whereas it is increased in hypophysectomized female rats but only to a level comparable to that in hypophysectomized male rats. *CYP2C11* levels are increased by intermittent administration of GH and decreased by continuous GH treatment. A corresponding pattern of response is seen for a female class I gene (e.g., *CYP2C12*), except that decreased expression occurs in hypophysectomized female rats and by intermittent GH administration. By comparison, hypophysectomy of female rats increases the expression of class II male-specific genes (i.e., *CYP2A2*, *CYP2C13*, and *CYP3A2*), whereas GH, in particular, continuous infusion of GH in male rats, represses the expression of these genes.

GH regulates the clearance of certain drugs that undergo CYP-mediated clearance [97–99], and also it appears that the sex-dependent pattern of GH secretion controls the expression of human CYP enzymes such as *CYP3A4*, which appears to be expressed at a somewhat greater level in livers from women than in livers from men (Section 4.4.2). As demonstrated in primary cultures of human hepatocytes, continuous GH treatment (to reflect the female pattern of GH secretion) increases *CYP3A4* expression [100,101], whereas intermittent GH administration (i.e., two daily doses, to mimic the pulsatile, male pattern of GH secretion) decreases *CYP3A4* expression, although the effect was

greater in hepatocytes isolated from males than in those from females [101]. The suppressive effect of intermittent GH has also been reported *in vivo*. In a human study conducted in male and female subjects with GH deficiency, GH infusion was associated with an increase in CYP3A activity, whereas GH administered as three daily doses was associated with a decrease in CYP3A activity, as assessed by the erythromycin breath test [102]. In another study conducted in GH-deficient boys and girls, administration of GH once daily for 30 days decreased CYP3A activity, as assessed by the urinary ratio of 6 β -hydroxycortisol/free cortisol, but this response was only observed in boys [103]. Interestingly, the administration of GH by constant infusion over a seven-day period increases CYP3A4 mRNA and protein levels in transgenic male mice [104]. Overall, it appears that the mode of GH administration (intermittent vs continuous) exerts differential effects on human CYP expression in a manner similar to that documented for rat CYP enzymes.

4.6 MOLECULAR DETERMINANTS OF SEX-DEPENDENT EXPRESSION OF HEPATIC CYTOCHROME P450s

Studies in the past decade have increased our understanding of the underlying molecular mechanisms through which GH controls the sex-dependent expression of hepatic CYP enzymes. GH exerts its action by binding to the growth hormone receptor (GHR), which is a member of the superfamily of cytokine receptors [105]. This leads to a conformational change of GHR and activation of a tyrosine kinase known as Janus kinase 2 (JAK2), which catalyzes the tyrosine phosphorylation of various intracellular proteins, including GHR and those belonging to the STAT family, such as STAT5b [106]. The GHR-bound, tyrosine phosphorylated STAT5b dimerizes and translocates to the nucleus where it binds to DNA response elements of target genes and stimulates gene transcription [105,107].

Whereas STAT5b is a direct transcriptional mediator of GH action, GH may also exert indirect effects on gene expression, through various HNFs, including HNF4 α [108], a member of the nuclear receptor superfamily [109]. The available evidence to date indicates that STAT5b and HNF4 α are molecular determinants of the sex-dependent expression of hepatic CYP enzymes [79]. These factors are likely to regulate hepatic CYPs and other sex-specific genes both directly and indirectly, via a complex transcriptional regulatory network, as indicated by the complex time course that characterizes the feminization of these sex-specific genes in male mice infused with GH continuously over a 14-day period [110]. Several other liver-expressed transcription factors show strong sex differences in their expression, including the female-specific factor Cux2 (Cutl2) [111] and the male-specific factor Bcl6 [112]. These factors are both dependent on GH and STAT5b for their sex-specific expression, suggesting that they may play a key role in this transcriptional regulatory network by regulating downstream targets of GH and STAT5b.

4.6.1 STAT5b

A hormonal basis for the tyrosine phosphorylation of STAT5b is provided by the experimental findings indicating that hepatic STAT5b activity is (i) greater in adult

TABLE 4.5 STAT5b and HNF4 α Regulate Sex-Dependent Mouse Hepatic Cytochrome P450 Gene Expression^a

Hepatic Cytochrome P450 Genes	Male		Female	
	STAT5b Deficiency	HNF4 α Deficiency	STAT5b Deficiency	HNF4 α Deficiency
	Effect on Gene Expression			
<i>Male-Specific Genes</i>				
<i>Cyp2d9</i>	↓	↓	↑	↓
<i>Cyp7b1</i>	↓	↓	↓	↓
<i>Cyp4a12</i>	↓	↓	No effect	No effect
<i>Female-Specific Genes</i>				
<i>Cyp2a4</i>	↑	↑	No effect	No effect
<i>Cyp2b9</i>	↑	↑	No effect	No effect

^aBased on Refs 110,118, and 119.

male rats than in adult female rats, (ii) abolished by hypophysectomy, (iii) stimulated by intermittent administration of GH but decreased by continuous administration of GH in hypophysectomized male rats, and (iv) stimulated by continuous GH infusion in hypophysectomized female rats [113,114]. Furthermore, there is a direct temporal relationship between the occurrence of a plasma GH pulse and the activation of hepatic STAT5b activity [115]. As illustrated in Fig. 4.1, in adult male rats, STAT5b is activated by each pulsatile secretion of GH, such that the time for peak STAT5b activity corresponds to the time for peak plasma GH levels and little or no STAT5b activity is detected during the interpulse, GH-free period. In contrast, in adult female rats, there is a very low but persistent level of STAT5b activity over time, similar to the temporal pattern of the low level, continuous release of plasma GH.

STAT5b is essential for the sex-dependent expression of hepatic CYP enzymes and many other sex-dependent genes, as determined in mice genetically deficient in STAT5b. Thus, the sexual dimorphism of hepatic CYP genes is abolished in the absence of STAT5b [116]. As shown in Table 4.5, when compared to the wild-type control male mice, STAT5b-deficient mice have decreased hepatic gene expression of male-specific CYPs (i.e., *Cyp2d9*, *Cyp4a12*, and *Cyp7b1*), but increased levels of female-specific CYPs (i.e., *Cyp2a4*, *Cyp2b9*, and *Cyp2b13*) [110,117]. In female mice, STAT5b deficiency leads to increased hepatic expression of the male-specific *Cyp2d9*, leads to decreases of the male-specific *Cyp7b1*, and has little or no effect on the female-specific *Cyp2a4* or *Cyp2b9* [118]. However, not all sex-dependent CYP genes are regulated by STAT5b because STAT5b deficiency does not influence the hepatic expression of the female-specific *Cyp3a16*, *Cyp3a41*, and *Cyp3a44* [118]. Moreover, hepatic *Cyp2d9* and *Cyp4a12* levels are increased in response to intermittent GH administration in hypophysectomized, STAT5b-expressing male mice but not in hypophysectomized, STAT5b-deficient male mice [110,117]. Collectively, these findings indicate positive regulation of male-specific CYP genes and negative regulation of a subset of female-specific CYP genes by GH pulse-activated STAT5b.

4.6.2 HNF4 α

HNF4 α (gene designation *NR2A1*) is a transcription factor initially known for its role in liver-specific gene expression [120]. It is expressed at high levels in liver and several tissues, and it is constitutively localized in the nucleus [121]. Among its many functions, HNF4 α has been shown to regulate the expression of CYP genes by interacting with various nuclear receptors including constitutive androstane receptor and pregnane X receptor [122,123]. HNF4 α also play a role in hepatic expression of sex-dependent CYP genes [110,124]. As summarized in Table 4.5, HNF4 α deficiency in male mice is associated with a decrease in hepatic levels of the male-specific genes *Cyp2d9*, *Cyp7b1*, and *Cyp4a12*, and an increase in that of the female-specific *Cyp2a4*. By comparison, HNF4 α deficiency in female mice decreases the low level of expression of *Cyp2d9*, whereas it has no effect on the male-specific *Cyp7b1* and *Cyp4a12* and the female-specific *Cyp2a4* and *Cyp2b9*. This pattern of response to HNF4 α deficiency is similar to that of STAT5b deficiency (Table 4.5) and is consistent with the idea that STAT5b is not the sole regulator of the sex-dependent hepatic *Cyp* genes [125,126]. The current view is that HNF4 α functions cooperatively with GH pulse-activated STAT5b to achieve the full expression of male-specific CYP genes [79], a view that is supported by microarray analysis showing global similarities in the dependence of sex-specific genes on STAT5b and HNF4 α in mouse liver [119].

4.7 CONCLUDING REMARKS

Sex differences in hepatic drug metabolism are well documented in rat and mouse models and are increasingly recognized as making a key contribution to sex differences in pharmacokinetics. The underlying mechanisms responsible for the sex differences in the hepatic expression of drug-metabolizing enzymes, in particular, CYPs, are not fully understood but involve complex patterns of hormonal control and intracellular signaling pathways. It is now well established that the sexually dimorphic pattern of plasma GH secretion is a major regulator that controls the sex-dependent expression of CYPs. Studies over the past few years have revealed that sex differences in the liver extend well beyond enzymes of drug metabolism to include more than 1000 genes [127,128], and the transcription factors STAT5b and HNF4 α have been identified as essential determinants of the sex-dependent expression of a large fraction of these genes, including many CYPs. Several other GH-regulated liver transcription factors that show strong sex differences in their expression, as well as a dependence on STAT5b, have also been identified and seem likely to be part of a complex transcriptional regulatory network that regulates the sex-dependent expression of CYPs and many other liver-expressed genes. Much has been learned but much more remains to be elucidated regarding the complex signaling pathways involved in the cellular and molecular actions of GH and how they contribute to sex differences in drug metabolism.

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