

Drug Glucuronidation in Clinical Psychopharmacology

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Glucuronidation is a phase II metabolic process and one of the most common pathways in the formation of hydrophilic drug metabolites. At least 33 families of uridine diphosphate-glucuronosyltransferases have been identified *in vitro*, and specific nomenclature similar to that used to classify the cytochrome (CYP) P450 system has been established. The UGT1 and UGT2 subfamilies represent the most important of these enzymes in human drug metabolism. Factors affecting glucuronidation include the following: cigarette smoking, obesity, age, and gender. In addition, several drugs have been found *in vitro* to be substrates, inhibitors, or inducers of UGT enzymes. Induction or inhibition of both UGT and CYP isoforms may occur simultaneously. Some important drug interactions involving glucuronidation have been documented and others can be postulated. This review summarizes the relevant literature pertaining to drug glucuronidation and its implications for clinical psychopharmacology. (J Clin Psychopharmacol 2001;21:500–515)

THE BIOTRANSFORMATION OF xenobiotics and endogenous compounds is accomplished by a variety of enzymatic reactions. Given humans' exposure to an enormous variety of substances of varying molecular weight and solubility, it should be expected that numerous enzymes of differing and overlapping functional capacity be required to accomplish drug elimination. Depending on the types of mediated metabolic pathways, these reactions have traditionally been classified as phase I or phase II.

Phase I reactions include oxidative, reductive, and hydrolytic processes. These metabolic reactions result in the attachment of polar functional groups to sub-

strates, for example, hydroxy or carboxyl groups produce more water-soluble compounds or provide a site for the attachment of a conjugate via a phase II reaction.¹ Of the phase I reactions, oxidative metabolism mediated by the cytochrome (CYP) P450 system, has stimulated the most recent research. Several reviews have summarized the drug interactions mediated by the CYP P450 system and the implications for clinical psychopharmacology.^{2,3}

Phase II reactions are conjugations. They include glucuronide (GLUC), sulfate, glycine and glutathione conjugation, acetylation, and methylation. Table 1 summarizes the characteristics of phase II drug metabolic reactions. Phase II processes, mediated by enzymes known as transferases, link a convenient moiety from a coenzyme to endogenous or exogenous substrates. Many drug metabolites undergo phase II conjugation after phase I reactions. For some drugs, conjugation can proceed without phase I functionalization of a drug molecule.^{4,5} Valproic acid (VPA), haloperidol, and olanzapine represent examples of psychotropic drugs that owe a significant portion of their metabolism to the direct glucuronidation of the parent compound.^{6–8} The formation of GLUC conjugates represents the most important of the various phase II reactions because of the numerous functional groups that are acceptors for GLUCs, the quantity of drugs that form GLUCs, and the ubiquitous nature of these conjugates in most vertebrate species.

The importance of phase II metabolic reactions, as pathways for elimination of drugs, has received little emphasis in the clinical psychopharmacology literature. Relative to the CYP P450 enzymes, less is known about the specificity of individual uridine diphosphate (UDP)-glucuronosyltransferases (UDPGT) toward the glucuronidation of specific medications. However, several extensively used anxiolytics, antidepressants, mood stabilizers, and antipsychotics are predominantly eliminated in the urine as GLUC conjugates. It is likely that the functional capacity of glucuronidation is a contributing factor to inter- and intrasubject variability in response to these psychoactive drugs. The *in vitro* interindividual variation in glucuronidation catalytic activity from different liver

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TABLE 1. Characterization of phase II metabolic reactions^a

Phase II Enzyme	Co-Factor	Location	Genetic Aspects	Common Reactions	Common Substrates	Inter-Hepatic Fold Activity Variation
UGT	UDPGA	Liver, kidney, intestine, lung, skin, prostate, brain	Mutations in exons UGT1A1, 2, 3, 4, and 5 associated with Crigler-Najjar type I and II and Gilbert syndrome Variant alleles exist for UGT2B7, and 2B15, no apparent effect on <i>in vitro</i> enzyme activity	Glucuronidation	Acetaminophen Bilirubin Ethinylestradiol Morphine Lorazepam Oxazepam Valproic acid Lamotrigine Olanzapine	3.1–10
SULT	PAPS	Liver, kidney, intestine	Variant alleles exist for SULT1A1 and 2A1, SULT1A1 genotype correlated with enzyme activity	Sulfate conjugation	Acetaminophen Albuterol Terbutaline Methyldopa	2–13
GST	Glutathione	Liver, kidney	Null genotypes exist GSTM1 and GSTT1, may be markers for cancer susceptibility	Glutathione conjugation	Azathioprine 6-Mercaptopurine Nitroglycerin Organophosphates	2
NAT	Acetyl coenzyme A	Liver, lung, spleen, gastric mucosa, RBCs, lymphocytes	Variant alleles exist for NAT1 and NAT2, slow acetylator status associated with increased isoniazid toxicity	N-acetylation	Dapsone Hydralazine Isoniazid Phenelzine Procainamide	3–18
MT	SAM	Liver, kidney, lung, CNS	Genetic polymorphism demonstrated for Catechol <i>O</i> -MT, Thiopurine MT, and Thio MT, low enzyme activity associated with thiopurine toxicity	Methylation	Dobutamine Dopamine Levodopa Norepinephrine Serotonin	2

^aUGT, glucuronosyltransferases; UDPGA, UDP-glucuronic acid; SULT, Sulfotransferases; PAPS, 3'-phosphoadenosine-5'-phosphosulfate; GST, Glutathione-S-transferases; NAT, N-acetyltransferases; RBC, red blood cell; MT, Methyltransferases; SAM, S-adenosyl-L-methionin; CNS, central nervous system.

banks was reported to range between 3.1- and 10-fold.⁹ Table 1 lists the reported interindividual fold variation of phase II catalytic processes. The purpose of this review was to examine the available data relating to glucuronidation, emphasizing potential drug-drug interactions involving psychoactive drugs, and summarize the importance in clinical psychopharmacology. A description of glucuronidation characteristics is followed by an explanation of the classification system. Finally, factors that can affect glucuronidation are discussed.

Glucuronidation Characteristics

Glucuronidation is a major detoxification pathway in mammals and represents the most common metabolic pathway in the formation of hydrophilic drug metabolites for excretion by the kidney or through biliary processes, or both.¹⁰ A general mechanism is shown in Figure 1 illustrating the metabolism of lamotrigine.¹¹

Uridine diphosphate glucuronic acid (UDPGA) mediates the conjugation of lamotrigine with glucuronic acid and acts as a cofactor. There is an attack at the activated UDPGA cofactor, resulting in the transfer of the glucuronyl moiety onto the substrate (aglycone), lamotrigine, with a subsequent release of UDP. The enzyme responsible for the formation of lamotrigine GLUC is UDPGT. UDPGTs are species-specific and vary widely in their substrate specificity. There is a large variability of functional groups that can be conjugated with glucuronic acid, including oxygen, nitrogen, sulfur, and carbon. Hydroxy and carboxy functional groups of primarily phenolic and alcoholic moieties form *O*-GLUCs. Many opioids and nonsteroidal antiinflammatory agents (NSAIDs) are primarily excreted as *O*-GLUCs. In general, planar phenolic molecules are glucuronidated more rapidly than drugs with bulkier structures.¹² Aromatic and aliphatic amines, amides, and sulfonamides can form *N*-GLUCs.^{1, 13}

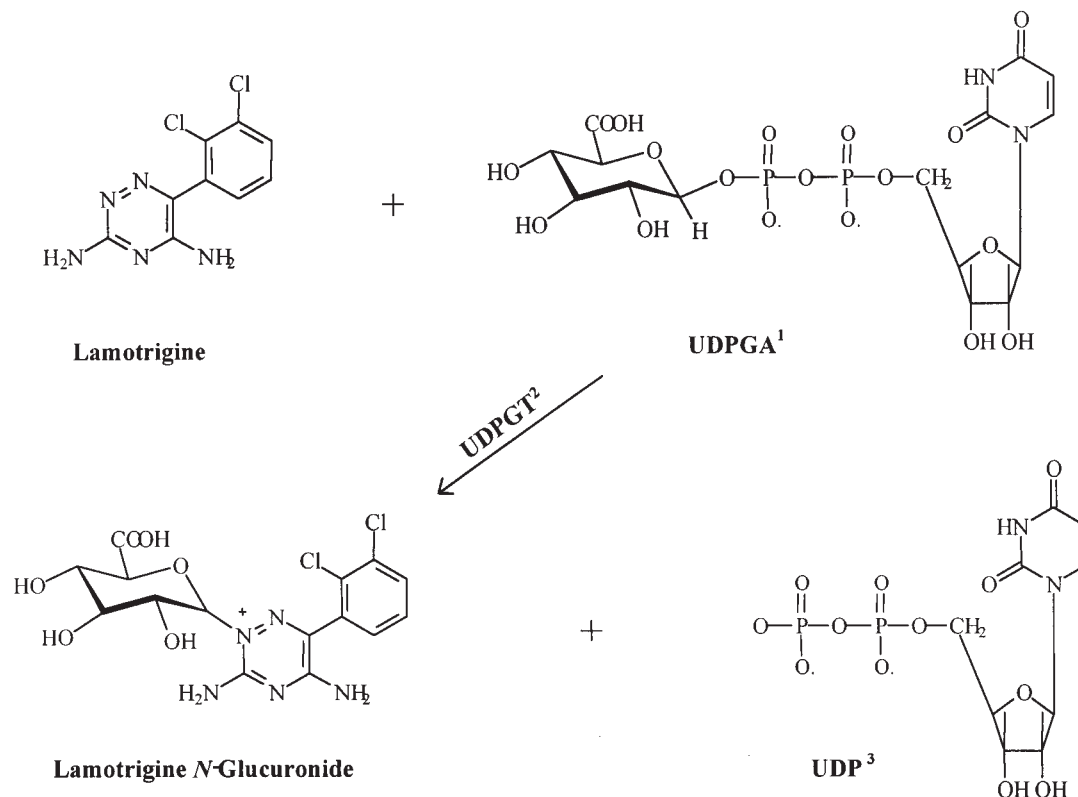


FIG. 1. Glucuronidation pathway of lamotrigine.

Tertiary amines including tricyclic antidepressants (TCAs), the atypical antipsychotics clozapine and olanzapine, the anticonvulsant lamotrigine, and the 5-hydroxytryptamine-2 antagonist cyproheptadine, form quaternary ammonium compounds via glucuronidation.¹⁴⁻¹⁸ *S*-GLUC and *C*-GLUC products are generally recognized as less common glucuronidation pathways in drug metabolism.¹

Although many pharmacologically active metabolites are known to be formed through phase I reactions, glucuronidated drugs have traditionally been regarded as inactive because of their hydrophilicity, low protein binding, and reduced volumes of distribution. Thus, GLUC metabolites have been often ignored in pharmacokinetic and pharmacodynamic studies. GLUCs can exert pharmacologic actions, and in some cases exhibit greater potency than the parent compound.⁴ For example, morphine-6-glucuronide (M6G) is a potent receptor agonist,¹⁹ digitoxin- and digoxin-GLUCs possess positive inotropic effects,²⁰ and fluorouracil GLUC was found to have antitumor activity.²¹ The glucuronidation of morphine (MOR) is of particular interest in clinical psychopharmacology.

MOR metabolism results in the formation of two major conjugates, M6G and morphine-3-glucuronide (M3G).²² Approximately 66% to 70% of a MOR dose is ex-

creted as these compounds.²³ M6G, a more potent analgesic than its parent drug, is 20-fold more active than MOR when injected into the periaqueductal gray matter of rats using the tailflick latency test.¹⁹ M3G has been shown to antagonize the antinociceptive effect of both MOR and M6G when administered via the intravenous, intrathecal, or intracerebroventricular routes in rats.²⁴ The development of tolerance after the administration of MOR may partially be explained by the formation of this GLUC.

β -Glucuronidases, esterases, and serum albumin catalyze the hydrolysis of GLUCs, converting them to the active aglycone, thus contributing indirectly to pharmacologic activity through the release of parent compounds (enterohepatic recycling).^{4, 25} β -Glucuronidase-resistant forms occur through the molecular rearrangement of primarily carboxylic acids containing GLUC conjugates. The intramolecular transesterification in hydroxyl groups of the glucuronic acid moiety is a phenomenon referred to as acyl migration.²⁵ Acyl GLUCs are reactive molecules that can bind irreversibly to plasma proteins.²⁶ This covalent protein binding can lead to toxicity and hypersensitivity reactions.²⁵ The acyl GLUC of zomepirac, an NSAID removed from the United States market because of a high incidence of anaphylaxis, undergoes irreversible protein binding *in vitro* and *in vivo*.²⁶

Analogous to phase I metabolic processes, stereospecific glucuronidation has been observed in a number of compounds. This may prove to be of therapeutic significance for compounds administered as racemic mixtures.^{10, 27-30} (+)MOR is preferentially glucuronidated over (-)MOR in human hepatic microsomes.³⁰ E-10-hydroxynortriptyline (E-10-OH-NT) undergoes stereoselective GLUC formation, with (+)E-10-OH-NT GLUC being preferentially formed in humans. Sixty-four percent of (+)E-10-OH-NT versus 35% of (-)E-10-OH-NT was recovered in the urine as GLUC conjugate after a 75-mg dose of 50:50 E-10-OH-NT racemate was administered to 10 healthy men. Comparably more (-)E-10-OH-NT was excreted unchanged.²⁸ *In vitro* analysis in human hepatic microsomal preparations demonstrated increased production of (*S*)-oxazepam over (*R*)-oxazepam GLUC. NSAIDS such as ketoprofen inhibited the formation of (*S*)-oxazepam over (*R*)-oxazepam significantly more than the (*R*)-isomer ($p < 0.01$).²⁹ Oxazepam pharmacodynamics could be altered in the presence of such inhibitors because the (*S*) enantiomer has been shown to be approximately 10 times more potent than the (*R*) enantiomer at inhibiting diazepam binding in rat synaptosomes.³¹

Glucuronidation is a low-affinity/high-capacity conjugative process^{4, 5} and seems to be under dual regulation, that is, (1) specific isozyme expression and (2) modulation of their functional state.³² The catalytic enzymes UDPGTs are located in the endoplasmic reticulum of cells of the liver, kidney, intestine, skin, lung, spleen, prostate, and brain.^{13, 33} Quantitatively, hepatic glucuronidation is most important.^{5, 13} Identification and classification of various UDPGTs has been accomplished in recent years.

UDPGT Classification

Thirty-three families of UDPGTs have been defined *in vitro* so far, and specific nomenclature similar to that used for the CYP P450 system has been established and recently refined.³⁴ The best examples are the mammalian UDPGTs and the three families of UGTs that have been identified in humans. As with the CYP P450 superfamily, the UGT superfamily is widely found among animals, plants, and bacteria, suggesting a common ancestral gene.^{34, 35} Among the UGTs, the UGT1 and UGT2 families seem to be the most important in human drug metabolism and are the focus of this review. Table 2 summarizes substrates of individual UGT isoforms. For a complete review detailing the background and nomenclature of the UGT system, refer to Mackenzie and colleagues.³⁴

UGT1 Family

At least 12 isoforms of UGT1 have been found to be encoded by this gene family^{3, 35, 36} and they share in

TABLE 2. *In vitro* UGT1 and UGT2 family substrates

IA1	IA3	IA4	IA6	IA9	2B7	2B15
Bilirubin	Amitriptyline	Amitriptyline	Acetaminophen	Acetaminophen	Androsterone	Androstane 3 α -diol
Buprenorphine	Buprenorphine	Chlorpromazine	5-Hydroxytryptophan	Dapsone	Buprenorphine	Dienestrol
Ethinylestradiol	Chlorpromazine	Clozapine	1-Naphthylamine	Ethinylestradiol	Chloramphenicol	Diethylstilbestrol
Irinotecan	Clozapine	Cyproheptadine	N-Hydroxynaphthylamine	Estrone	Codeine	6-Hydroxyquinoline
Nalorphine	Cyproheptadine	Desmethyloclozapine		4-Hydroxyestrone	Clofibric acid	8-Hydroxyquinoline
Naltrexone	Doxepin	Doxepin		Furosemide	Fenoprofen	5-(<i>p</i> -Hydroxyphenyl)-5-phenylhydantoin
	Fenoprofen	Imipramine		Ibuprofen	5-Hydroxytryptophan	5-(<i>m</i> -Hydroxyphenyl)-5-phenylhydantoin
	Hydroxymorphone	Lamotrigine		Ketoprofen	Hydromorphone	SCH23390
	2-Hydroxyestrone	Loxapine		Labetalol	Ibuprofen	
	2-Hydroxyestrrol	Olanzapine		Naproxen	Ketoprofen	
	Ibuprofen	Promethazine		Propranolol	Morphine	
	Imipramine	Retigabine		Propofol	Nalmefene	
	Ketoprofen			3,3',5'-Triiodo-L-Thyronine	Nalorphine	
	Loxapine			Retinoic acid	Naloxone	
	Morphine				Naltrexone	
	Nalorphine				S-Naproxen	
	Naloxone				Oxazepam	
	Naproxen				Oxymorphone	
	Promethazine				Propranolol	
	Valproic Acid				Valproic acid	
					Temazepam	

common exons 2 through 5.³⁷ The UGT1 family includes enzymes that can glucuronidate bilirubin, quinone, and phenol derivatives such as anthraquinones, flavones, and estrogens.^{37, 38}

At least five human UGT1A complementary deoxyribonucleic acids have been identified in hepatic tissue. They are UGT1A1, UGT1A3, UGT1A4, UGT1A6, and UGT1A9. UGT1A9 is also found in the kidney.^{35, 37, 39, 40} The human colon has been found to express all isoforms present in the liver, in addition to UGT1A7, UGT1A8, and UGT1A10.^{41, 42} UGT1A activities differ significantly depending on the location and substrate.

UGT1A1

UGT1A1, previously known as HUGBr1 or UGT1.1, is the enzyme catalyzing the glucuronidation of bilirubin.³⁷ This isoform is inducible by phenobarbital,⁴⁰ which has been used in the treatment of hyperbilirubinemia.⁴³ Human-expressed UGT1A1 was found to catalyze the glucuronidation of the opioids buprenorphine, nalorphine, and naltrexone. Of the compounds investigated, buprenorphine exhibited the most reactivity with UGT1A1, demonstrating a glucuronidation efficiency more than 1000-fold greater than that for naltrexone and nalorphine. Glucuronidation rates of buprenorphine were significantly reduced (75%) in liver microsomes from patients with type I Crigler-Najjar disease.⁴⁴ Type I Crigler-Najjar disease affects fewer than 1% of the population; however, it can lead to lethal hyperbilirubinemia as a result of defects in the UGT1A1 gene. Patients with Gilbert syndrome, a milder form of Crigler-Najjar, could potentially experience adverse effects from medications primarily metabolized by this isoform. This impairment does not generally seem to be of clinical significance¹⁰; however, two cases of severe toxicity characterized by neutropenia and diarrhea have been reported in patients with Gilbert syndrome after treatment with irinotecan, a chemotherapeutic agent.⁴⁵ UGT1A1 was found to catalyze the glucuronidation of irinotecan, which explains the observed toxicity in the two patients with Gilbert syndrome.⁴⁶

Drugs sharing this metabolic pathway, such as ethynyl-estradiol (a component of most oral contraceptives [OCs]) and naltrexone,^{38, 44} may competitively inhibit one another, resulting in pronounced or adverse pharmacologic effects. Likewise, drugs that inhibit UGT1A1 could result in hyperbilirubinemia or other adverse effects from coadministered drugs metabolized by this enzyme. Specific inhibitors of UGT1A1 that could be useful in human volunteer studies have not been identified.

Other xenobiotics glucuronidated by UGT1A1 include 2-hydroxyestrogenic catechols, anthraquinones, coumarins, flavonoids, and phenolic compounds.^{38, 47} Many of these agents are components of herbal products, foods,

and food preservatives and could also competitively inhibit the metabolism of UGT1A1 substrates such as bilirubin.³⁸

UGT1A3

Human-expressed UGT1A3 catalyzes the glucuronidation of primary, secondary, and tertiary amines, coumarins, flavonoids, anthraquinones, estrones, and phenolic and carboxylic acid-containing substrates.^{15, 39, 47} As with UGT1A1, 2-hydroxyestrone and 2-hydroxyestradiol are substrates of UGT1A3, whereas the 4-hydroxycatechol estrogens are not.⁴⁷ Apparent K_m values for primary amine substrates of UGT1A3 are approximately 100-fold higher than K_m values obtained for UGT1A4, making the glucuronidating ability of UGT1A3 one of low efficiency in comparison to UGT1A4. O-glucuronidated substrates of UGT1A3 include VPA; the opiates MOR, hydromorphone, nalorphine, buprenorphine, norbuprenorphine; the opioid antagonists naltrexone and naloxone; and various NSAIDs. Amitriptyline and cyproheptadine exhibited the greatest quaternary ammonium GLUC formation of the tertiary amines investigated by Green and associates.¹⁵

UGT1A4

Little structural difference exists between UGT1A4 and UGT1A3 with more than 90% similarity in their amino acid sequences.³⁴ Unlike UGT1A3, UGT1A4 exerts minimal activity toward phenolic compounds and is not involved in the metabolism of carboxylic acid-containing substrates such as NSAIDs.^{15, 48} Green and colleagues,^{14, 48} demonstrated that expressed human UGT1A4 protein efficiently catalyzes the N-glucuronidation of many primary, secondary, and tertiary amines. The highest glucuronidating efficiency has been demonstrated for the primary amines, 2- and 4-aminobiphenyl.⁴⁸

UGT1A4 was found to catalyze the tertiary amines including imipramine, amitriptyline, doxepin, promethazine, chlorpromazine, loxapine, and cyproheptadine to quaternary ammonium GLUCs.^{14, 48} Olanzapine has also been found to be a UGT1A4 substrate in vitro. (personal communication, T. Tephly, 6/15/2001). Two *N*-GLUC metabolites of clozapine have been observed after incubation with human UGT1A4 protein.⁴⁸ Formation of the first GLUC likely resulted from glucuronidation at the secondary amine position, whereas the quaternary ammonium GLUC represented the second metabolite. The first reaction is kinetically favored, with a reaction rate approximately six times greater than that for the quaternary ammonium GLUC.⁴⁸ The primary metabolite of clozapine desmethyl clozapine, a secondary amine, was found also to be glucuronidated by UGT1A4, whereas nortriptyline and desipramine were not.⁴⁸ Similarly, the 10-*N*-GLUC of olanzapine has been found to be preferentially formed

over the quaternary ammonium GLUC and represents the predominant circulating metabolite in human plasma.¹⁷ In contrast, another atypical antipsychotic, risperidone, is metabolized primarily by nonconjugative pathways, and its plasma concentration was not influenced by coadministration in human volunteers with probenecid, a broad-spectrum UGT inhibitor (Markowitz and associates, unpublished data).

The glucuronidation of lamotrigine is also catalyzed by this isoenzyme (Fig. 1). Because lamotrigine possesses both heterocyclic nitrogen and an amine side chain, glucuronidation could occur at both sites. Previously, the quaternary ammonium GLUC has been isolated and shown to be the major metabolite in patients receiving lamotrigine.¹¹

Additionally, UGT1A4 has been found to be the major isoform responsible for the N-glucuronidation of the structurally related anticonvulsant retigabine.⁴⁹ Significantly, human plasma concentrations of retigabine-N-GLUC exceeded those of the parent compound by a factor of 24, and the metabolite is likely to be subject to enterohepatic cycling.⁴⁹

UGT1A6

UGT1A6 is involved in the glucuronidation of primarily planar phenols.⁵⁰ 2-Naphthylamine and 4-aminobiphenyl are two carcinogenic arylamines found in small quantities in cigarette smoke. Glucuronidation of these compounds may contribute to a reduction in toxicity or determine the target of adverse effects by hydrolysis to the parent compound or hydroxy metabolite in such environments as the urinary bladder.^{51,52} Acetaminophen (APAP) glucuronidation rates were obtained in homogenates of both UGT1A6 and UGT1A9 transfected cells. APAP affinity was greater for UGT1A6 than UGT1A9 with K_m values of 2 mM and 50 mM, respectively.⁵⁰ Expression of UGT1A6 in the human central nervous system (CNS) has been demonstrated, specifically in the cerebellum.⁵³ King and associates⁵³ investigated whether the neurotransmitter 5-hydroxytryptamine was a potential substrate for this enzyme in HK293 cells. 5-Hydroxytryptamine was found to be efficiently glucuronidated by UGT1A6 with K_m and maximum velocity (V_{max}) values of 165 μ M and 55 $\text{nmol}\cdot\text{min}^{-1}\cdot\text{mg}^{-1}$, respectively. The significance of this observation remains to be explained.

UGT1A9

Although UGT1A6 favors reactions with planar phenols, UGT1A9 has been shown to be conjugated with both planar and nonplanar phenols, anthraquinones, flavones, aliphatic alcohols, aromatic carboxylic acids, and steroids.¹² Pharmacologic agents that were found to be substrates of UGT1A9 expressed in Chinese hamster V79 cells included propranolol, labetalol, propofol,

ethynylestradiol, 4-hydroxytamoxifen, ibuprofen, ketoprofen, naproxen, furosemide, and dapsone. Endogenous substrates, included 3,3',5'-triiodo-L-thyronine, estrone, 4-hydroxyestrone, and retinoic acid.¹²

UGT2 Family

The UGT2 gene products, unlike UGT1, likely arise from separate genes.^{54,55} The UGT2A subfamily represents glucuronosyltransferases present in rat and bovine olfactory systems that have not been well characterized.⁵⁶ The UGT2B family contains phenobarbital-inducible genes and others that are involved in the glucuronidation of endogenous steroids, biogenic amines, and various opioids.^{54,55,57-60} Although the majority of enzyme activity rests within the liver, the presence of UGT2B7 has been found in human intestinal mucosa.⁶¹

UGT2B7

There is evidence to support genetic variants of UGT2B7, as well as UGT1A1. Two forms have been identified, UGT2B7Y contains a tyrosine, and UGT2B7H contains a histidine residue at position 268.^{58,62}

MOR metabolism to both the M3G and M6G has been shown to be catalyzed by human embryonic kidney cells expressing UGT2B7.^{57,58} Other opioids that are also glucuronidated by this enzyme include dihydrocodeine, hydromorphone, oxymorphone, nalorphine, nalmefene, buprenorphine, and the antagonists naloxone and naltrexone.^{57,58} Codeine was found to be glucuronidated to the codeine-6-glucuronide (C6G), but at only 1% the rate of MOR.⁵⁷ The ratio of V_{max} for the formation of M3G versus M6G was found to be comparable to those reported in human hepatic microsomes. Because of this, it has been suggested that UGT2B7 is likely the major UGT that catalyzes MOR glucuronidation.⁵⁷ Like UGT1A6, UGT2B7 has also been isolated from the human brain and found to glucuronidate 5-hydroxytryptamine, although with much less efficiency.⁵³ Human brain microsomes were shown to catalyze the glucuronidation of naloxone and MOR.⁶³ This observed metabolism may contribute to the exertion and termination of opiate pharmacologic action within the CNS.

The UGT2B7 variant demonstrated specificity for a number of carboxylic acid substrates including various NSAIDs, VPA, and clofibrilic acid.⁶² Other substrates not containing a carboxylic acid included propranolol, testosterone, temazepam, chloramphenicol, and oxazepam, although activities for these substrates was substantially reduced.^{58,62}

UGT2B15

UGT2B15 expressed in HK293 cells was found to catalyze the glucuronidation of planar phenols, coumarins, anthraquinones, and flavonoids, as well as various andro-

genic and estrogenic steroids.⁶⁴ Other xenobiotics found to be substrates of this enzyme included a metabolite of diethylstilbestrol, dienestrol, the *para*- and *meta*-hydroxylated metabolites of phenytoin, 6- and 8-hydroxyquinoline, phenolphthalein, and the dopamine D₁ receptor antagonist SCH23390.⁶⁴ A variant of UGT2B15(D⁸⁵), UGT2B15(Y⁸⁵) has been characterized, and substrate specificity between the two forms was found to be similar.⁶⁵ The expression of the UGT2B15 gene was found in several extrahepatic tissues including kidney, testes, mammary glands, placenta, adipose, skin, uterus, prostate, and lung, suggesting it is a significant glucuronidating enzyme in humans.⁶⁵

Genetic Polymorphism

Genetic polymorphism has been demonstrated for UGT1A1, UGT1A4, UGT2B15, and UGT2B7.^{40, 65} Although little is known about the effects of mutations in the later three UGTs, UGT1A1 defects lead to hyperbilirubinemia conditions such as Crigler-Najjar disease and Gilbert syndrome.^{66, 67} Studies in Japanese, Eastern Scottish, and Canadian Inuit populations have revealed differences in the percentage incidence of homozygous (TA_n)-TAA alleles associated with Gilbert syndrome of 3%, 12%, and 17%, respectively.^{66, 68, 69}

Evidence supports polymorphic distribution of N-glucuronidation of nicotine and its principal metabolite, cotinine, in African American population.⁷⁰ The percentage of radiolabeled nicotine and cotinine excreted as the N-glucuronide was found to be 25% and 21%, respectively, in 51 African American subjects compared with 40% and 35%, respectively, in 57 Caucasian subjects. In addition, a trimodal distribution for nicotine and a bimodal distribution for cotinine N-glucuronidation was demonstrated in the African American group.⁷⁰ In a previous report, African Americans glucuronidated less of the carcinogenic nicotine derivative, nitrosamine 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanol (NNAL).⁷¹ A decrease in glucuronidating ability in African Americans is a potential cause of the increased incidence of lung cancer observed in this population. The specific UGT responsible for this observed genetic polymorphism has not yet been identified.

Yue and associates⁷² compared codeine metabolism in 149 Swedish Caucasian patients and 133 Chinese. The mean proportion of C6G excreted by the Caucasian subjects was 62%. This was significantly higher than the 44% excreted by the Chinese subjects. Comparably, the Chinese group excreted more unchanged codeine, norcodeine, and less norcodeine-GLUC. These results parallel those found with the smaller codeine doses required to treat pain in the Chinese population. APAP clearance (CL) has also been demonstrated to be 20% lower in patients of Asian descent than those of European descent.⁷³

The incidence of neonatal hyperbilirubinemia in the first few days of life was reported to be significantly higher in a group of Navajo American Indians than in a control group made up of African Americans, Hispanic Americans, and Caucasians. The breast milk of Navajo women was shown *in vitro* to significantly inhibit the glucuronidation of bilirubin; however, this alone could not account for the observed incidence of hyperbilirubinemia because several affected infants were bottle fed.⁷⁴

Factors Affecting Glucuronidation

As noted above, ethnicity can impact glucuronidating ability. Other factors that have been shown to alter glucuronidation include age, gender, body habitus, some disease states, and cigarette smoking. Table 3 summarizes these factors and their effects.⁷⁵⁻⁹⁹

Inducers and Inhibitors of Glucuronidation

There are fewer data on drug-drug interactions involving glucuronidation in contrast to an extensive documentation for CYP P450-mediated drug interactions. One reason for this situation is the lack of specific marker substrates or inhibitors of individual UDPGT isoforms analogous to debrisoquine and quinidine, a substrate and inhibitor, respectively, of CYP2D6 used in urine CL studies. In addition, many drugs are metabolized through multiple pathways, and substantial overlap of substrate specificity of UGTs exists.

The following section discusses the glucuronidation of drug substrates by specific isoforms mainly in expressed human cells, as well as interactions reported *in vivo*. Potential drug-drug interactions associated with the UGT enzyme system are summarized in Table 4.

Any substrate of UDPGT has the potential to competitively inhibit glucuronidation of other substrates metabolized by the same enzyme. Unlike the CYP P450 system, no specific inhibitors of individual UGT isoforms have been identified that may be safely administered clinically for use in inhibition studies.^{13, 100} Furthermore, no substrates that are glucuronidated exclusively are available for human studies. Although APAP is frequently used as a marker for glucuronidation activity, it is also subject to other metabolic pathways including sulfate conjugation and, to a minor degree, oxidation through the CYP P450 system.¹⁰¹ Nonetheless, several examples of competitive inhibitors that have relevance to clinical psychopharmacology exist.

Antidepressants

The 10-hydroxy metabolites of nortriptyline are often found in higher concentration than the parent drug in humans.¹⁰² The elimination of the E-10-hydroxy metabolite is primarily through glucuronidation.¹⁰² *In vitro* inhibition

TABLE 3. Factors affecting glucuronidation^a

Factor	Effect on Glucuronidation	Drugs or Compounds Affected
Age		
Infant	↑	Chloramphenicol, MOR, APAP, bilirubin, steroids. ⁷⁵⁻⁷⁸
Elderly	↔ or ↑	No change found for APAP, oxazepam, temazepam, or propranolol; decreased CL found for C6G and decreased unbound CL for oxazepam in the very elderly. ⁷⁹⁻⁸³
Gender		
Men	↑	CL higher in men for APAP, oxazepam, temazepam, and propranolol. Possible additive role with CYP1A2 resulting in higher clozapine and olanzapine concentrations in women. ^{80,81,84-90}
Women	↓	
Body habitus		
Overweight	↑	CL of lorazepam, oxazepam, temazepam, and APAP, likely the result of an increase in liver size and quantity of enzyme. ⁹¹
Underweight, malnourished	↓	Chloramphenicol, APAP. ⁹²
Disease states		
Fulminant hepatitis, cirrhosis	↓	Zidovudine, oxazepam, lamotrigine (glucuronidation by affected liver microsomes).
Hypothyroidism	↓	Oxazepam, APAP. ⁹⁴
HIV	↓	APAP. ⁹⁵
Cigarette smoking	↑	Propranolol, oxazepam, lorazepam, APAP. ^{80,86,96,97} Possible additive role with CYP1A2 induction resulting in decreased clozapine and olanzapine concentration. ^{88,89,98}
Ethanol	↓	<i>In vitro</i> inhibition of harmol, 2-naphthol, 4-methylumbelliform, phenothalein ⁹⁹

^aMOR, morphine; APAP, acetaminophen; CL, clearance; C6G, codeine-6-glucuronide; HIV, human immunodeficiency virus.

studies were performed on the formation of (+)-E-10-hydroxynortriptyline GLUC in human liver microsomes. Both amitriptyline and 2-hydroxydesipramine inhibited the glucuronidation of (+)-E-hydroxynortriptyline.¹⁰³

In human hepatic microsomes, nortriptyline noncompetitively inhibited the formation of M3G and M6G, whereas amitriptyline and clomipramine exerted competitive or mixed inhibition of MOR glucuronidation.¹⁰⁴ The K_i values for nortriptyline, amitriptyline, and clomipramine for the formation of M3G were 0.033, 0.160, and 0.090 mM, respectively. Amitriptyline has previously been shown to inhibit the glucuronidation of both codeine and MOR in human liver microsomes.¹⁰⁵ In 24 patients with cancer, the mean MOR area under the curve (AUC) was shown to increase approximately 2-fold after the addition of either clomipramine or amitriptyline.^{106, 107} Inhibition of glucuronidating enzymes by TCAs may contribute to the potentiation of opioid analgesic, as well as adverse effects.¹⁰⁷

In human liver microsomes, both amitriptyline and imipramine were found to inhibit testosterone UDPGT activity with percentage inhibitions of 75% and 68%, respectively.¹⁰⁸ Androsterone UDPGT activity was inhibited by 70% and 45% by amitriptyline and imipramine, respectively. Other endogenous substrates including estriol and 1-naphthol were also inhibited, although to a lesser degree. The inhibition of amitriptyline seemed to be non-competitive, whereas that of imipramine was competitive.¹⁰⁸ Alterations in endogenous steroid metabolism may play a role in the occurrence of adverse effects related to sexual function often associated with these agents.

Two cases of drug-drug interactions involving sertra-

line and lamotrigine have been reported suggesting sertraline inhibition of lamotrigine glucuronidation.¹⁰⁹ Lamotrigine concentration doubled after the addition of sertraline in the first case. In the second case lamotrigine concentration decreased from 19.3 $\mu\text{g/mL}$ to 9.8 $\mu\text{g/mL}$ when the sertraline dose was reduced to 50 mg/day despite a simultaneous increase in the lamotrigine dose.¹⁰⁹ Although sertraline possesses mild CYP2D6 inhibitory capability, lamotrigine is primarily metabolized by glucuronidation, making inhibition of this pathway a potential explanation.¹¹⁰ A single case report involving citalopram suggests the selective serotonin reuptake inhibitor (SSRI) may inhibit the glucuronidation of 8-hydroxydesmethylclomipramine (8-OHDCMI). After the addition of 40 mg of citalopram, a 5-fold increase in the 8-OHDCMI concentration was observed. Only slight increases occurred in the clomipramine, desmethylclomipramine, and 8-hydroxyclopramine concentrations. The increase in the 8-OHDCMI metabolite was observed despite a decrease in the patient's clomipramine dose from 150 mg/day to 75 mg/day. Because CYP P450 does not further metabolize 8-OHDCMI, citalopram may have inhibited the glucuronidation of this metabolite.¹¹¹

Anticonvulsant and antimanic agents

The majority of anticonvulsants undergo partial glucuronidation in the course of hepatic metabolism. Gabapentin and lithium, the only significantly renally eliminated agents in this class, seem devoid of the potential for drug-drug interactions associated with glucuronidation.¹¹² VPA is a widely used anticonvulsant

TABLE 4. Potential *in vitro* and *in vivo* substrates, inhibitors, and inducers of glucuronidation

Substrate	Inhibitor	Inducer
Morphine	Amitriptyline Clomipramine Clonazepam Diazepam Flumitrazepam Lorazepam Nitrazepam Oxazepam	Rifampin
Oxazepam	Ethinylestradiol Fenoprofen Ibuprofen Ketoprofen S-Naproxen	Phenobarbitone Phenytoin
(+)-E-10-Hydroxynortriptyline	Amitriptyline 2-Hydroxydesipramine	Pentobarbital
Bilirubin		Phenobarbital
Acetaminophen	Ethinylestradiol Probenecid Propranolol	
Androsterone	Promethazine Chlorpromazine	
Carbamazepine-10,11-transdiol	Valproic acid	
Codeine	Amitriptyline	
Lamotrigine	Sertraline Valproic acid	
Lorazepam	Ethinylestradiol Probenecid Valproic acid	
Temazepam	Probenecid	
Testosterone	Amitriptyline Chlorpromazine Imipramine Promethazine	
Zidovudine	Probenecid Valproic acid	

and antimanic agent implicated as a broad inhibitor of drug metabolism, including glucuronidation. VPA undergoes extensive glucuronidation with approximately 20% to 70% of a daily dose excreted in the urine as the GLUC.⁶ The percentage of the drug and GLUC excreted seems to increase with increasing dose, whereas oxidative metabolism decreases.¹¹³ The half-life ($t_{1/2}$) of lamotrigine was found to be 164% higher in subjects who were coadministered VPA (69.6 vs. 26.4 hours).¹¹⁴ The manufacturer's product information for lamotrigine recommends that the dose of lamotrigine be reduced to half of the normal dose in patients who are concurrently receiving VPA.¹¹⁵ The clinical importance of this interaction is the potentiation of the development of a severe rash or Stevens-Johnson syndrome when these agents are combined, because this dermatologic effect is proposed to be dose- or concentration-mediated.¹¹⁶

Combining anticonvulsant medications in the treatment of affective disorders has become more prevalent. The coadministration of VPA and carbamazepine caused an increase in plasma concentrations of carbamazepine-

10,11-epoxide. This was attributed to VPA inhibition of epoxide hydrolase; however, VPA also seems to inhibit the glucuronidation of carbamazepine-10,11-trans-diol.¹¹⁷ Because increased plasma concentration of the epoxide is associated with potential toxicity, careful monitoring of this combination is necessary.

The percentage of dose excreted as *p*-hydroxyphenobarbitone GLUC was similar between patients treated with and without VPA. However, the percentage of drug excreted as unconjugated metabolite significantly increased, (5.7% to 16.0%). In addition, the excretion of phenobarbitone *N*-glucoside was significantly decreased with concomitant VPA therapy (16.2% vs 1.9%).¹¹⁸ The potential mechanism for this observation may be depletion of both UDP-glucuronic acid and UDP-glucose by VPA.

The effects of VPA were studied on the disposition of amitriptyline in healthy volunteers.¹¹⁹ A 50-mg dose of amitriptyline was given with and without VPA 500 mg, twice daily. Subjects demonstrated a 31% mean increase in the AUC of amitriptyline and a 55% increase in its principal metabolite, nortriptyline.¹¹⁹ Both nortriptyline and

amitriptyline are partially eliminated by glucuronidation. The observed effects of VPA suggest inhibition of oxidative metabolism, conjugation, or both.

Recently, a case report documented increased clomipramine and desmethylclomipramine concentrations associated with adverse effects in a 46-year old woman, after the addition of VPA. The authors postulated the observed increase was caused by the inhibition of CYP2C or UGT by VPA.¹²⁰

In healthy subjects, VPA has been shown to decrease lorazepam CL by 40% and lead to increased lorazepam plasma concentration and decreased lorazepam GLUC formation.¹²¹ Samara and co-workers¹²² confirmed these findings in a randomized, double-blind, placebo-controlled crossover study. Under concomitant treatment with VPA 500 mg twice daily, subjects demonstrated increases in the lorazepam maximum concentration (8%), AUC (20%), and trough lorazepam plasma concentration (31%). Despite these increases, sedation scales revealed no significant effect under coadministration conditions. The disposition of VPA was not affected by lorazepam.¹²²

Significant sedation, confusion, and slurred speech was reported in a patient receiving clozapine, lithium, and VPA. Symptoms resolved after discontinuation of VPA but emerged again when the anticonvulsant was reintroduced.¹²³ A separate investigation reported a mean 41% decrease in total clozapine concentration after the initiation of VPA in four patients.¹²⁴ The combination of clozapine plasma protein-binding displacement and the inhibition of GLUC formation may explain the emergence of adverse effects reported above because of a potential increase in the unbound fraction of clozapine.¹²³

Like probenecid, VPA has been proposed to inhibit the glucuronidation of zidovudine. Because there is a higher frequency of adverse effects including bone marrow suppression associated with increased plasma zidovudine concentration, the significance of this interaction has been studied *in vitro* and correlated with *in vivo* data. VPA at a concentration level of 100 $\mu\text{g}/\text{mL}$, a concentration comparable to levels achieved *in vivo* for the treatment of acute mania, produced 50% inhibition of zidovudine glucuronidation in human hepatic microsomes.¹²⁵ Inhibition of zidovudine glucuronidation occurred in a concentration-dependent manner with complete blockade of glucuronidation by VPA at a concentration of 1,000 $\mu\text{g}/\text{mL}$.¹²⁵ At least one report demonstrated a significant increase in both plasma and cerebrospinal zidovudine concentrations, after the addition of VPA. Peak plasma zidovudine concentration levels increased to 344 ng/mL from a baseline of 119 ng/mL and cerebrospinal fluid concentration levels increased to 47 ng/mL from a baseline of 26 ng/mL. This change was accompanied by a zidovudine GLUC plasma concentration reduction of 50%.¹²⁶

The ratio of chloramphenicol GLUC to total nitro-

compounds excreted in the urine was shown to be significantly higher in two of six patients with schizophrenia receiving both a phenothiazine and phenobarbital.¹²⁷ The average phenobarbital dose for the group was 62 mg/day. The chloramphenicol $t_{1/2}$ in these two patients was 45 and 95 minutes compared to an average $t_{1/2}$ of 130.6 minutes in the control group.¹²⁷

The *in vitro* glucuronidation rate of (+)-E-10-hydroxynortriptyline in human hepatic microsomes was found to be higher in livers from two patients who had received prior treatment with pentobarbital.¹⁰³ GLUC formation was 31 pmol/mg per minute and 23.8 pmol/mg per minute, respectively, versus the mean activity of 18.8 pmol/mg per minute found in microsomes from all 13 patients studied.¹⁰³ Phenobarbital induces the glucuronidation of endogenous bilirubin catalyzed by UGT1A1⁴⁰ and is used therapeutically in the treatment of hyperbilirubinemia in newborns.⁴³

Phenytoin alone and in combination with phenobarbitone significantly increased oxazepam CL in nine patients with epilepsy.¹²⁸ Average phenytoin and phenobarbitone doses were 267 mg and 100 mg, respectively. Mean oxazepam AUC and $t_{1/2}$ were significantly reduced from matched controls (1864 vs. 1030 ng/mL and 6.99 vs. 3.31 hours, respectively) after a 15-mg dose. Bilirubin concentrations were also significantly lower in treated patients than in controls (6.1 vs. 10.9 $\mu\text{mol}/\text{L}$) reflecting phenobarbitone or phenytoin induction of UGT1A1.¹²⁸

Antipsychotics

Promethazine and chlorpromazine were both found to inhibit testosterone, androsterone, estriol, and 1-naphthol activity in human hepatic microsomes.¹⁰⁸ Chlorpromazine exhibited the most profound inhibition of testosterone, androsterone, and 1-naphthol with inhibition values of 77%, 86%, and 62%, respectively. Inhibition of testosterone and androsterone UDPGT activity by promethazine was 56% and 59%, respectively.¹⁰⁸ The mean circulating concentrations of haloperidol GLUC (35.4 ng/mL) have been shown to be higher than those of the parent drug (16.2 ng/mL), reduced haloperidol (14.1 ng/mL) or reduced haloperidol GLUC (12.7 ng/mL) in 39 hospitalized patients with schizophrenia.⁷ Increased extrapyramidal symptoms (EPS) were reported after the addition of olanzapine to existing haloperidol treatment in a 67-year old man.¹²⁹ The increase in EPS may be explained by a pharmacodynamic interaction at the D_2 receptor level; however, inhibition of haloperidol conjugative or oxidative metabolism could also be involved.¹²⁹

The plasma nortriptyline concentration was doubled to 185 ng/mL in a 38-year old man with schizoaffective disorder after the addition of clozapine.¹³⁰ The patient demonstrated symptoms congruent with an anticholinergic delirium including confusion, slurred speech, and

short-term memory loss. Both drugs were discontinued, and clozapine was gradually reinstated after 5 days. Ten days after the discontinuation of nortriptyline, the plasma concentration was 44 ng/mL. Concentration of the 10-hydroxy metabolite was not measured.¹³⁰ Competitive inhibition of oxidative enzymes might have accounted for the observed increase in plasma nortriptyline concentration; however, inhibition of glucuronidation is also possible. Although the affinity of the 10-hydroxynortriptyline metabolite for muscarinic receptors is approximately one ninth and one eighteenth that of nortriptyline's for the Z-10 and E-10-hydroxy metabolite, respectively,¹⁰² excessive concentrations of these metabolites, secondary to reduced GLUC formation, could have contributed to the adverse effects.

Antihypertensives

The β -adrenergic antagonist, propranolol is used in the treatment of cardiovascular disorders, performance anxiety, and in the symptomatic management of antipsychotic-induced akathisia.¹³¹ When administered at a dosage of 80 mg, twice daily, during a 4-day period, it was found to inhibit both the oxidation and glucuronidation of APAP in 10 healthy volunteers. Propranolol decreased the formation of the APAP GLUC by 27%, whereas sulfation was unaffected.¹³² Propranolol had no effect on the pharmacokinetics of lorazepam; however, CL was prolonged for diazepam, which undergoes metabolism through several oxidative pathways.¹³³

Guanfacine, a centrally acting α_2 agonist, is used in the treatment of disruptive and impulsive behavior in children.¹³⁴ It undergoes both oxidative and conjugative metabolism with 3-hydroxy guanfacine GLUC and sulfate conjugates accounting for approximately 50% of the metabolites found in the urine.¹³⁵ In a case report, VPA plasma concentration was found to decrease by 20% to 40% when guanfacine was decreased in two children and resulted in the reemergence of symptoms in one patient. This interaction may reflect competitive inhibition of glucuronidation.¹³⁶

Benzodiazepines

Diazepam inhibited codeine glucuronidation when incubated with UDP-glucuronic acid in human liver and kidney microsomes.¹⁰⁵ In addition, diazepam and oxazepam were found to inhibit the formation of both M3G and M6G. The degree of inhibition was concentration-dependent, with the most profound inhibition occurring with diazepam at a concentration of 1,000 μ M. The observed percentage of control activity in the presence of diazepam was 13% and 9% for the formation of M3G and M6G, respectively.¹⁰⁵ In separate studies, oxazepam has also been shown to competitively inhibit MOR glu-

curonidation in human fetal and adult liver microsomes.^{30, 137} Oxazepam was found to preferentially inhibit the formation of M3G of the (-)enantiomer, although it had the least inhibitory effect on the formation of (+)M6G.³⁰

An apparent pharmacodynamic interaction occurred after the addition of lorazepam to existing clozapine treatment in two patients. Marked sialorrhea and ataxia occurred with combined therapy. Previous administration of lorazepam with haloperidol in one of the patients had not resulted in such adverse effects. Although clozapine concentration was not measured, lorazepam may have competitively inhibited the glucuronidation pathway of clozapine.¹³⁸

NSAIDS

(*S*)-Oxazepam was found in one study to be preferentially glucuronidated over the (*R*)-isomer by a variant of UGT2B7,^{29, 139} although others have found low activity for UGT2B7 toward oxazepam.^{58, 62} Ketoprofen was found to competitively inhibit the glucuronidation of (*S*)-oxazepam to a greater degree than (*R*)-oxazepam.²⁹ Other NSAIDs that have demonstrated competitive inhibition included (*S*)-naproxen, ibuprofen, and fenoprofen.²⁹

Oral contraceptives

Ethinylestradiol is a substrate of UGT1A1³⁷ and has been found to participate in interactions of agents subject to glucuronidation. The effect of OCs on the pharmacokinetics of lorazepam, oxazepam, and chlordiazepoxide was investigated in eight healthy women who had been receiving OCs for a period of 6 months.¹⁴⁰ The CLs of lorazepam and oxazepam were significantly enhanced in the OC users compared with nonusers (288.9 vs. 77.5 mL/min for lorazepam ($p < 0.01$) and 251.2 vs. 97.86 mL/min for oxazepam) ($p < 0.01$). The CL of chlordiazepoxide, which undergoes oxidative metabolism, was found to be significantly impaired. It seems that OCs induced the glucuronidation of lorazepam and oxazepam, whereas the oxidative metabolism of chlordiazepoxide was inhibited.¹⁴⁰ In contrast, Abernethy and coworkers¹⁴¹ found OCs had no effect on the pharmacokinetics of either lorazepam or oxazepam in 17 women who had received OCs containing 50 μ g of ethinylestradiol or less for a minimum of 3 months. The larger sample of women in this study or the inclusion of subjects receiving less than 50 μ g of ethinylestradiol may reflect the absence of effect on benzodiazepine CL. Fifty μ g of ethinylestradiol taken alone or in combination with norethindrone for 21 days increased propranolol CL through glucuronidation significantly.¹⁴² Propranolol GLUC CL in women receiving ethinylestradiol alone

was 71% higher, whereas those receiving the combination of ethynylestradiol with norethindrone experienced a 32% increase in propranolol glucuronidation, suggesting that the progestin component exerts an inhibitory effect on glucuronidation.¹⁴²

OCs were also found to have an impact on APAP CL.¹⁴³ OC users had a higher CL rate of the analgesic (5.81 mL/min) compared with controls (4.12 mL/min). Likewise, OCs were found to impair the oxidative metabolism of antipyrine.¹⁴³ Given the widespread use of OCs concomitantly with psychotherapeutic agents, the potential significance of OC drug interactions deserves additional investigation.¹⁴⁴

Probenecid

Probenecid is a uricosuric agent used in the treatment of gout to competitively inhibit the reabsorption of uric acid in the renal proximal convoluted tubule. It also inhibits the active tubular secretion and urinary excretion of penicillin and cephalosporin antibiotics. Besides undergoing oxidative metabolism, approximately 20% of a probenecid dose is glucuronidated into the acyl-GLUC.¹⁴⁵

The glucuronidation of APAP and lorazepam was competitively inhibited *in vitro* by probenecid in hepatic microsomes.¹⁴⁶ In 11 patients who received concurrent probenecid and APAP, APAP CL was decreased from 329 to 178 mL/min ($p < 0.001$).¹⁴⁰ The 24-hour urinary excretion of APAP GLUC decreased from 260 to 84 mg, whereas the APAP sulfate excretion was increased proportionally so that the sum of the two conjugated metabolites remained constant. Similarly lorazepam CL decreased from 80.3 to 44.7 mL/min ($p < 0.001$) in nine subjects who received probenecid.¹⁴⁷

The pharmacokinetics of nitrazepam and temazepam were investigated after the addition of probenecid in 16 healthy subjects.¹⁴⁸ The total CL of antipyrine and nitrazepam were decreased by 22% and 25%, respectively. For norantipyrine and 4-OH-antipyrine, this decrease was caused by a reduction in GLUC formation. The sulfate fraction of norantipyrine and 4-OH-antipyrine was found to be increased. The fraction of temazepam excreted as GLUC was decreased by 33%; however, no changes in the pharmacokinetic parameters of temazepam were observed, as a result of increases in both the parent and sulfate fractions.¹⁴⁸

The antiretroviral zidovudine undergoes extensive glucuronidation. The exact isoforms responsible for the glucuronidation of zidovudine are unknown, but UGT1A6 and UGT2B7 have been excluded.¹⁰ Seven patients with AIDS-related complex who were receiving treatment with zidovudine were administered probenecid. The mean AUC of zidovudine doubled, along with a decrease in CL and an increase in $t_{1/2}$ after the coad-

ministration of probenecid.¹⁴⁹ Theoretically, psychoactive drugs extensively glucuronidated, including clozapine, olanzapine, and lamotrigine, could be similarly affected by probenecid.

Rifamycins

Rifampin is an antimicrobial known to induce drug metabolism through effects on human CYP isoforms, resulting in the increased CL of several drugs including OCs, warfarin, quinidine, digoxin, cyclosporine, protease inhibitors, nortriptyline, midazolam, and triazolam.¹⁵⁰ Interactions with zidovudine have also suggested an inductive effect on glucuronidation.¹⁵¹ The CL of nitrazepam was increased by 83% in healthy subjects who received rifampin at a dosage of 600 mg/day for 7 days. However, the pharmacokinetics of temazepam were not significantly affected by concurrent rifampin treatment.¹⁴⁸ The percentage of dose excreted as propranolol GLUC did not change after 22 days of treatment with rifampin 600 mg/day.¹⁵² This was noted in both extensive and poor metabolizers of debrisoquine; however, the fractional CL of propranolol glucuronidation increased after treatment because of a higher total propranolol CL.¹⁵² After 13 days of treatment with rifampin, analgesic effects of MOR were lost as measured by the cold pressor test in 10 subjects.¹¹³ Furthermore, MOR AUC values were significantly reduced by nearly 28% with proportional reductions in both M3G and M6G AUCs. The urinary recovery of the MOR GLUCs was also reduced.¹⁵³

Clozapine serum concentrations were significantly decreased (600%) after the addition of rifampin for the treatment of tuberculosis in a 33-year-old patient with schizophrenia. This reduction resulted in the reemergence of psychotic symptoms.¹⁵⁴ Clozapine undergoes both oxidative and conjugative metabolism; therefore, it is unlikely that the observed decrease in clozapine concentration can be solely accounted for by rifampin's induction of CYP P450 isozymes.

Conclusion

Relative to published clinical investigations of phase I metabolic processes and interactions, there has been less systematic study of changes in the pharmacokinetic parameters of psychotropic agents as a result of inhibition or induction of phase II processes. Substantial inter- and intra-individual variability in the pharmacokinetics of many psychotropic drugs has come to be a widely recognized, yet frequently unexplained phenomena associated with psychotropic drug therapy. The identification of genetic polymorphisms in CYP isoforms has explained the causes of significant variability in drug metabolism and response to many drugs, yet does not adequately

explain all observations. Glucuronidation as a major metabolic pathway is being increasingly recognized as a source of variability in the dose/concentration/effect relationship of several psychoactive drugs. The UGT enzyme system can be inhibited or induced and at least some UGT isoforms seem to be subject to genetic polymorphism. Additionally, stereospecific conjugation has been observed for a number of compounds. Taken together, these factors likely contribute to intersubject variability in psychoactive drug metabolism and, potentially, important drug-drug interactions.

The ability of some xenobiotics to simultaneously *coincide* or *coincide* both UGT and CYP isoforms (i.e., polycyclic aromatic hydrocarbons) seems to be a potential confound to accurate interpretation of drug-drug interactions. Even drug interaction case reports suggestive of cimetidine inhibition of clozapine metabolism¹⁵⁵ and rifampin induction of clozapine metabolism,¹⁵³ purportedly through CYP mechanisms, could also represent inhibitory¹³⁹ and inductive¹⁵¹ effects of these drugs on glucuronidation, respectively. Most SSRI antidepressants are likewise subject to extensive glucuronidation and may also affect the disposition of coadministered medications.¹⁰⁸ This is an intriguing possibility that has undergone little study. Future *in vitro* and *in vivo* drug interaction studies involving both old and new drug entities will undoubtedly investigate the effects and consequences of glucuronidation on psychotropic medications.

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