

Enterohepatic Circulation

Physiological, Pharmacokinetic and Clinical Implications

Michael S. Roberts,¹ Beatrice M. Magnusson,¹ Frank J. Burczynski² and Michael Weiss³

- 1 Department of Medicine, University of Queensland, Princess Alexandra Hospital, Brisbane, Queensland, Australia
 2 Faculty of Pharmacy and Department of Pharmacology, Faculty of Medicine, University of Manitoba, Winnipeg, Manitoba, Canada
 3 Section of Pharmacokinetics, Department of Pharmacology, Martin Luther University Halle-Wittenberg, Halle, Germany

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Abstract

Enterohepatic recycling occurs by biliary excretion and intestinal reabsorption of a solute, sometimes with hepatic conjugation and intestinal deconjugation. Cycling is often associated with multiple peaks and a longer apparent half-life in a plasma concentration-time profile. Factors affecting biliary excretion include drug characteristics (chemical structure, polarity and molecular size), transport across sinusoidal plasma membrane and canaliculae membranes, biotransformation and possible reabsorption from intrahepatic bile ductules. Intestinal reabsorption to complete the enterohepatic cycle may depend on hydrolysis of a drug conjugate by gut bacteria. Bioavailability is also affected by the extent of intestinal absorption, gut-wall P-glycoprotein efflux and gut-wall metabolism.

Recently, there has been a considerable increase in our understanding of the role of transporters, of gene expression of intestinal and hepatic enzymes, and of hepatic zonation. Drugs, disease and genetics may result in induced or inhibited activity of transporters and metabolising enzymes. Reduced expression of one transporter, for example hepatic canalicular multidrug resistance-associated protein (MRP) 2, is often associated with enhanced expression of others, for example the usually quiescent basolateral efflux MRP3, to limit hepatic toxicity. In addition, physiologically relevant pharmacokinetic models, which describe enterohepatic recirculation in terms of its determinants (such as sporadic gall bladder emptying), have been developed.

In general, enterohepatic recirculation may prolong the pharmacological effect of certain drugs and drug metabolites. Of particular importance is the potential amplifying effect of enterohepatic variability in defining differences in the bioavailability, apparent volume of distribution and clearance of a given compound. Genetic abnormalities, disease states, orally administered adsorbents and certain coadministered drugs all affect enterohepatic recycling.

In enterohepatic recycling, foreign chemicals entering the alimentary tract are absorbed into portal venous blood by enterocytes, removed from blood by uptake into hepatocytes, secreted into the bile, and then deposited back into the intestinal lumen where they may be reabsorbed by intestinal cells and available for recycling. Enterohepatic cir-

culcation may be significantly influenced by the formation of metabolites during absorption by the enterocytes and hepatocytes and during transit through the biliary system, intestinal tract or the systemic circulation (figure 1). For either the chemical or its metabolite, enterohepatic circulation can be terminated by elimination into the fae-

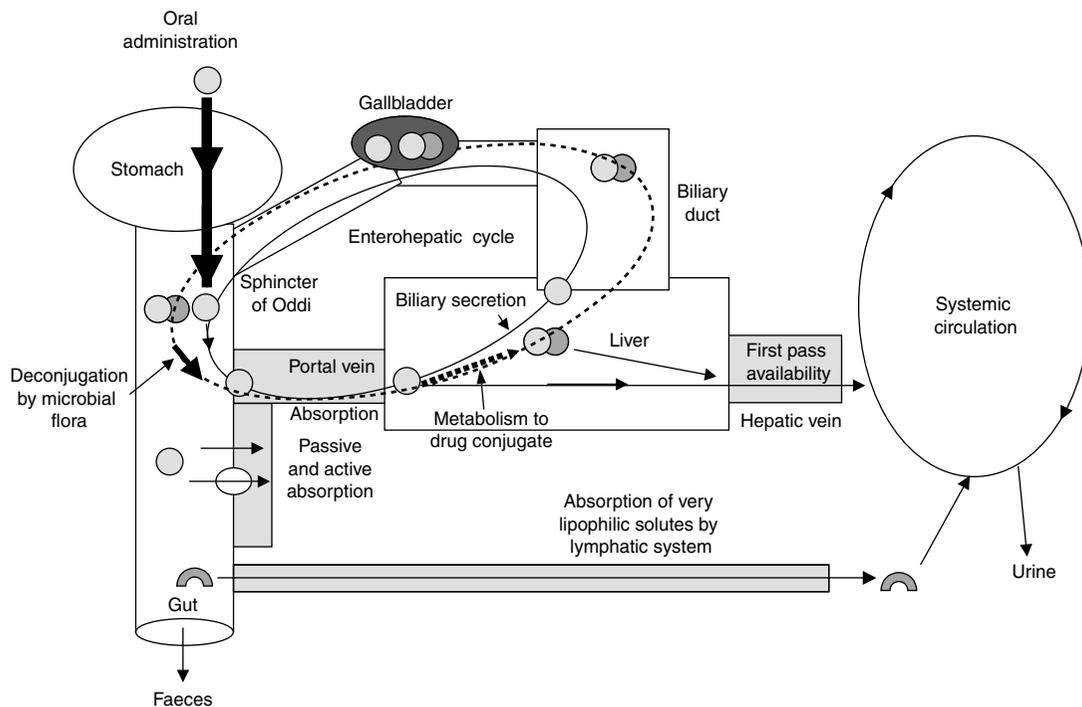


Fig. 1. Enterohepatic recirculation. Drugs entering the intestinal tract may be absorbed into the portal circulation where they can be removed from systemic circulation by hepatic uptake. The compound may then be excreted into the bile and pass back into the intestinal tract and become available for enterohepatic cycling. Biotransformation in enterocytes, hepatocytes, and the intestinal tract and throughout the body can convert the drug into metabolites, which may undergo enterohepatic cycling or escape into the urine and faeces. Some very lipophilic solutes may bypass the portal circulation and be absorbed into the systemic circulation via the lymphatic system.

ces or, if the compound has entered the systemic circulation, the urine.^[1] Enterohepatic circulation can also be interrupted by reversible storage in various reservoirs. Therefore, an unequivocal understanding of the enterohepatic circulation of chemicals requires an appreciation of the influence of biotransformation and membrane transport at each potential site.

The liver is the major site in the body for xenobiotic metabolism. An important function of the liver is the clearance of drugs by either biotransformation or biliary excretion. It therefore also follows that hepatocytes are at risk for exposure to the toxic bioactivated metabolites that result from the metabolism of some toxicants.^[2] The direct routing of blood to the liver from the gastro-

intestinal tract from which ingested xenobiotics are absorbed, as well as the tendency for such compounds to undergo enterohepatic cycling, also increase the vulnerability of liver cells to assault from toxicants.^[3]

Various endogenous as well as exogenous compounds are known to undergo enterohepatic circulation, which may serve a physiological function, for example in the recycling of bile acids. For drugs that are excreted extensively into the bile, insight into the magnitude of enterohepatic circulation is of crucial importance, as it will significantly affect pharmacokinetic parameters such as plasma half-life and area under the plasma concentration-time curve (AUC) as well as estimates of bioavailability. The complete study of enterohep-

atic circulation requires determination of the rate of biliary excretion and re-excretion, intestinal absorption, and faecal and urinary elimination of parent compound and all metabolites.^[4]

A major difficulty in reviewing this area is that much of the literature referring to enterohepatic recirculation has defined its contribution indirectly, often putting forward its contribution as a means to explain, for instance, the lack of excretion of certain compounds in urine. As an example, Jonsson et al.^[5] suggested that the small amounts of active metabolites of glibenclamide (glyburide) [4-*trans*-hydroxy-glibenclamide and 3-*cis*-hydroxy-glibenclamide] excreted in the urine of diabetic patients with impaired renal function may indicate the involvement of complementary nonrenal elimination routes. These routes may include the shunting of metabolised glibenclamide to the biliary excretion route and/or enterohepatic recycling of both metabolites and unmetabolised glibenclamide.

Other studies have defined enterohepatic recirculation indirectly from urine studies following interruption of the enterohepatic cycle. For instance, Hellstern et al.^[6] showed that although the fraction of lormetazepam and its glucuronide excreted in the urine during a 24-hour period was normally 9 to 35% of the dose, during interrupted enterohepatic recirculation this increased to 23 to 58% of the dose. However, the 24-hour post-dose bile fraction contained only 0.3 to 2.8% of the oral lormetazepam dose in the form of the drug and its glucuronide.

The use of nasobiliary catheters or biliary T-tubes to collect bile samples is an effective way to measure enterohepatic recirculation, but is often conducted using only small numbers of patients. Melnik et al.^[7] showed that the mean bile/plasma ratios of trovafloxacin in three patients for peak plasma concentration and AUC were 13.9 and 14.9, respectively, with concentrations of the *N*-acetyl metabolite in the bile about one-tenth that of the parent. More information is available on the excretion of endogenous compounds such as the bile acids (for example, Lenzen et al.^[8]). Given that

the principles of enterohepatic recycling for these compounds are also likely to apply to drugs, some aspects of the kinetics of their enterohepatic recycling are considered in this review.

In this overview, we examine enterohepatic cycling using predominantly human data, but also data from animal studies as appropriate, to assist in our understanding of the clinical pharmacokinetics of enterohepatic recycling. Consideration is also given to the range of relatively recently discovered transport processes affecting the recycling of drugs.

1. Anatomy and Physiology of the Liver and Bile Formation

The liver has an important role in many metabolic and excretion processes. Nutrient-containing blood from the gastrointestinal tract travels first to the liver via the portal vein, where nutrients such as carbohydrates, lipids and vitamins can be removed and stored until needed.^[9] The liver also synthesises and secretes bile, which provides a route for the excretion of endogenous and exogenous compounds such as bile acids, bilirubin, phospholipid, cholesterol, drugs and toxins.^[9] Bile is an aqueous solution, and hence is more suitable for the excretion of water-soluble compounds. Many compounds are metabolised to more water-soluble products prior to excretion in bile. In addition, the presence of micelle-forming bile acids above their critical micellar concentration allows solubilisation of lipid-soluble compounds in bile. Thus, the liver can excrete water-soluble as well as lipid-soluble compounds.^[10] Bile salts are later actively reabsorbed from the terminal ileum into the portal circulation to complete their enterohepatic circulation.

Hepatic bile acid secretion is a key determinant of the secretion of bile and biliary lipids. Bile acids are the major determinants of bile flow under physiological conditions (figure 2). The average bile flow in humans is 1.5 to 2.0 ml/min/kg body-weight.^[11] The ability of the liver to concentrate bile acids in bile is, therefore, an important aspect of bile formation. Bile also assists in fat digestion

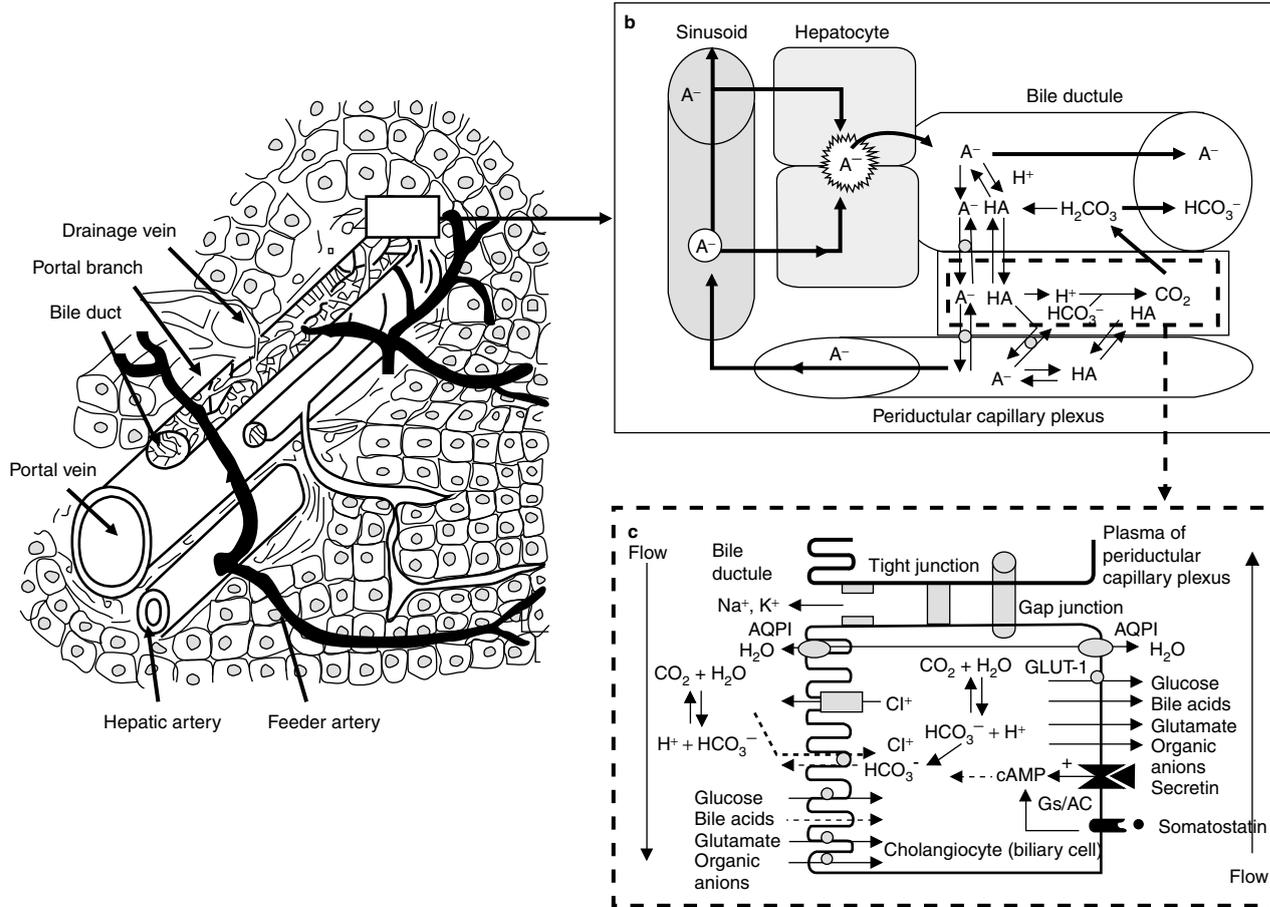


Fig 2. Schematic drawing of the bile duct and the peribiliary vascular plexus (reproduced from Roberts et al.,^[12] with permission). **(b)** Cholehepatic circulation; and **(c)** Transport mechanisms in the cholangiocyte. **AQPI** = aquaporin I; **Gs/AC** = GTP-binding protein/adenyl cyclase.

and absorption by providing bile acids and phospholipids to the duodenum, and it partially neutralises acidic chyme from the stomach and plays an immunological role by delivering IgA to the intestine. Because of its location and ability to efficiently extract a wide variety of compounds from the portal circulation, the liver plays a central role in removing toxic materials before their entry into the systemic circulation. Many of these agents are excreted into bile following detoxification and/or metabolism by the liver. Osmosis is considered to be the major mechanism of water movement during bile formation, although hydrostatic pressure may affect bile formation under experimental and pathological conditions. It is believed that the osmotic gradient is provided by organic and inorganic solutes secreted into bile. Bile formation is, thus, intimately related to hepatobiliary transport of biliary constituents.

The liver is composed of two types of epithelial cells: hepatocytes, which account for approximately 60% of the nuclear population, and cholangiocytes (epithelial cells that line intrahepatic bile ducts), which account for 3 to 5% of the liver cell population.^[12,13] Hepatocytes have been characterised as 'polarised' cells in that they absorb substrates from the blood and secrete metabolites in the bile. They are the only cells in the body that convert cholesterol to bile acids, and their canalicular membrane secretes bile acids vectorially into bile. In addition, hepatocytes excrete endogenous metabolites and endogenous xenobiotics and degrade hormones. This variety of hepatocellular functions is at variance with the apparent morphological homogeneity of liver cells. The constantly changing metabolic demands on the liver are such that hepatocytes must coordinately express an unusually diverse array of structural, enzymatic and secretory proteins.^[14] Cholangiocytes exhibit some morphological heterogeneity in that they tend to be cuboidal in small ducts and columnar in large ducts.^[15,16]

Bile is made as the result of active transport of its constituents into the biliary space and it is formed primarily by hepatocytes and secreted at

the bile canaliculus. The canalicular space is surrounded by two hepatocytes, and is separated from intercellular space by permselective tight junctions. The location of tight junctions also serves as a morphological and functional demarcation point between the apical (canalicular) and basolateral (sinusoidal and lateral plasma membrane) domains of hepatocytes. These two membrane domains differ in their lipid composition,^[17] and tight junctions, which are believed to provide effective barriers against lateral movement of lipids and proteins, maintain these differences.^[18] Tight junctions, along with desmosomes and gap junctions present at the lateral domain, are considered to represent important diffusional barriers between the interstitium and bile. In addition, microfilaments and microtubules in hepatocytes may be involved in canalicular bile formation.^[11,19,20] Circumstantial evidence indicates that microfilament dysfunction may alter bile formation (see section 4.8.2). Cytochalasin B, an agent that destroys microfilaments, causes cholestasis in the isolated rat liver preparation.^[21]

Bile formed at canaliculae is modified downstream in the bile ducts (ductular bile) by reabsorption and/or secretion of electrolytes and water. Although it has been suggested that water moves passively into bile,^[22] recent studies with microperfused intrahepatic bile ductules suggest that that ducts both secrete and absorb water in response to osmotic gradients.^[23] Bile entering the duodenum includes a canalicular secretion, elaborated by the hepatocytes, and a ductular/ductal secretion, elaborated in the biliary channels. Bile is concentrated by the gall bladder by reabsorption of electrolytes and water and secreted into the small intestine via the Sphincter of Oddie. The volume and composition of bile are modified within the lumen of intrahepatic bile ducts as a result of the absorption and secretion of water and solutes by cholangiocytes (figure 2). Cholangiocytes provide a large surface area for transport between blood and bile and play a significant role in bile formation.^[24]

The fate of unconjugated bile acids entering the canaliculus depends on their lipophilicity. For example, ursodeoxycholate is secreted in the unconjugated form, protonated from H_2CO_3 into ursodeoxycholic acid with generation of HCO_3^- (figure 2). The lipophilic acid is reabsorbed by the biliary epithelial cells into the periductule plexus and returned to the liver.^[25] The cholehepatic cycling causes a bicarbonate choleresis, because intraluminal carbonic acid donates the proton, permitting passive bile acid absorption.^[26] Other bile acids have similar bile secretory effects to ursodeoxycholate.^[27]

2. Component Processes in Enterohepatic Circulation

2.1 Absorption from the Gastrointestinal Tract Into the Portal Blood Supply

Enterohepatic circulation occurs after the process of absorption of a solute from the gastrointestinal tract into the portal blood (figure 3). The amount of solute involved in enterohepatic circulation will be dependent on the availability of solutes to the liver and in turn on absorption from the gastrointestinal tract. The factors affecting oral absorption include the formulation and characteristics of the drug product as well as patient characteristics such as pH of lumen, gastric emptying time, intestinal transit time, surface area, gastrointestinal disease and mesenteric blood flow. Other factors are the presence of food and drugs in the gastrointestinal tract and the pharmacokinetic characteristics of the drug in terms of metabolism by the gut wall and gut microflora.^[28] These factors will define the disintegration and dissolution properties of the drug formulation as well as the solubility of the drug molecule in the luminal environment of the gastrointestinal tract.

The absorption of drugs across the gastrointestinal wall membrane may be affected by different rate-determining steps, and may be limited by the unstirred water layer, by the characteristics of the membrane or by portal blood flow.^[29] The actual rate-determining step is defined by the physico-

chemical properties of the drug (pKa, water/lipid solubility, structural mimicry of endogenous substrates for transport proteins) and the physiology of the gastrointestinal tract. Some solutes such as di- and tri-peptides, amino acids, inorganic ions, glucose and galactose may be actively transported across the gut wall. The intestinal peptide transport system has broad substrate specificities for peptidomimetic drugs, including β -lactam antibiotics, ACE inhibitors, renin inhibitors, bestatin, thrombin inhibitors and thyrotropin-releasing hormone (protirelin) and its analogues.^[30]

Drug absorption into the portal circulation can occur down the length of the gastrointestinal tract to the superior rectal vein. Absorption from the lower region of the rectum (middle and inferior rectal veins) bypasses the portal circulation.^[31] Some highly lipophilic solutes, for example longer chain fatty acids (>12 carbon atoms), also bypass the portal blood supply by being absorbed via the intestinal lymph system and circulated via the thoracic duct back into the systemic circulation.^[32] Examples of compounds transported by this pathway include cyclosporin, naftifine, probucol, mepitiostane, lipophilic vitamins, halofantrine, DDT and lipophilic prodrugs.^[32]

2.2 Metabolism and Efflux by the Gut Wall

Transport across the wall is also affected by the susceptibility of the molecule to various plasma membrane transporters and intra- or extracellular biotransformation enzymes. The overall fraction of the dose reaching the liver (f_{liver}) will be a product of the net fractions of the dose absorbed across the apical membrane of the epithelial cell (f_{ab}) and escaping first-pass intestinal metabolism (f_{nm}), i.e. $f_{\text{liver}} = f_{\text{ab}} \times f_{\text{nm}}$.

Drug-metabolising enzymes found in the intestine usually have concentrations much lower than have been found in the liver.^[29] The variable and low oral bioavailability of cytochrome P450 3A (CYP3A) substrates such as cyclosporin and verapamil is caused to a considerable extent by the wide intra- and inter-individual variation in the activity of the intestinal enzymes.^[33] Both phase I

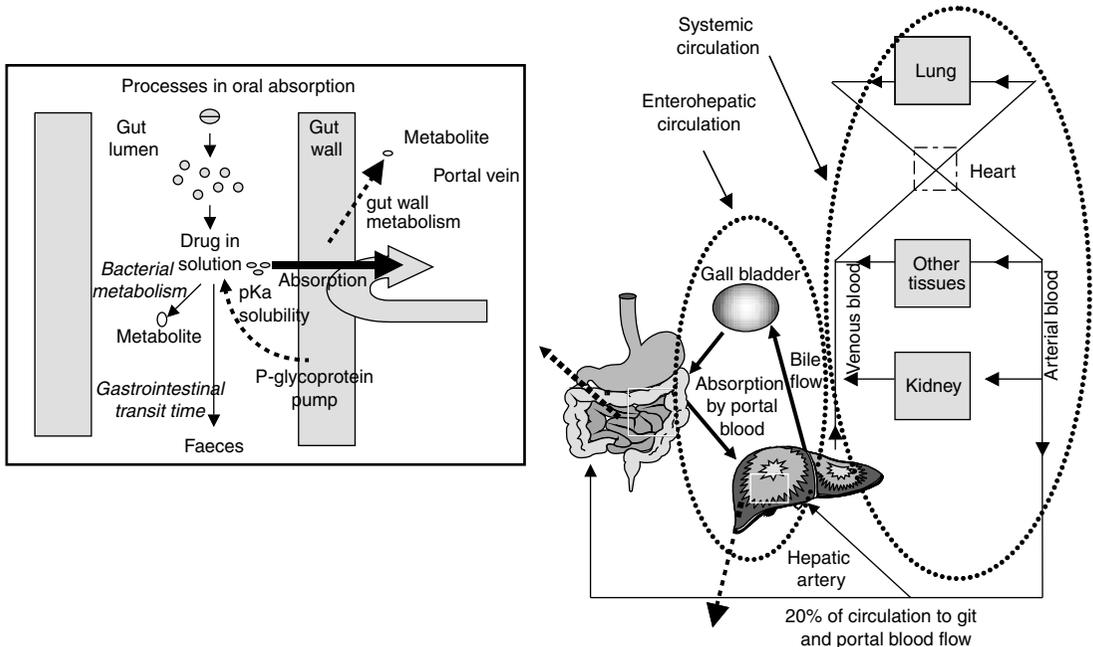


Fig. 3. Oral absorption into the enterohepatic and systemic circulation (see figure 4 for transport processes in the liver).

(oxidation and hydrolysis) and phase II (conjugation) metabolism exists in the gut wall, and the resulting metabolites may become involved in enterohepatic recirculation. The types of solutes metabolised in the gut wall have recently been reviewed by Ilett et al.^[34] The phase I metabolic reactions occurring in the gut wall are oxidation (e.g. ethanol), *C*-hydroxylation (e.g. ethinylestradiol), *N*-dealkylation (e.g. flurazepam), *O*-dealkylation, deamination, *S*-oxidation, desulfuration, decarboxylation and hydroxamate reduction. The corresponding phase II reactions are esterification (e.g. hydoxycholic acid), ether formation (e.g. ethinylestradiol), sulfation (e.g. ethinylestradiol), glutathione conjugation (isosorbide dinitrate), glycine conjugation, *N*-acetylation, *O*-acetylation (e.g. *N*-hydroxy-aminofluorene), and *O*-methylation.^[34]

P-glycoprotein within the brush border on the apical (luminal) surface of the intestinal epithelium reduces the absorption of a range of drugs (for example cyclosporin, tacrolimus, paclitaxel,

diltiazem, dexamethasone, lidocaine, erythromycin and HIV protease inhibitors) by pumping the xenobiotic from the enterocyte back into the intestinal lumen.^[35] One of the most abundant gut wall enzymes, CYP3A4, is located near the glycoprotein and has similar substrate specificity, leading to a more effective intestinal barrier to certain xenobiotics.^[35]

2.3 Transport from the Portal Blood Across Hepatocyte Plasma (Sinusoidal) Membranes

Delivery of drugs from the circulation to the hepatocyte membrane surface has been the subject of intense investigation for many decades. The role of extracellular binding proteins on the uptake of highly protein-bound drugs also has been a controversial issue. Recent studies into this area have provided evidence that uptake may occur from both the unbound and protein-bound fractions.^[35]

Once in the portal circulation, drugs are absorbed from the sinusoids by a transport process,

or they may diffuse across the hepatocyte plasma membranes (figure 4). For highly lipophilic drugs, traversing the membrane involves several steps. The first is reversible binding of the drug to the plasma side of the cell membrane leaflet. Traversing the lipid bilayer itself could occur by a flip-flop process of the hydrophilic region of the molecule. To aid in this process, the carboxyl group of fatty acids becomes protonated by hydrogen bonding, either with constituents in the membrane/water in-

terface or constituents within the membrane itself.^[35] The uptake process is completed by dissociation of the drug from the cytosolic side of the cell membrane leaflet. Any one of these barriers has the potential for being rate-limiting in the overall uptake process.

Transmembrane flux rate and the subsequent rate of metabolism are key determinants in the hepatic clearance of compounds. Accordingly, hepatic extraction is determined by hepatic blood flow,

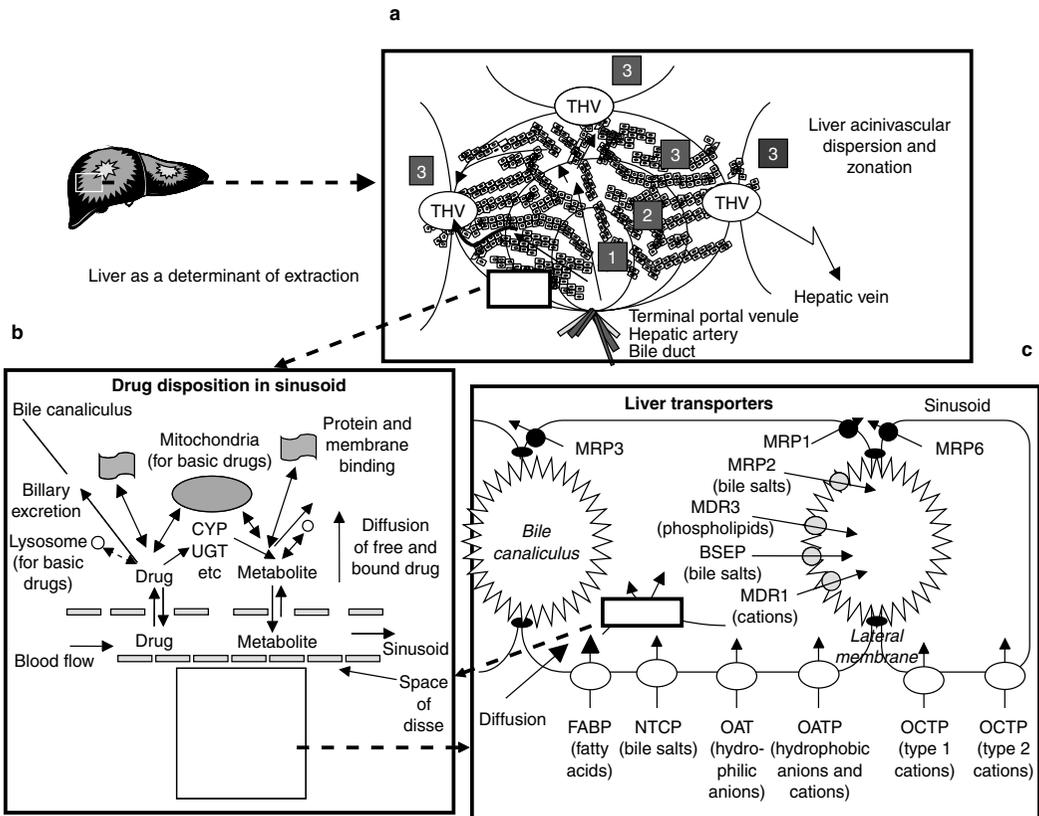


Fig. 4. Transport processes in the liver determining enterohepatic recirculation kinetics of drugs and their metabolites. **(a)** Uptake and clearance of drugs depends on the vascular dispersion in the acini (variation in times taken by blood cells from portal to hepatic venules; two pathlengths shown) and zonation of liver function (three zones shown). **(b)** Ion trapping, binding and transport in the hepatocyte. **(c)** Transporters into and from the hepatocyte into the bile (via biliary canaliculus) and back into sinusoid. **BSEP** = bile salt export pump; **CYP** = cytochrome P450; **FABP** = fatty acid binding protein; **MDR** = multidrug resistance; **MRP** = multidrug resistance-associated protein; **NTCP** = Na^+ -taurocholate cotransporting polypeptide; **OAT** = organic anion transporter; **OATP** = organic anion transporting polypeptide; **OCTP** = organic cation transporting polypeptide; **THV** = terminal hepatic venule; **UGT** = uridine 5'-diphosphate glucuronosyltransferase.

binding to blood constituents, hepatocyte permeability and biliary and/or metabolic elimination. Hepatic clearance is also a function of vascular dispersion caused by the heterogeneity in sinusoidal blood flows and interconnection between sinusoids (figure 4).^[36,37] Vascular dispersion is used in the scaling of *in vitro* liver microsomal or hepatocyte data to predict *in vivo* availabilities.^[38,39] Vascular dispersion is also being increasingly used in the modelling of hepatobiliary excretion.^[40]

Compounds undergoing enterohepatic recirculation are usually transported as parent solutes through different carrier-mediated systems or by diffusion as uncharged solutes, which are later biotransformed in the liver to suitable polar solutes. Specific plasma membrane transporters exist, including Na⁺-dependent and Na⁺-independent anionic transporters as well as cationic transporters (figure 4). Many of the liver plasma membrane organic anion transporters into hepatocytes have only recently been cloned and their transport properties identified.^[41,42] At this time, much more information is available on rat than human transporters, especially on drug uptake characteristics,^[43] and so both are mentioned here. Human transporters include Na⁺-taurocholate cotransporting polypeptide (NTCP), human organic anion transporting polypeptide (OATP1), human liver specific transporter (LST1)/OATP2, human organic anion transporter (hOAT1), human prostaglandin transporter (hPGT) and various organic cation transporters, including the organic cationic transporter associated with the uptake of hydrophilic aliphatic and certain aromatic cations (hOCT1). In addition, certain multidrug resistance-associated protein (MRP) transporters (MRP1, MRP3 and MRP6) exist at the plasma membrane but usually only show significant expression in certain diseased livers.

Proost et al.^[44] have shown that the carrier-mediated uptake of the cationic aminosteroidal neuromuscular blockers is related to plasma protein binding and to the partition coefficient between octanol and aqueous buffer (log P). The cellular transmembrane electrical potential difference (membrane potential) also has been suggested to

greatly affect cellular substrate availability. Membrane potential has been reported to influence the hepatic uptake of organic anions such as the fatty acids^[45-47] and the bile acid taurocholate,^[47,48] as well as other drugs into tissues other than liver.^[49] Canalicular secretion of taurocholate also was shown to be dependent upon membrane potential.^[47,48] Canalicular membrane hyperpolarisation resulted in the stimulation of taurocholate-dependent fluid secretion.

2.4 Transport from the Hepatocyte Membrane to Metabolising Sites and to the Bile Canaliculae

There is some conjecture on the processes by which solutes traverse the hepatocyte to reach drug-metabolising sites such as the endoplasmic reticulum and the bile canaliculae. In most pharmacokinetic studies, it has been traditionally assumed that most solutes move 'instantaneously' in the liver in the unbound form in accordance with the principles of a well-stirred tissue compartment in the radial direction to flow. More recent evidence suggests that transport in the liver may be limited by radial diffusion,^[50,51] binding to proteins or membranes in transport through the liver,^[52] or transport into pools.^[53] Zucker et al.^[54] have suggested that intrahepatocellular drug transport may occur through several processes. These include cytoplasmic diffusion, protein-mediated diffusion, cytoplasmic flow, vesicular transport and drug transfer from intracellular membranes to intracellular proteins.

Cytoplasmic flow has been postulated to result from the actions of intracellular motor proteins and filaments (myosin and actin) in directing flow (also known as cytocirculation). Although detailed studies have not been undertaken in hepatocytes, canalicular contractility may be expected to set up a cytocirculatory movement of drugs to the canalicular membrane. Although vesicular transport may play a minor role in the overall transport of drug molecules to the canalicular membrane, it is important in protein trafficking.

Cytoplasmic diffusion and protein-mediated diffusion of drugs have been suggested to play an important role in overall intracellular drug disposition. The transfer of bile acids from the plasma membrane to the canalicular membrane involves several different steps. These include lateral diffusion of the unbound ligand through the cytoplasm, lateral diffusion of the unbound ligand through membranes, vesicular transport and transport by cytosolic binding proteins. Cytosolic bile acid binding proteins include glutathione transferases, the Y' bile acid binding protein and, to a lesser extent, fatty acid binding proteins (FABP).^[55,56]

Vesicular transport of bile acids to the canalicular membrane was suggested over 30 years ago.^[57] Using confocal scanning fluorescence microscopy, El-Seaidy et al.^[58] reported a lack of evidence for microtubule-dependent trafficking of bile acids or the intracellular sequestration of bile acids prior to their canalicular secretion. The conclusions were based upon examining the effect of the microtubule inhibitor colchicine on the biliary secretion of the fluorescent bile salts cholylyl-lysyl-fluorescein and chenodeoxycholylyl-lysyl-fluorescein in hepatocyte couplets. The previously reported microtubule-dependent trafficking events were suggested to be important in bile salt-induced delivery of bile salt transporters to the membrane rather than have any direct role in the biliary secretion process of bile salts.

It is also possible that, although largely undefined at this time, transporters also facilitate the uptake of some solutes by intrahepatic organelles such as lysosomes and mitochondria. Lipophilic basic drugs appear to accumulate in the relatively acidic hepatic lysosomes as a result of the pH difference with the cytoplasm (pH-partition theory) but also possibly due to binding (partition or adsorption) to lipophilic substance(s) and/or aggregation within lysosomes.^[59] Imipramine and other weak lipophilic bases such as chlorpromazine and propranolol increase intralysosomal pH, and it has been shown that the accumulation of imipramine in lysosomes can be inhibited by coadministration of lipophilic basic drugs.^[59] However, it has been

suggested that the uptake of psychotropic drugs by liver tissue may depend more on phospholipid binding than on lysosomal trapping.^[60] We have recently shown that the sequestration of cationic drugs in the liver occurs by both slow and fast binding to protein and membrane sites as well as ion trapping into mitochondria (20% of hepatocyte, pH 6.67 in fasted state) and lysosomes (1% of hepatocyte and pH 4.70).^[61] The extent of ion trapping is determined by the pH of the organelles, pKa of the drug and fraction of the hepatocyte volume occupied by the organelles.

The overall hepatic extraction ratio for cationic drugs is determined by perfusate flow rate, permeability surface area product and intrinsic clearance and, in turn, by the lipophilicity of the solute.^[61] In liver disease, the extent of fibrosis and amount of metabolising protein present are other determinants affecting hepatic extraction ratio and other parameters.^[62] The accumulation of cationic lipophilic drugs in the liver, probably due to trapping in mitochondria and lysosomes as well as binding, results in a slow elution and a low availability of cationic lipophilic drugs for excretion in the bile and enterohepatic transport.

2.5 Biotransformation in the Liver

Biotransformation is the cellular process that produces pharmacologically or toxicologically active metabolites, as well as inactive ones, by altering the structure of a xenobiotic or by conjugating it with an endogenous compound, thereby changing the aqueous solubility and eventual distribution of the chemical.^[63] The ability to biotransform and eliminate potentially toxic chemicals from the body is a well-known protection mechanism^[64] and is an important determinant of the rate of elimination of lipophilic chemicals, which can accumulate in the body.^[65] Factors that influence biotransformation at each of several potential sites include species, age and development stage, gender, nutritional status and prior or concurrent exposure to various chemicals.

In general, blood-borne drugs and toxicants are most capable of crossing cell membranes and bind-

ing to protein targets if they are lipophilic, small and neutrally charged.^[66] Subsequent detoxification is most efficient when the parent compound can be altered to a metabolite that is hydrophilic, large and charged,^[67,68] so that it has the appropriate properties for biliary secretion (section 3). These changes typically occur in two stages. First, in what is often referred to as phase I reactions, the parent compound is typically hydrolysed or oxidised (or in some cases, reduced). This leads to formation of a metabolite that may be conjugated to a larger, much more hydrophilic, molecule, often containing a charged moiety. This second conjugation step is often referred to as a phase II reaction. Usually, this stepwise biotransformation is necessary due to the lack in the parent compound of a functional group for easy conjugation.

The phase I drug-metabolising enzymes are predominantly the CYP superfamily. Over 35 human CYPs identified have been classified into families, subfamilies and specific isoforms by amino acid sequence homology. Masimirembwa et al.^[69] have suggested that their relative abundances in humans are CYP1A2 (13%), 2A6 (4%), 2B6 (<1%), 2C (20%), 2D6 (2%), 2E1 (7%), and 3A4 (30%), but can vary because of induction (CYP1A1/1A2, 2A6, 2E1, 2C and 3A4), inhibition (all CYPs) and genetic polymorphism (CYP2A6, 2C9, 2C19 and 2D6). The clinical pharmacokinetics associated with CYP3A have recently been reviewed.^[70]

Probably the most common phase II (conjugation) reaction is glucuronidation, catalysed by the uridine 5'-diphosphate (UDP)-glucuronosyltransferases (UGTs), consisting of at least 10 different UGT isoforms.^[71] Most drug [morphine, paracetamol (acetaminophen)] and endogenous (bilirubin, estradiol) glucuronides are excreted in the bile and undergo enterohepatic recirculation. Many drug conjugates formed by phenol sulfotransferase and cytosolic glutathione *S*-transferase (GST) can also undergo enterohepatic recirculation.

From a pharmacokinetic viewpoint, drug metabolites (especially conjugates) are usually more polar than the parent drugs and will have difficulty backdiffusing across the sinusoidal barrier.^[53]

Some of these metabolites, therefore, have longer hepatic disposition half-lives and there exists the potential for them to accumulate in the hepatocyte to a greater extent than the parent drug if diffusion is the main means of transport.^[72] Larger polar metabolites are cleared in the bile (section 3) and others may be actively transported back into the sinusoid (figure 4). At this time, it is not possible to predict which metabolites will experience a diffusion barrier and accumulate or be actively transported out of the liver into the bile or into sinusoids.^[73]

The major metabolic pathways in the liver show acinar gradients.^[74] Hence, oxidative energy metabolism (for example fatty acid oxidation), amino acid catabolism, ureagenesis, gluconeogenesis, cholesterol metabolism and certain types of protective metabolism occur in the periportal region. Nonparenchymal cells such as Kupffer, stellate, endothelial and pit cells are generally also predominant in this region. In contrast, glycolysis, liponeogenesis, ketogenesis, glutamine formation and xenobiotic metabolism are preferentially located in the perivenous (also called centrilobular or zone 3) region. Accordingly, endoplasmic reticulum (microsomes) is more prevalent in the perivenous region. Zonation is determined by a range of gradients including oxygen tension, bile acids, hormones and cell expression.^[74]

Most studies on the zoned expression of CYP genes have been in rat liver.^[74] In general, the genes are mainly distributed in the perivenous region but there are some exceptions, such as CYP2C7 and FABP, which are mainly periportal (figure 5). Translation of such data to human livers is complicated by different CYP genes being present in rat and human livers (CYP2C7, CYP2C11 and CYP2C12 are common rat CYP2C genes whereas the more common human CYP2C genes are CYP2C9 and CYP2C19), and the limited amount of data on zonation of human liver metabolising enzymes. Palmer et al.^[75] suggest that members of the CYP3A subfamily are preferentially expressed in hepatocytes in the perivenous region, whereas genes coding for CYP1A2,

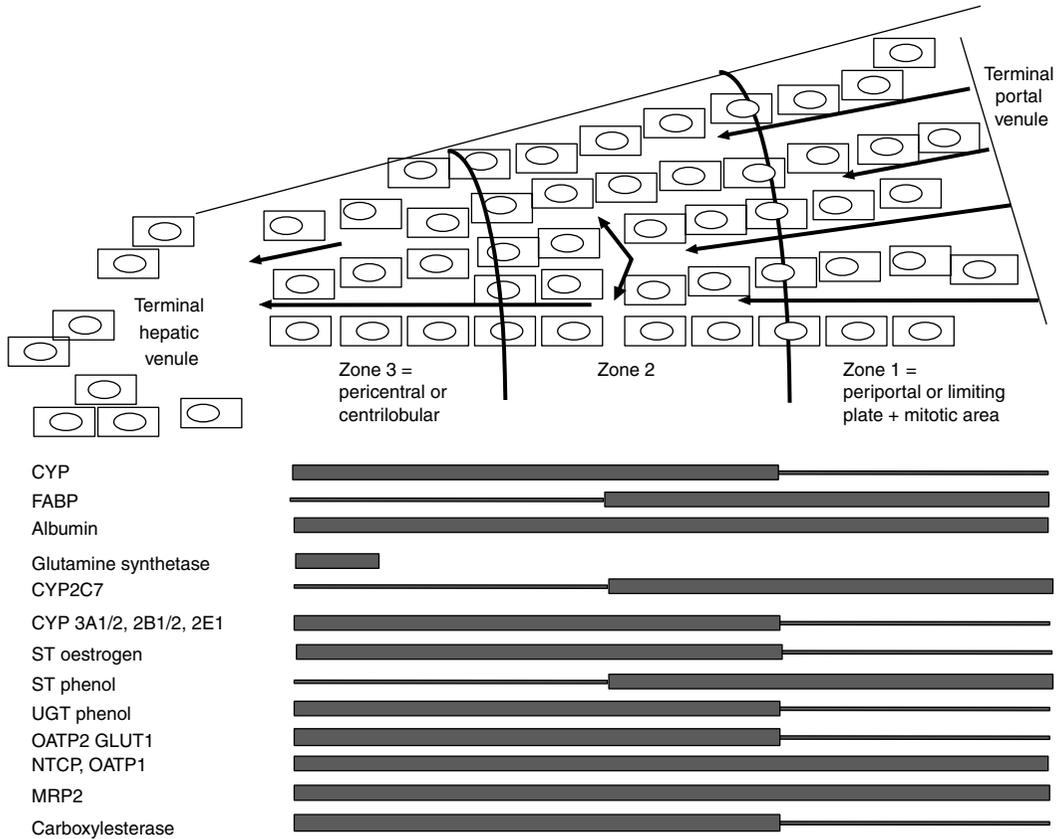


Fig. 5. Examples of heterogeneity of gene expression in the zones of the rat hepatic acinus.^[74,81-100] A thick line indicates gene expression. **CYP** = cytochrome P450; **FABP** = fatty acid binding protein; **GLUT1** = glucose transporter protein 1; **MRP** = multidrug resistance-associated protein; **NTCP** = Na⁺-taurocholate cotransporting polypeptide; **OATP** = organic anion transporting polypeptide; **ST** = sulfotransferase; **UGT** = uridine 5'-diphosphate glucuronosyltransferase.

CYP2A, 2B and 2C are expressed uniformly throughout the liver acinus. In contrast, McKinnon et al.^[76] reported higher levels of CYP1A1 and CYP1A2 in the perivenous region. CYP2E1 is normally localised within the perivenous region.^[77] In human liver disease, the acinar distribution of CYP2E1 and CYP2A follows that of fatty deposits and is most abundant in zone 3.^[78] Zonation of drug-metabolising enzymes in the direction of flow from the periportal to perivenous regions of the hepatic acinus (figure 5) may affect the pharmacokinetics of generated metabolites, but has

been studied only to a limited extent to date.^[72,79] Zonation of CYP drug-metabolising enzymes is absent from the fetal liver.^[80]

The higher expression of CYPs in the perivenous region, especially after xenobiotic induction, together with reduced 'mopping-up' mechanisms for reactive metabolites in this region, may account for increased perivenous xenobiotic toxicity.^[74] Gumucio and Miller^[101] have reported preferential pericentral toxicity for paracetamol, halothane and carbon tetrachloride, among others. The toxicokinetics associated with the production

and clearance of reactive metabolites in terms of zonation appears to have been studied only to a limited extent.

2.6 Transport from the Hepatocytes Across the Canalicular Membrane Into the Bile

Active transport of solutes across the canalicular membrane into the bile by several membrane proteins has been demonstrated. Evidence from the microperfusion of intrahepatic bile duct units suggests that the bile ducts both secrete and absorb water in response to osmotic gradients, actively absorb bile acid through a sodium-dependent mechanism, and transport HCO_3^- .^[23]

Mechanistically, biliary elimination of anionic compounds, including glutathione *S*-conjugates, is mediated by MRP2, whereas bile salts are excreted by a bile salt export pump (BSEP)^[102] (figure 4). MRP2, previously called canalicular anion transporter, uses ATP to transport a number of drug conjugates that undergo enterohepatic recycling, including glucuronide, sulfate and glutathione conjugates. In humans, MRP2 function is defective in Dubin-Johnson syndrome. It has been suggested that MRP2 expressed on the canalicular membrane is homogeneously distributed within the acinus,^[103] as shown in figure 5.

P-glycoprotein, the multidrug resistance (MDR) gene product, is exclusively located on the canalicular membrane of hepatocytes.^[104] Class 1 P-glycoprotein expression from the MDR1 gene mediates the excretion of hydrophobic, mostly cationic, metabolites, whereas class 2 P-glycoprotein expression from the MDR3 gene is involved in phospholipid transport.^[102] Acridine orange, doxorubicin, verapamil, propranolol, phosphatidylcholine, estradiol glucuronide and daunomycin have been shown to inhibit the uptake of daunomycin by P-glycoprotein in isolated rat canalicular liver plasma membranes.^[105]

As discussed later in this review (section 2.7), the expression of hepatic transporters and their activity are regulated in various situations, such as ontogenesis, carcinogenesis, cholestasis, cellular stress and after treatment by hormones and xeno-

biotics. Interestingly, the defective transport of many solutes directly correlates with the absence of BSEP, MDR3, P-glycoprotein and MRP2 transporters in specific liver diseases. Alkaline phosphatase plays an active role in downregulating the secretory activities of the intrahepatic biliary epithelium.^[106] From a clinical pharmacokinetic perspective, as discussed later, various compounds and diseases affect transporter activity and the resulting enterohepatic recycling of drugs. Hence, the enterohepatic recycling of indomethacin glucuronide, which is mediated by canalicular multispecific organic anion transporter/multidrug resistance associated protein 2 (cMOAT/MRP2) is reduced in livers where cMOAT/MRP2 function is hereditarily defective.^[42] Recently, inducible MRP3 on the central membrane has been shown to play a significant role in the cholehepatic circulation of bile salts and in the transport of organic anions such as 17β -estradiol 17β -D-glucuronide in rats.^[107] It is clear that there is much more to learn about transporters, which will greatly increase our understanding of enterohepatic recirculation and modifications of drug kinetics as a result of drug structure, drug interactions and patient disease.

2.7 Active Transport from the Hepatocytes Across the Lateral Membrane Into the Sinusoid

As shown in figure 4, the active transporters MRP1, MRP3 and MRP6 also exist on the lateral membrane and can transport certain drugs and metabolites from hepatocytes back into the sinusoids. In some cases, as has been shown for intracellularly formed morphine glucuronides,^[108] there is preferential excretion of the solutes back into the sinusoids rather than into the bile, thus minimising potential enterohepatic recycling. Other drug anion conjugates are also preferentially excreted into the sinusoids by saturable transport mechanism.^[43] In general, transporters such as MRP1 are present in the lateral membrane only at very low levels in quiescent cells.^[109] However, as discussed later (section 4.8.2), particular conditions such as cholestasis may induce the expression of the MRP3

transporter on the basolateral membrane and facilitate the cellular efflux of drug or drug metabolite anions back into the sinusoids.^[43]

2.8 Bile Transport Into Duodenum

A significant fraction of bile may flow into the duodenum during fasting. Generally, bile is stored in the gall bladder and released into the small intestine via the Sphincter of Oddi on the sight, smell or ingestion of food. As a consequence, gallbladder emptying is sporadic, leading to complex pharmacokinetics when enterohepatic recycling is substantial. Patrick et al.,^[110] in recognising that the multiple peaks in the plasma concentration-versus-time profiles of ezetimibe, conjugated ezetimibe and total ezetimibe were suggestive of enterohepatic recycling, commented that the multiple secondary peaks corresponded to the approximate time of meals. This timing was consistent with food intake stimulating the emptying of the gallbladder.

2.9 Gut Flora

The enterohepatic circulation of many solutes depends on the gut flora metabolism of the solutes and their metabolites excreted in bile (table I). Various enzymes, such as the gut flora cysteine conjugate β -lyase associated with the metabolism of cysteine conjugates, have been identified in the last decade.^[111] Furthermore, drug metabolism by the intestinal microflora contributes to the pharmacological profile of various drugs. The metabolism of orally administered drugs can result in pharmacodynamic changes, and the hydrolysis of biliary drug conjugates is responsible for the enterohepatic circulation of drugs.^[34] The β -glucuronidase activities in humans for the proximal and distal regions of small intestine are 0.02 and 0.9 μmol of substrate degraded/h/g content, respectively.^[34] Activity of bacterial enzymes varies markedly among different host species.

3. Properties of Compounds Undergoing Enterohepatic Recycling

3.1 Size of Drug or Metabolite

The biliary route of excretion plays a major role in the elimination of anions, cations and non-ionised molecules containing both polar and lipophilic groups above a threshold molecular weight. The molecular weight threshold for humans is estimated to be 500 to 600, but it should be emphasised that this is a generalisation. Dyes such as bromosulphophthalein (BSP), indocyanine green (ICG) and various iodinated cholecystographic media and their respective metabolites, which are known to be excreted in human bile, have molecular weights greater than 600. Compounds with molecular weights less than this threshold are excreted primarily in the urine. It is important to recognise in scaling up that the molecular weight threshold differs between species. A minimum molecular weight of between 200^[112] and 325 is required for significant biliary excretion in the rat, and 400 and 475 are the approximate values in the guinea pig and rabbit, respectively.^[113]

Many exogenous substances are secreted into bile and undergo some degree of enterohepatic cycling. These include commonly used drugs (morphine, warfarin, indomethacin, cardiac glycosides), several antimicrobial agents (clindamycin, rifampicin, erythromycin, metronidazole, ampicillin, ceftriaxone and doxycycline), radiocontrast media and potentially toxic heavy metals. Table II lists examples of compounds reported to be excreted in the bile and therefore possibly available for enterohepatic recycling.

3.2 Biliary Excretion of Bile Acids

Bile contains a wide spectrum of anionic organic compounds, among which bile acids are the most abundant. Bile acids undergoing enterohepatic circulation are efficiently extracted from portal blood by liver and are concentrated in bile. Active solute transport into canaliculi is primary responsible for canalicular bile formation. Of all the compounds in bile, bile acids are the most con-

Table I. Examples of processes, drugs and enzymes involved in enterohepatic recirculation (adapted from Ilett et al.,^[34] with permission)

Processes	Drug and compound examples	Enzyme	Sample tested and microflora containing enzymes
Hydrolysis			
Glucuronides	Chloramphenicol glucuronide Estradiol-3-glucuronide	β -Glucuronidase	Human faeces, <i>Escherichia coli</i>
Glycosides	Cycasin	β -Glucosidase	Human intestinal content, <i>Streptococcus faecalis</i> , <i>Eubacterium rectale</i>
Amides	Sennosides	Amide hydrolase	<i>Clostridium sphenoides</i> , <i>E. coli</i> , <i>Proteus vulgaris</i> , <i>Bacillus subtilis</i> , <i>B. mycoides</i>
	Sulfonamides		
Sulfamates	Penicillin Chloramphenicol	Arylsulfotransferase	Human faeces, <i>Streptococci</i>
	Estrone sulfate		
Reductions			
Ketones	Estrone	Hydrogenase	Human faeces
Elimination reactions			
Demethylation	Methylmercury	Demethylase	Human faeces, <i>Streptococci</i> , <i>Lactobacilli</i> , <i>Clostridium</i>
<i>N</i> -Demethylation	Methylamphetamine	<i>N</i> -Demethylase	Human intestinal content, <i>Clostridium</i> , <i>Bacteroides</i>
	Imipramine Choline		
Deamination	Putrescine	Deaminase	<i>Bifidobacteria</i> , <i>E. coli</i> , <i>Bacteroides</i> , <i>Clostridium</i>
	Cadaverine Amino acids		
Decarboxylation	Caffeic acid Phenolic acids	Decarboxylase	Human faeces, <i>S. faecalis</i>
	Amino acids		
Dehydrogenation	Bile acids	Hydroxysteroid dehydrogenase	Human faeces, <i>Clostridium welchii</i>
	Cholanic acids		
Other reactions			
Ring	O-Heterocyclic flavones	Tryptophase	Rat faeces
R-C-S-R'	Propachlor-cysteine	C-S lyase	Pig cecal content, <i>Eubacterium aerofaciens</i> , <i>Fusobacterium necrophorum</i>

centrated. Hepatic bile acid secretion involves uptake at the sinusoidal pole of the hepatocyte, intracellular transport to the canalicular pole and secretion into the canalicular lumen. At least three different transport mechanisms are involved in the uptake of bile acids:^[141-143] (i) Na⁺-dependent bile salt uptake system; (ii) Na⁺-independent carrier-mediated transport; and (iii) nonionic diffusion fol-

lowed by intercellular binding to cytosolic proteins and amidation with taurine or glycine or conjugation with sulfate or glucuronide. The first two mechanisms are responsible for concentrative uptake.^[144]

Most studies on biliary excretion of anions have been reported with bile acids. In general, conjugated bile acids are transported by the Na⁺-bile

acid cotransport mechanism, whereas unconjugated bile acids are transported via the Na⁺-independent mechanism and/or by passive diffusion. Biliary secretion across the canalicular membrane is also a concentrative process mediated by an electrogenic, carrier-mediated transport mechanism.^[142,145] This is consistent with the finding that only negatively charged bile acids are efficiently secreted into bile.^[146] Uptake of conjugated bile acids is Na⁺-dependent and thought to be mediated by a Na⁺-bile acid co-transport (or symport) system, energised by the transmembrane Na⁺ gradient maintained by the Na⁺, K⁺-ATPase. Bile acid binding proteins have been identified but their role in transport has not been established. There is a lobular gradient in bile acid transport, periportal hepatocytes removing most of the bile acid load at physiological concentrations. For some unconjugated bile acids, conjugation may be the limiting factor in overall transport from blood to bile. Canalicular secretion occurs via carrier-mediated diffusion sensitive to the membrane po-

tential. Hepatic extraction of bile acids and bile salts from portal blood plasma is an extremely efficient process in animals as well as in humans, and the extraction efficiency appears to be inversely related to the extent of bile acid binding to serum albumin.^[147,148] Hence, while the latter increases with increasing lipophilicity, the hepatic extraction efficiency is higher for the relatively hydrophilic cholates (taurocholate > glycocholate > cholate) than for the more hydrophobic dihydroxy (e.g. chenodeoxy- and deoxycholate) and mono-hydroxy (e.g. lithocholate) derivatives.^[147,149,150] Furthermore, conjugation with taurine or glycine significantly increases the maximal uptake velocity (V_{max}) via a saturable and Na⁺-dependent transport process.^[146,151] In contrast, the relatively hydrophobic unconjugated dihydroxy bile acids are largely taken up into hepatocytes by nonsaturable (probably nonionic) diffusion.^[149,152,153] Finally, a Na⁺-independent saturable uptake process has been repeatedly suggested to be involved in overall

Table II. Examples of drugs, other foreign chemicals and their metabolites excreted in human bile

Amiodarone ^[114]	Estrone ^[115]	Phenol red ^[113]
Ampicillin ^[113,116]	Ezetimibe ^[110]	Phenolphthalein ^[117]
Benzylpenicillin ^[118]	2-Fluoro- β -alanine ^[119]	Phenytoin ^[117]
Bilirubin ^[120]	Gentamycin ^[118]	Pivampicillin ^[118]
Bromocresol green ^[121]	Glibenclamide (glyburide) ^[5]	Rifamide ^[113]
Bromosulphothalein ^[122]	Gliclazide ^[123]	Rifamycin ^[113]
Cefixime ^[124]	Imipramine ^[125]	Roquinimex ^[126]
Ceftriaxone ^[127]	Indocyanine green ^[122]	Rose bengal ^[121]
Cefatrizidime ^[128]	Indomethacin ^[125]	Spirolactone ^[125]
Cephaloridine ^[113]	Irinotecan ^[129]	Sulfamethoxazole ^[113]
Cephmandole ^[118]	Lanatocid C ^[118]	Sulindac ^[125]
Cephazolin ^[118]	Lorazepam ^[130]	Subactam ^[116]
Chenodeoxycholic acid ^[122,125]	Lormetazepam ^[6]	Temafloxacin ^[131]
Chloramphenicol ^[118]	Methotrexate ^[113]	Testosterone ^[125]
Chlortetracycline ^[118]	Metronidazole ^[117]	Tetracycline ^[113,118]
Clindamycin ^[117]	Mezlocillin ^[132]	Thiamphenicol ^[113]
Demethylchlortetracycline ^[118]	Morphine ^[133]	Tolfenamic acid ^[134]
Diazepam ^[113]	Mycophenolic acid ^[135]	Toremifene ^[136]
Digitoxin ^[113]	Mycophenolate mofetil ^[137,138]	Troglitazone ^[139]
Digoxin ^[113]	Nortriptyline ^[113]	Trovafoxacin ^[7]
Doxycycline ^[118]	Novobiocin ^[113]	Ursodeoxycholic acid ^[122,125]
Erythromycin ^[113]	Oltipraz ^[140]	Valproic acid ^[125]
Estradiol ^[125]	Pethidine (meperidine) ^[113]	Warfarin ^[133]

bile acid uptake, especially for the unconjugated trihydroxy bile salt cholate.^[141,153]

The processes responsible for vectorial transport of organic anions from the sinusoidal side to the canalicular pole of the hepatocyte are still poorly understood. Part of the transfer may be directed by intracellular protein binding (see section 5.2). Within the cell, organic anions are largely bound to the cytosolic protein ligandin, also known as Y protein or glutathione *S*-transferase B,^[154,155] FABP or Z protein,^[156] and Y'binders. The last group of identified proteins, which includes among others 3 α -hydroxysteroid dehydrogenase and a steroid sulfotransferase, has been claimed to play a role in the translocation of bile acids.^[55]

3.3 Bilirubin

Bilirubin is the main degradation product of haem catabolism in mammals and has ideal properties for biliary excretion. The major source is haemoglobin, which is degraded in the reticuloendothelial system of the spleen, bone marrow and liver. The molecule is an open-chain tetrapyrrole (molecular weight 585) with two vinyl, four methyl and two propionic acid side chains. Because the pKa value is around 4.4,^[157] the carboxylic acid groups would be expected to ionise almost completely at physiological pH in aqueous medium, so an unconjugated bilirubin would have polar properties. Surprisingly, the pigment is almost insoluble at pH 7.4.^[158] This unexpected finding can be explained by intramolecular hydrogen bonding, which involves all polar groups of the molecule.^[159] The formed bilirubin is reversibly bound by plasma albumin. The liver takes up bilirubin circulating in the plasma rapidly and almost exclusively.^[120] The uptake is consistent with a carrier-mediated membrane transport.^[160] Albumin is important in restricting bilirubin to the plasma compartment and directing it to the liver for its metabolic clearance. In albuminemic rats, the plasma disappearance of an injected bolus of radiolabelled bilirubin is higher than in normal rats, but the recovery in bile is slower.^[161] This demonstrates that in the absence of albumin the pigment

repartitions indiscriminately in all body tissues and less is available for hepatic uptake.

Early work provided strong evidence that a characterised BSP- and bilirubin-binding protein in liver plasma membranes was responsible for the uptake of these anions.^[162] A monospecific antibody against this membrane protein reduced the uptake of bilirubin and BSP, in contrast to free fatty acids and bile acids.^[163] Bilirubin, BSP and ICG share a common transport system, which appears to be distinct from that for bile acids.^[164-166] Specific high-affinity binding sites in the plasma membrane maintain a plasma/cell concentration gradient of the ligands. The cytosolic binding protein ligandin also serves to maintain a concentration gradient within the cytosol from the plasma membrane to canalicular membrane.^[167] At the present time, although bilirubin monoglucuronide is known to be transported by OATP1 and human liver-specific transporter (HLST), the transporter of unconjugated bilirubin in the sinusoidal membrane has not as yet been identified.^[41] Kamisako^[41] note that unconjugated bilirubin may also cross the hepatocyte sinusoidal membrane by a diffusion process.

In the liver cell, bilirubin is largely bound to the two cytosolic binding proteins with high affinity, ligandin (Y-protein) and Z-protein.^[155] Ligandin (molecular weight 46 000), identical with glutathione *S*-transferase B,^[154] binds a variety of compounds including bilirubin, haem, steroids, BSP, ICG and carcinogens.^[168] Approximately 30% of the bilirubin present in cytosol is bound with low affinity to Z-protein,^[155] which seems mainly to be involved in the transport of fatty acids.^[160] Both binding proteins are supposed to play a role in the transport of bilirubin from the liver cell membrane to the conjugating enzyme system in the endoplasmic reticulum.

In the endoplasmic reticulum, bilirubin is conjugated by bilirubin UGT (UGT1A1) to form mono- and diglucuronides of bilirubin.^[41] The bilirubin conjugates formed in the endoplasmic reticulum are secreted in bile. Virtually all bilirubin excreted in bile is in the form of glucosidic conju-

gates. Bilirubin conjugates are secreted across the liver canalicular membrane by MRP2, a member of the ATP-binding cassette transporter family.^[41] Finally, there is regurgitation of bilirubin glucuronides into blood under conditions of impaired biliary bilirubin excretion. MRP3, which is located in the lateral membrane, transports bilirubin glucuronides into blood.^[41]

3.4 Other Anions and Anionic Drugs

Beside bile acids and bilirubin, bile contains varying amounts of endogenous organic anions, such as steroid hormones. In addition, anionic xenobiotics, including dyes like ICG and BSP, cholecystographic contrast agents and certain antibacterials, are readily excreted into bile.^[122] Biliary excretion of organic anions is the result of a complex sequence of events, involving uptake by hepatocytes at the sinusoidal side of the cell, intracellular trafficking with or without biotransformation, and secretion at the canalicular pole of the cells, as discussed in details in section 2.

Most nonsteroidal anti-inflammatory drugs (NSAIDs) are metabolised by the liver, with subsequent excretion into urine or bile. Enterohepatic recirculation occurs when a significant amount of an NSAID or its conjugated metabolites are excreted into the bile and then reabsorbed in the distal intestine.^[169]

The clinical pharmacokinetics of therapeutic bile acids, chenodeoxycholic and ursodeoxycholic acids, have also been reviewed.^[122] The absorption of chenodeoxycholic acid by the intestine is better than that of ursodeoxycholic acid. Both solutes are extracted efficiently by the liver, conjugated with glycerine and taurine, secreted in bile, and then undergo enterohepatic circulation with the endogenous bile acids. These therapeutic bile acids are metabolised by intestinal bacteria to lithocholic acid, which is mainly excreted with faeces.

A significant enterohepatic recirculation has been reported to delay the final elimination of a large fraction of tricyclic antidepressants taken in large quantities during suicide attempts.^[170]

The compound calcitriol (1,25-dihydroxy-vitamin D₃), critical in calcium absorption, is formed by the sequential hydroxylation of calciferol (vitamin D₃) in the liver and kidney, and excreted in the bile as polar metabolites (such as glucuronides and, possibly, sulfates and neutral polar steroids) which undergo enterohepatic recirculation in both humans and experimental animals.^[171]

The topoisomerase I inhibitor SN-38, a minor metabolite of irinotecan, is glucuronidated in the liver, then excreted in the bile. Enterohepatic recirculation occurs after the glucuronide is deconjugated to SN-38 by β -glucuronidase produced by the intestinal flora, and subsequently reabsorbed.^[129] The major metabolite of 5-fluorouracil, 2-fluoro- β -alanine, and its conjugates are also believed to undergo enterohepatic recycling.^[119]

3.5 Organic Cations

Many drugs are compounds with one or more tertiary or quaternary amine functions. Quaternary ammonium compounds are permanently positively charged, whereas tertiary amines can be protonated and therefore become positively charged at physiological pH depending on the compound's pK_a value. Among the monoquaternary drugs are anticholinergic, antineoplastic and antihelminthic drugs, whereas the biquaternary drugs represent predominantly neuromuscular blocking agents. Cationic drugs can pass membranes by carrier-mediated transport, and such processes have been characterised in kidney tubular cells,^[172] in the intestine,^[173] in the choroid plexus,^[174] and in the liver.^[10,143,175] Several routes for hepatic transport of organic cations have been described based on the structure-kinetic relationship of the compound. Factors involved in the extent of biliary excretion of organic cations are molecular size,^[176] number of charged groups and lipophilicity of the agents. Protein binding, metabolism and the chemical structure may also influence the relative contribution of this excretion route.^[177]

Organic cations of relatively low molecular weight (<200) tend to be poorly excreted in bile. Therefore a so-called 'molecular weight threshold' has been proposed, above which the biliary excretion of drugs becomes appreciable (>0% of the dose). For monovalent organic cations, this threshold is suggested to be 200 ± 50 in the rat.^[112] Lipophilicity of the organic cations seems to be a good physicochemical parameter to predict the extent of biliary output. The biliary excretion rate of the type I cations (for example triethylmethylammonium, tripropylmethylammonium, procainamide ethobromide, thiazinamium, and *N*-methyldepropine) generally increases with increasing lipophilicity.^[176,178] A molecular weight of 500 to 600 has been proposed as a threshold for hepatobiliary excretion of most bivalent type II organic cations (for example D-tubocurarine, trimethyltubocurarine, stercuronium, vecuronium, *N*-propyldeoxyajmalinium, pancuronium and hexafluronium) in the rat, guinea pig and rabbit.^[179] Similar to the type I cations, the balance between lipophilicity and hydrophilicity appears to be an important factor determining hepatic transport at the uptake and secretion level.^[180]

Carrier-mediated transport of cationic drugs in uptake into and secretion from cells is, in principle, saturable, so that a nonlinear kinetic pattern is to be expected. Potential transport interactions between drugs can be anticipated based on chemical structure. However, many drugs can be present in various forms (dissociated, undissociated, complexed) and therefore transported by various carrier systems at the same time. Carrier-mediated transport is not restricted to the plasma membranes of the secretory cells; various organelles within the cells can accumulate drugs via similar translocation mechanisms, leading to persistent storage of such drugs, changes in organelle function or local toxicity.^[180,181] Apart from carrier-mediated uptake processes, an alternative uptake route is proposed for some hydrophilic cations. For instance, stercuronium,^[182] decamethonium and hexamethonium^[179] might be solely accumulated by an endocytotic uptake mechanism. As discussed in sec-

tion 2.4, intrahepatic binding is a major determinant in the hepatic disposition of the lipophilic cations.

3.6 Other Drug Examples – Hormones and Antimicrobial Agents

The extent of biliary excretion of endogenous and exogenous hormones (natural or synthetic) differs significantly depending upon the compound investigated. In general, it may be assumed that steroids are excreted into bile to different extents depending on their structure: metabolites of less polar steroids, e.g. progesterone, show a greater tendency to be excreted into bile than metabolites of polar steroids (e.g. hydrocortisone). In humans, biliary excretion of exogenously administered estradiol, estrone or estriol was found to be 65, 48 or 23% of the dose per 12 hours, respectively.^[183-185] With progesterone, approximately 30% of the applied dose was excreted in bile following intravenous injection, whereas 13% of an intravenous dose of testosterone and only 4% of intravenously administered hydrocortisone were reported to be excreted into bile by patients with biliary fistula.^[183,186,187] Biliary metabolites of hormones are at least partly reabsorbed from the intestine and may undergo extensive enterohepatic circulation. This results in a prolongation of their overall elimination half-lives. Intestinal metabolism, especially deconjugation reactions of glucuronide and sulfate adducts, plays a central role in this process.^[188] The pharmacokinetics and enterohepatic cycling of estradiol have been recently reported after three oral single-dose administrations of equimolar doses of estradiol alone, estradiol plus desogestrel and estradiol valerate in a three-way cross-over mode in 18 healthy postmenopausal women.^[115]

The mechanisms responsible for the biliary excretion of antimicrobials are poorly understood quantitatively. It cannot be predicted from the molecular weight or other physicochemical properties whether an antibacterial agent will be excreted to a significant extent by the hepatobiliary route. The molecular weights of the third generation cephalosporins are very similar: ceftazidime (molecular

weight 546) is almost exclusively excreted by the renal route in contrast to ceftriaxone (molecular weight 554), which is excreted about 40% by the hepatobiliary route.^[127,132] Similar figures can be given for azlocillin (5%) and mezlocillin (25%). In humans, the biliary route is not the predominant route of elimination despite the fact that hepatic elimination plays an important role for many drugs. For new antibacterial agents, a detailed balance of urinary and faecal excretion, preferably after intravenous administration, should be determined. Data on concentrations in common duct bile and gallbladder bile should also be provided.

Amiodarone, a type III antiarrhythmic, is poorly bioavailable (20 to 80%) and undergoes extensive enterohepatic circulation before entry into a central compartment.^[114]

3.7 Formation of Glutathione and Glutathione Adducts

The concentration of glutathione (GSH) in bile depends on the species studied, the assay conditions and the intrahepatic glutathione status. Most of the glutathione in bile is in form of GSH, and only a small fraction is oxidised glutathione (GS-SG). The excretion of GSSG and glutathione *S*-conjugates is an important mechanism to eliminate potentially toxic compounds from the organism and the liver. The role of GSH in bile is less clear. It clearly plays an important role in the elimination of potentially toxic metals from the liver. The biliary excretion of many metals is closely coupled to the secretion of GSH.^[189] Zinc, methylmercury, cadmium, chromium and lead are excreted as GSH complexes, and depletion of hepatic GSH decreases the biliary excretion of the compounds together with the excretion of GSH.^[190-192] In species where substantial concentrations of GSH are preserved throughout the biliary tree, the tripeptide might play a role in the detoxification of potentially mutagenic or carcinogenic compounds generated in the upper gastrointestinal tract.^[193]

3.8 Biliary Excretion of Glucuronide and Glutathione Conjugates

Biliary excretion of paracetamol glucuronide has been shown by using MRP2-deficient rats be facilitated by cMOAT/MRP2.^[194] MRP2 is responsible for the biliary excretion of a variety of endogenous and exogenous organic anions, including many glucuronide conjugates, some glutathione conjugates and a few sulfate conjugates.^[195] The majority of MRP2 substrates identified to date have at least two negative charges.^[195] Glucuronide conjugates, depending on their reactivity, can covalently bind to proteins within hepatocytes. However, glucuronides are also likely to be secreted into the bile across the canalicular plasma membrane, where they could react with targets in the biliary tree. Hepatobiliary transport of anionic xenobiotics, including many types of glucuronide and glutathione conjugates, normally occurs by ATP-dependent transport, which is mediated by the multispecific conjugate export pump cMOAT.^[195-197] Indomethacin glucuronide is an example of one metabolite excreted into the bile by MRP2.^[42] As discussed in section 2.7, MRP3 is also involved in glucuronide excretion. A direct transport function of MRP2 for glutathione conjugates and leukotriene C₄ has recently been demonstrated by expression of the carrier in COS-7 cells and in *Xenopus laevis* oocytes.^[198] Glutathione conjugates appear to be a poor substrate for MRP3.^[107]

It should be noted that most of our information on transporters has come from rats. Overall, the primary active transport of organic anions across rat bile canalicular membrane is greater than in humans, although the difference between the species is less marked for glucuronides.^[199]

4. Factors Affecting Biotransformation and Enterohepatic Circulation

The absorption, biotransformation and enterohepatic circulation of a xenobiotic can be influenced by factors including the role of bile acids; the biological influences of species variation, gen-

der, age and developmental stage; nutritional status; disease states; and pharmacological effects of other chemicals after intentional administration or inadvertent exposure.

4.1 Bile Acids

The primary metabolic pump for enterohepatic circulation is the active excretion of bile acids across the canalicular membrane into bile. The rate-limiting step is the transfer of bile acids derived either from blood or from synthesis into bile. Bile acid depletion, induced by either prolonged biliary diversion or administration of cholestyramine, markedly decreases the biliary excretion of digitoxin and troglitazone,^[139,200] bilirubin,^[201] ICG, bromocresol green, rose bengal, BSP and eosin.^[202,203] In contrast, increasing the bile acid pool, as occurs with diabetes, increases the biliary excretion of bromocresol green, ICG, phenol red and rose bengal.^[121,204]

4.2 Chemical Adsorbents

Chemical adsorbents may stimulate movement of endogenous compounds and xenobiotics through the intestine.^[205] In rats and in humans, cholestyramine markedly increased the faecal excretion of the pesticide chlordecone.^[206-208] This can only partly be attributed to interruption of enterohepatic circulation, because chlordecone is also excreted into the intestine by transmucosal transport.^[209] In rhesus monkeys, dietary supplementation with 4% cholestyramine enhanced faecal excretion of pentachlorophenol from 3 to 54%.^[210] Consequently the urinary excretion decreased from 35 to 5%, and thus the net effect of treatment was a 40% enhancement of the total excretion of pentachlorophenol.^[211]

Cholestyramine, a nonabsorbable anion exchange resin, effectively binds bile acids and anionic drugs. Oral administration of cholestyramine was shown to decrease the absorption and biliary excretion of bile acids in rats.^[202,212] Oral administration of cholestyramine or activated charcoal diminished the urinary recovery of an intravenous dose of paracetamol by 20 to 30%, indicating sig-

nificant biliary excretion of paracetamol conjugates and almost complete reabsorption.^[213] Moreover, cholestyramine treatment was effective in preventing paracetamol-induced hepato- and nephrotoxicity.^[214] Activated charcoal adsorbs paracetamol,^[213] carbamazepine, phenobarbital, and phenylbutazone^[215] in the intestinal tract, preventing their absorption and future enterohepatic circulation.

Oral ingestion of activated charcoal also appears to inhibit enterohepatic recycling of estriol, as shown in figure 6. Recently, Elomaa^[216] examined the extent to which oral activated charcoal affected the bioavailability of norethisterone acetate and gestodene during steady-state conditions by inhibiting their enterohepatic recirculation. They suggested that the enterohepatic circulation of the steroids was not of clinical importance and that women on oral contraceptives can take activated charcoal for the treatment of diarrhoea when administered 3 hours after and at least 12 hours before pill intake.

4.3 Species and Age Differences

Variations among species in the capacity for hepatobiliary transport^[10] are primarily responsible for species differences in enterohepatic circulation.^[218] Interspecies variations in hepatic and extrahepatic biotransformation are well known.^[219-221] Moreover, species variations in gastrointestinal microflora may result in significant qualitative and/or quantitative differences in microbial biotransformation.^[222] Studies in several species indicate that β -glucuronidase activity is higher in intestine from rats and mice than intestine from humans.^[223]

In addition to metabolic differences, the anatomical, physiological and biochemical differences between the gastrointestinal tracts of humans and the common laboratory animals can cause significant variation in drug absorption from the oral route. Among the physiological factors, pH, bile, pancreatic juice, and mucus and fluid volume and content can modify dissolution rates, solubility, transit times, and membrane transport of drug mol-

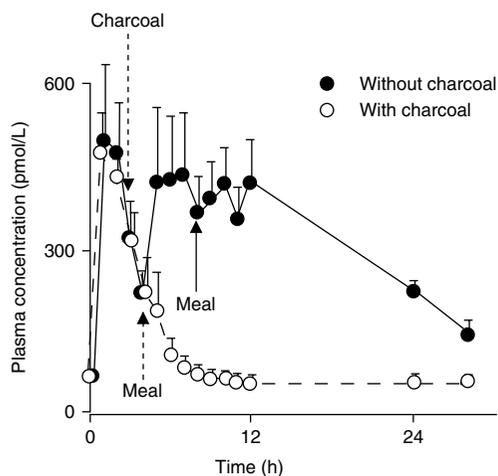


Fig. 6. Plasma levels of estriol after oral administration of estriol 12mg without and with 20g of activated charcoal. Values are means \pm standard error for $n = 8$ (reproduced from Heimer and Englund,^[217] with permission).

ecules.^[224] Enterohepatic circulation is also subject to age-related effects in the development of, for example, hepatic uptake mechanisms for bile acids,^[225] exogenous organic acids^[226] and hepatic excretory function.^[227] Aging also decreases the rates of phase I and phase II biotransformations in rat colon.^[228]

4.4 Xenobiotics

Exposure to xenobiotics may alter the biotransformation, hepatobiliary transport and intestinal absorption of a given chemical. Exposure of animals and humans to microsomal enzyme-inducing agents may increase biliary excretion by augmenting biotransformation and increasing bile formation. A specific interaction is the increased conjugation with glucuronic acid induced after exposure to numerous drugs which facilitates the biliary excretion of paracetamol,^[229] alfentanil and sufentanil,^[230] bilirubin,^[231] and diethylstilbestrol and phenolphthalein.^[232]

4.5 Intravenous Colloidal Nanoparticles

Enterohepatic recirculation may also be facilitated by the intravenous administration of a drug

(for example, indomethacin) formulated using colloidal nanoparticles, which are then taken up by liver Kupffer cells to facilitate the drug's absorption into liver parenchymal cells.^[233]

4.6 Coadministration of Drugs

Drug interactions can occur at any part of the enterohepatic cycle, through effects on absorption, distribution, transporters, metabolism or other mechanisms. Sections 4.2 and 4.4 examined the effects of adsorbents and certain xenobiotics on the enterohepatic recycling process. A number of studies have been carried out on inhibition of carrier-mediated plasma membrane transport. For instance, the hepatic membrane transport of 1-*O*-gemfibrozil- β -D-glucuronide is carrier-mediated and inhibited by the organic anion dibromosulphophthalein.^[234] Clofibrac acid increases the hepatic clearance but decreases the biliary clearance of 1-*O*-gemfibrozil- β -D-glucuronide.^[234]

Kim et al.^[235] have suggested that a number of drugs, including peptidomimetic renin inhibitors, propranolol, cyclosporin and progesterone, are potent inhibitors of the bile acid uptake transporter NTCP, responsible for the uptake of bile acids from the portal circulation into hepatocytes. Anti-arrhythmic agents, including bupivacaine, lidocaine and quinidine, were found to enhance the polypeptide activity.

The modulator LY-335979 is among the most potent modulators of P-glycoprotein, and does not modulate resistance mediated by MRP1 or MRP2. LY-335979 significantly enhances the survival of mice implanted with P-glycoprotein-expressing murine leukaemia (P388/ADR) when administered in combination with anticancer agents (daunorubicin, doxorubicin or etoposide).^[236]

Shipkova et al.^[135] have recently suggested that cyclosporin attenuates the enterohepatic recirculation of mycophenolic acid glucuronide/mycophenolic acid in kidney transplant recipients. The reduction in the bioavailability of mycophenolate mofetil in liver transplant patients treated with antibacterials for selective bowel decontamination is

probably due to an interruption of enterohepatic cycling.^[137]

4.7 Genetic Defects and Biliary Transport

Three (bile acids, phospholipids and bilirubin) of the four (the other is cholesterol) major substrates transported out of hepatocytes have their own transport mechanisms. Bile acids and phospholipids each have a specific transporter, and that for bilirubin is multispecific. In each of these transporters, isolated autosomal recessive defects have now been identified and have helped to confirm the physiological role of these proteins.^[22] An abnormality in plasma membrane aminophospholipid distribution leading to a secondary defect in bile acid transport has also been identified.^[22] Studies using isolated perfused livers from MRP2-deficient Wistar rats have shown that the hepatobiliary disposition of paracetamol is impaired relative to normal rats.^[194] There was also upregulation of a basolateral organic anion transporter in the MRP2-deficient rat livers. The mutant strain Eisai-hyperbilirubinaemic rats (EHBR) used in certain pharmacokinetic studies by Kato et al.^[197] have a hereditary defect in cMOAT/MRP2. Using this same model, Morikawa et al.^[237] showed that the transport of 17 β -estradiol 17 β -D-glucuronide across the bile canalicular membrane is predominantly mediated by cMOAT/MRP2.

4.8 Effect of Disease

The biliary excretion of drugs is a unique route of elimination; enterohepatic circulation and intestinal metabolism may also determine the fate of a drug in the organism. Liver and kidney diseases might influence the biliary excretion of drugs by decreasing the uptake into hepatocytes and altering the distribution and metabolism within hepatocytes. The disease might also decrease the transfer mechanism into bile. In animals, cholestasis results in an inhibition of hepatic uptake^[238,239] and drug metabolism,^[240] whereas hepatocellular damage decreases the transfer from the hepatocyte into bile.^[241] In rats with ureter ligation, the biliary excretion of total paracetamol was enhanced to 38%

of the dose compared with 28% in normal rats, mainly due to an enhanced sulfation of paracetamol.^[241]

Relevant to the clinical pharmacokinetics of enterohepatic recirculation is the coordinated regulation and function of sinusoidal and canalicular transporters to produce bile. In the diseased state, downregulation of the unidirectional sinusoidal uptake system NTCP will protect the hepatocyte from further intracellular accumulation of bile salts. In contrast, expression of the canalicular bile salt export pump maintains bile salt secretion even in complete biliary obstruction.^[242] In cholestasis, there is also a strong downregulation of canalicular MRP2 and a converse upregulation of hepatocyte basolateral efflux systems such as MRP3 and MRP1.

Interruption of enterohepatic circulation can be used in patients with toxic drug concentrations.^[125] α -Amanitin, a toxic substance from *Amanita phalloides*, which potently inhibits DNA-dependent RNA polymerase II,^[243] undergoes extensive enterohepatic circulation.

4.8.1 Cirrhosis

Cirrhosis is known to be associated with alterations in both hepatic blood flow and cellular function, including biliary secretion. Thus, it is not surprising that the hepatic uptake rate of many substrates is reduced.^[244,245]

Two hypotheses have been proposed to account for the decreased uptake.^[244,246] The intact cell hypothesis attributes the reduced uptake to changes in the liver microcirculation. These changes include capillarisation of hepatic sinusoids, collagenisation of the space of Disse and intrahepatic shunting, resulting in reduced perfusion of liver cells that contain normal enzymatic activity.^[247-250] The sick cell hypothesis attributes the decreased uptake to a reduction in cellular transport capacity. In cirrhosis, the lipid composition of the hepatocyte plasma membrane and membranes from intracellular organelles is altered.^[251,252] As well, Na⁺-K⁺ ATPase activity has been reported to be decreased.^[253,254] Recently, the effect of CCl₄-induced experimental hepatic failure on the sequen-

tial hepatobiliary transport of the model organic cations triethylmethylammonium and tributylmethylammonium in a rat model was reported.^[255] It was suggested that the hepatic failure affected sinusoidal membrane transport systems (uptake and/or efflux) rather than those on the bile canalicular membrane (excretion). We have reported that transport can be related to the degree of fibrosis.^[62]

4.8.2 Cholestasis

The term 'cholestasis' literally means stagnation of bile flow and can be due either to physical obstruction of the biliary tree, termed extrahepatic cholestasis, or a decrease in the secretion of bile by the hepatocyte, termed intrahepatic cholestasis. Extrahepatic cholestasis involves a mechanical obstruction to the bile duct that may be caused by duct stones, sclerosing cholangitis or pancreatic carcinoma.

Intrahepatic cholestasis is defined as a syndrome with decreased bile flow and regurgitation of biliary constituents such as bile acids, conjugated bilirubin and lipids into plasma. Intrahepatic cholestasis is caused by an impairment of normal biochemical processes within the hepatocyte. Permeability of the canalicular membrane and tight junctions to bile acids are affected. Significant changes result in many subcellular membranes, including the actin-containing microfilaments (implicated in bile duct contraction) and the endoplasmic reticulum. Also affected are the intracellular bile salt transport proteins responsible for transporting bile salts from the plasma membrane to the canalicular membrane (intracellular trafficking) and bile salt metabolism. Interruption of bile flow results in elevated serum bile acid levels as well as cytosolic levels.^[256] Cholestasis also affects the CYP system, possibly through bile salt retention. Bile salts have been shown to inhibit CYP *in vitro*, which may lead to hypertrophy of the smooth endoplasmic reticulum.^[257]

The morphological features of cholestasis, both intra- and extrahepatic, include the presence of biliary material (primarily bile pigments) in hepatocytes and Kupffer cells, and bile thrombi in dilated

bile canaliculi.^[258] Electron microscopy shows a decreased number of canalicular microvilli, and distorted microvilli with a blunted or oedematous appearance. Hypertrophy of the Golgi apparatus and smooth endoplasmic reticulum is seen in addition to increased lysosomes and dilation and vesiculation of the rough endoplasmic reticulum.^[259] It is not possible to distinguish extra- and intrahepatic cholestasis on the basis of morphological features alone, substantiating the idea that many of the morphological changes are the result of, rather than the cause of, cholestasis.

The alterations of hepatobiliary transport that occur in cholestasis can be divided into transporter defects (such as mutations of transporter genes or acquired dysfunctions of transport systems that cause defective canalicular or cholangiocellular secretion) and those that result from biliary obstruction.^[242] Multiple alterations of hepatocellular transporter expression can be caused by the hepatocellular accumulation of toxic cholephilic compounds in progressive familial intrahepatic cholestasis or by the direct inhibition of transporter gene expression by cytokines.

α -Naphthylisothiocyanate (ANIT) produces cholestasis in a reproducible and dose-dependent manner following its acute administration, whereas chronic administration results in bile duct hyperplasia and biliary cirrhosis.^[260] Plaa and co-workers^[261,262] have shown that a single dose of ANIT to the rat causes retention of BSP, an organic anionic dye excreted primarily in bile, within 2 to 4 hours and hyperbilirubinaemia at 12 to 24 hours. Bile flow decreases within 16 to 24 hours and remains depressed for about 5 days.^[263] A dose of ANIT greater than 150 mg/kg causes complete cholestasis in the rat, whereas the mouse is more sensitive, the hamster less sensitive, and the dog and rabbit resistant to ANIT-induced cholestasis.^[262]

Phalloidin, cytochalasin and colchicine are thought to induce cholestasis by their toxic actions on microfilaments and microtubules in the hepatocyte. These organelles are thought to be important for maintenance of cell shape, motility and secre-

tion of various substances, including lipoproteins, proteins and bile. Phalloidin is taken up into the hepatocytes, apparently by the bile acid transport system and is thus concentrated in the liver.^[264,265] Chronic treatment of rats with low doses of phalloidin (500 µg/kg/day for 3 or more days) causes a marked accumulation of actin microfilaments in the pericanalicular region of the hepatocyte^[266] and a decrease in bile flow.^[267] Many of the bile acids are hepatotoxic and will induce cholestasis if infused at rates that exceed the transport maximum for their excretion in bile. Thus, infusion of taurocholate at increasing concentrations increases bile flow to a maximum. Further infusion decreases bile flows rather than maintains it at the maximal rate.

Cholestasis has been shown to have a number of effects on transporters involved in the biliary excretion process. Overall, there appear to be adaptive processes whereby bile salts not able to exit the hepatocyte in cholestasis are effectively removed across the basolateral membrane.^[268] Transcriptional and post-transcriptional mechanisms are involved in this high level of regulation, with experimental evidence supporting communication between hepatocytes and cholangiocytes, especially in primary biliary cirrhosis. In cholestatic rats, the multispecific organic anion transporter MRP2 has a decreased expression. The expression of the bile salt export pump BSEP and the phospholipid transporter MDR2 are less affected. Interestingly, as a balance to NTCP, the carrier protein for hepatic uptake of bile salts, being sharply downregulated, a basolateral ATP-dependent transporter for glucuronides and bile salts (MRP3) is upregulated.^[43]

4.8.3 Diabetes

Diabetes mellitus is a commonly occurring disease, characterised by the inability of the body to produce or use insulin, and often accompanied by various pathological states including coronary heart disease, renal insufficiency, cerebrovascular disorders and neuropathy. In addition to these disorders, there is some evidence that the course of diabetes may also alter the biotransformation of

pharmaceutical agents and increase the risk of drug toxicity and adverse effects. Early researchers found an increased incidence of gallstones at autopsy of patients with diabetes,^[269,270] as well as an increased frequency of cholelithiasis among diabetic patients.^[271] Other workers have shown changes in biotransformation due to diabetes, both in humans and in animal models,^[272,273] as well as alterations in pharmacokinetic parameters for various drugs and xenobiotics.^[274,275] Other studies have shown elevated mRNA expression of some plasma membrane proteins.^[276] All these studies suggest that diabetes mellitus has a profound influence on hepatobiliary structure and function.

The influence of insulin on hepatic protein synthesis may impact on the protein carriers controlling hepatic uptake and canalicular secretion, and on the enzymes involved in hepatic biotransformation. For example, the rate of uptake of isoniazid into liver, lung, diaphragm and brain is enhanced in the presence of insulin.^[277] Recent studies suggest that hepatic protein synthesis is also affected by insulin-dependent diabetes.^[278]

4.8.4 Renal Disease

Renal disease has been suggested to affect biliary transport through the accumulation of some endogenous P-glycoprotein substrates/modulators in the plasma in disease states.^[279] Using glycerol-induced acute renal failure, Huang et al.^[279] showed that, associated with an increase in P-glycoprotein level in the kidney, P-glycoprotein function in the liver and brain was suppressed.

4.8.5 Impaired Biliary Bilirubin Excretion

Impaired bilirubin secretion in the bile will lead to active transport of bilirubin glucuronides back into blood via MRP3.^[41] Interestingly, Kato et al.^[197] found that the biliary excretion of telmisartan is relatively unaffected in Eisai-hyperbilirubinemic rats relative to normal rats, and suggested that telmisartan glucuronidation is enhanced in the Eisai-hyperbilirubinemic rats so that, overall, there is comparable excretion of glucuronide in both the mutant and normal animals.

4.8.6 Transporter Malfunction and Disease

In patients with inflammation-induced icteric cholestasis (mainly cholestatic alcoholic hepatitis), the mRNA levels of the basolateral transporters NTCP, OATP2 and BSEP are reduced by about 40% when compared with controls.^[280] In addition, mRNA levels for the apical transporter MRP2 remained unchanged, but canalicular immunolabelling for MRP2 was also decreased. The mRNA expression of MRP3, MDR1, MDR3, and familial intrahepatic cholestasis gene 1 (FIC1) remained unchanged. Zollner et al.^[280] suggest that reduced expression of hepatobiliary transporters for bile salts and other organic anions may contribute to inflammation-induced cholestasis in humans.

5. Pharmacokinetics of Enterohepatic Recycling

Because of the recycling of solutes between the intestinal tract, portal circulation and liver via the biliary tract, drugs subject to enterohepatic recirculation are often characterised by a long half-life and multiple peaks. As an example of this process, the enterohepatic cycling of furosemide acyl glucuronide, followed by hydrolysis, results in a second and slow elimination phase with a half-life of 20 to 30 hours.^[281] A similar increase in half-life is seen for estrone after administration of estradiol (figure 7). The pharmacokinetics of the immunosuppressant mycophenolate mofetil also show secondary peaks of mycophenolic acid, suggesting a significant enterohepatic cycling process.^[282] Bullingham et al.^[282] suggested that enterohepatic cycling must involve colonic bacterial deconjugation of phenolic glucuronides and, based on an oral cholestyramine interaction study, calculated that the mean contribution of enterohepatic cycling to the AUC_{∞} of mycophenolic acid was around 40%. More recently, Funaki^[283] used a population analysis approach to define the pharmacokinetics of mycophenolate mofetil.

5.1 Compartmental Models

The first models developed to describe enterohepatic cycling were classical compartmental

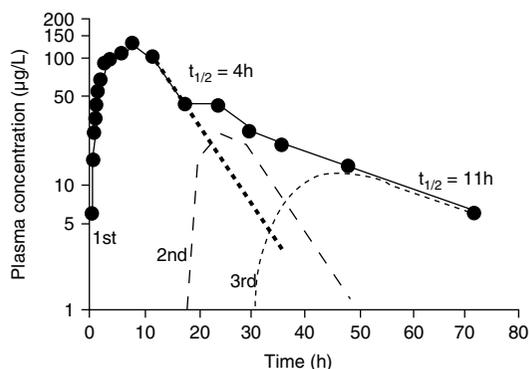


Fig. 7. Plasma concentration-time profile of estrone after oral administration of estradiol 1.5mg in a desogestrel combination to 18 healthy postmenopausal women. Also shown are the second and third enterohepatic recycling phases leading to secondary peaks and an increase in the half-life ($t_{1/2}$) of oestrone (reproduced from Vree and Timmer,^[115] with permission).

models. Thus, two- (or three-) compartment models of drug disposition with first-order absorption after oral administration (gut as a well-mixed compartment) were extended by a chain of compartments describing the transport of the drug from the central compartment to the gut.^[284-286] In its simplest form, this chain consists of only one compartment, the gallbladder (figure 8).

Recirculation loops of more than one compartment allowed a better simulation of the delay caused by the recirculation process. Such a model was successfully applied to describe enterohepatic cycling of morphine,^[287] valproic acid^[288] and morphine-3-glucuronide^[289] in the rat. Although these models with a continuous delay time appear sufficient for animals without gallbladder (like the rat), they are not appropriate to describe the discontinuity in the enterohepatic cycling process caused by gallbladder emptying. Enterohepatic cycling models that account for the effect of gallbladder emptying (lag-time models) can further be classified in models where emptying is assumed to occur at regular intervals (τ , 2τ , ...) and models with specific irregular emptying times (τ_1 , τ_2 ,... τ_n). Steimer et al.^[290] showed that a simple two-compartment lag-time model already accounts for the occurrence of second peaks in the plasma concen-

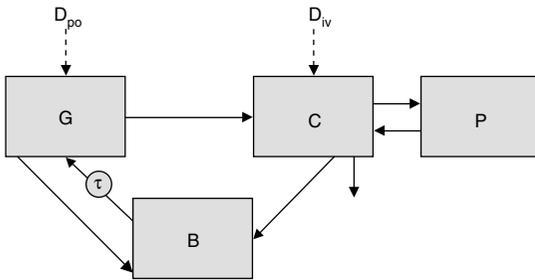


Fig. 8. Compartment model for enterohepatic cycling. **B** = gallbladder; **C** = central compartment; **D** = dose; **G** = gastrointestinal tract; **iv** = intravenous; **P** = peripheral compartment; **po** = oral; τ = lag-time.

tration-time profile. A general treatment of enterohepatic recirculation of drugs can be based on the fraction of drug in the systemic circulation that is excreted in the bile and the fraction of drug reabsorbed from the gut that reaches the systemic circulation in each enterohepatic cycle.^[291] A four-compartment enterohepatic model for unchanged drug described by Wang and Reuning^[292] contained central, peripheral, gallbladder and intestinal compartments and an intermittent gallbladder-emptying rate constant.

Models based on irregular lag-time intervals are more near to physiological reality since gallbladder emptying starts when food begins to be digested in the upper gastrointestinal tract. Furthermore, the extent of biliary emptying may depend on the quantity of fat in the meal. Models with regular biliary emptying have been studied extensively in the literature,^[293,294] but models with irregular intervals are mathematically more complex. Although the latter were first developed for only one or two recirculations,^[295] these models have been later formulated for any number of cycles occurring at various times.^[296,297] However, their application through data analysis has been restricted to two recirculations, τ_1 and τ_2 . As shown by Plusquellec et al.^[297] for the pharmacokinetics of an antihypertensive drug in humans, this may not be a limitation for many drugs as only two (or one) recirculations could be detected (fitting the plasma concentration-time data by analytical func-

tions derived from the model). In using compartmental models with regular (or continuous) gallbladder emptying, the resulting differential equations are usually solved numerically. For certain compartmental structures, equations for mean residence time in the body (MRT), mean absorption time (MAT), the area under the concentration-time curve (AUC), clearance (CL) and bioavailability (F) have been derived.^[64,285,286,298]

5.2 System Dynamics Approach

A generalisation of these compartmental models is possible on the basis of a system theoretical approach using the method of numerical inverse Laplace transformation.^[299] The integration of the physiological dispersion model description of vascular transit times in the liver^[36] with enterohepatic recirculation has been described using this approach.^[300] This approach uses subsystems to model the processes of intestinal absorption, systemic disposition and enterohepatic cycling, and is much more flexible. The basic structure of the recirculatory system describing enterohepatic cycling (figure 9) is analogous to that of pharmacokinetic models of circulatory drug transport, and the corresponding equation for the plasma concentration-time profile $[C(t)]$ after intravenous bolus administration can readily be written in the Laplace domain^[299,301] (equation 1):

$$C(s) = \frac{D_{iv}}{CL^1 F_B} \frac{f_D(s)}{1 - f_{EHC}(s) f_D(s)}$$

where D_{iv} is the intravenous dose, CL^1 is the total clearance without enterohepatic cycling and F_B denotes the biliary excretion ratio.

Any appropriate mathematical function can be assumed – only their Laplace transforms $f_D(s)$ and $f_{EHC}(s)$ must exist – for the systemic drug disposition function $f_D(t)$ [processes inside the body] and the transit time density describing enterohepatic cycling $f_{EHC}(t)$ [processes ‘outside’ the body, i.e. intestinal tract, portal vein, liver and bile duct]. Note that for a system without enterohepatic cy-

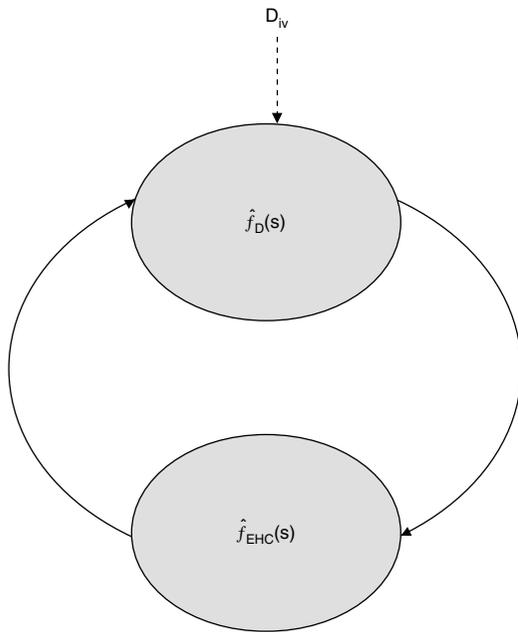


Fig. 9. Modelling of enterohepatic cycling (EHC) in pharmacokinetics by a circulatory arrangement of subsystems characterised by residence time density functions $\hat{f}_D(s)$ and $\hat{f}_{EHC}(s)$ describing drug disposition and EHC in the Laplace domain.^[299]

cling (e.g. bile duct cannulation), equation 1 reduces to equation 2:

$$C^I(s) = \frac{D_{iv}}{CL^I F_B} f_D(s)$$

Calculating the moments of $C(t)$ from its Laplace transform $C(s)$ [equation 1], equations for pharmacokinetic parameters such as MRT, AUC and F can be derived that are independent of the specific forms of the disposition and enterohepatic cycling functions. Thus, enterohepatic cycling causes AUC and MRT to increase (equation 3 and equation 4):^[299]

$$AUC_{iv} = AUC^I \frac{1}{1 - F_{EHC} F_B}$$

$$MRT_{iv} = MRT^I + \frac{(MRT^I + MTT_{EHC}) F_{EHC} F_B}{1 - F_{EHC} F_B}$$

where $AUC^I = D_{iv}/CL^I$ and MRT^I denote the measures without enterohepatic cycling, and F_{EHC} and MTT_{EHC} are the availability and mean transit time from the intestinal tract to the systemic circulation, respectively. In other words, $E_{EHC} = 1 - F_{EHC}$ is the extraction in a single enterohepatic circulation. Note that ‘without enterohepatic cycling’ means blockade by cholestyramine and antibacterials or biliary cannulation, situations where one can estimate the extent of enterohepatic recirculation from a comparison of the respective AUC values (figure 10).

This concept (equation 1) was used to study the disposition of cefixime^[299] and diclofenac^[302] in rats. The basic model equation can also be readily extended to the case of oral administration and/or multiple doses. When the enterohepatic cycling function contains a lag time, it accounts for discontinuous (but regular) gallbladder emptying and secondary peaks in the $C(t)$ curve appear if the lag time increases. Recently, Yasui et al.^[301] have applied the recirculatory concept to evaluate the stereoselective enterohepatic cycling and the unidirectional chiral inversion of ketoprofen enantiomers in the rat. Although the system approach proposed by Yamaoka et al.^[299] was up to now only applied to analyse the pharmacokinetics of drugs after intravenous administration (equation 1), it can be extended to account for the first-pass effect after oral administration as suggested by Peris-Ribera et al.,^[291] who developed more general equations for AUC, F and CL based on mass-balance considerations. This would allow, for example, the assumption of a more realistic absorption process where the fractional absorption rate is a function of time instead of a constant (as in the case of the first-order absorption process in compartmental modelling).

Equations that describe the effect of enterohepatic cycling on drug bioavailability, F , are more complex than equation 3. It has been shown that enterohepatic cycling increases F and that this effect can be pronounced for drugs with high biliary excretion, where small changes in the pharmaco-

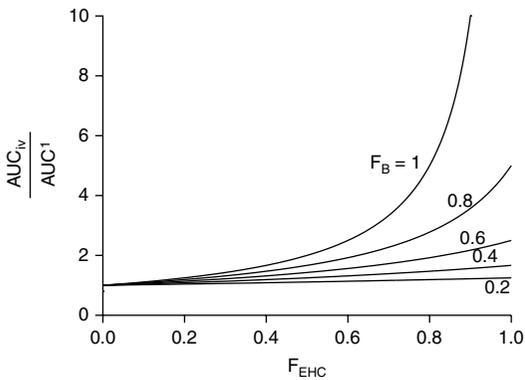


Fig. 10. Effect of the drug absorption ratio, F_{EHC} , on the ratio of areas under the concentration-time curve with (AUC_{IV}) and without (AUC^1) enterohepatic recycling for different values of the excretion ratio into the bile (F_B) [equation 3].

kinetic parameters of the drug can cause large changes in F .^[64]

5.3 Parameter Estimation

The construction of a model that is sufficiently realistic from a pharmacokinetic (or physiological) point of view is one problem, the identifiability of the model, i.e. the reliability of parameter estimation based on available concentration-time data, is another. Experimental design strategies for quantifying pharmacokinetic models with enterohepatic cycling were studied by Wang and Reuning.^[292] These authors also discussed the advantage of getting additional information from bile sampling (a necessary condition for the identification of more complex models). Recently, a simple three-compartment enterohepatic cycling model (based on a one-compartment disposition model) has been applied in a population pharmacokinetic study of mycophenolate mofetil.^[283] Figure 11 is an illustration of that modelling, showing the secondary peak for mycophenolate mofetil after gallbladder emptying. Recently, Ploeger et al.^[303] used scaling from a rat to a human physiologically-based pharmacokinetic model to describe the enterohepatic cycling of glycyrrhetic acid metabolites.

5.4 Enterohepatic Circulation, or Some Other Mechanism?

Multiple peaks in the concentration-time profile could be due to mechanisms other than enterohepatic circulation. A double-peak behaviour could, for instance, be due to biphasic gastric emptying behaviour. Wang et al.^[304] attributed the double peak of alprazolam to a reduction in gastric motility caused by its muscle relaxant effect. They suggested that enterohepatic recycling be precluded from being the underlying mechanism as the presence of double peaks was not evident after an intravenous dose. Solutes undergoing enterohepatic recirculation are frequently characterised by a relatively high hepatic extraction and low plasma concentrations, as illustrated by the pharmacokinetics of chenodeoxycholic and ursodeoxycholic acids, where majority of bile acid is confined within the enterohepatic circulation. When the enterohepatic recirculation is efficient and dominant, the solutes may also be characterised by a high apparent volume of distribution and a low clearance.

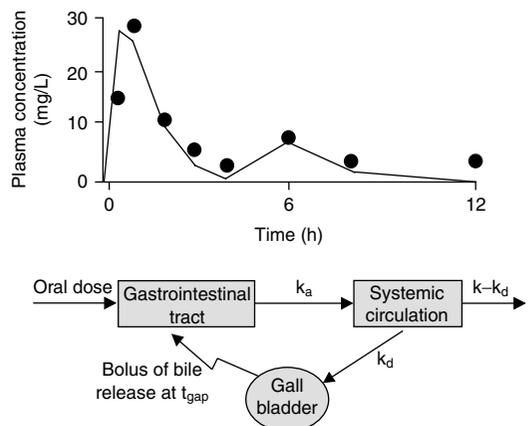


Fig. 11. NONMEM fit of mycophenolate mofetil plasma concentration-time profiles for a typical patient and the associated enterohepatic circulation model used (reproduced from Funaki,^[283] with permission). k = elimination rate constant; k_a = absorption rate constant; k_d = excretion rate constant into gallbladder; t_{gap} = expulsion time of gallbladder.

6. Clinical Implications of Enterohepatic Recirculation

The impact that enterohepatic cycling has on the pharmacokinetics and pharmacodynamics of a drug depends on: (i) the importance of biliary excretion of the compound relative to renal and metabolic clearance processes; and (ii) the efficiency with which the compound is absorbed into the circulation from the gastrointestinal tract. Enterohepatic circulation influences drug pharmacokinetics by increasing drug bioavailability and influencing plasma concentration curves of the drug.^[297,305] The absorption of a drug is usually more rapid than its elimination. If drug elimination includes enterohepatic circulation, drug secreted in bile can serve as a secondary source for drug absorption. Enterohepatic cycling may influence drug pharmacokinetics by prolonging elimination. This has been shown to be of clinical significance for spironolactone.^[306] The clinical significance of the enterohepatic circulation depends on the pharmacological or toxicological properties of the biliary excretory products, their availability for absorption, and whether the absorbed products are re-extracted by the liver or pass into the general circulation. In some cases, the enterohepatic circulation of a drug may be a therapeutic advantage. For example, sulindac, an NSAID, is recycled through the biliary system to a greater degree than its active sulfide metabolite.^[307] Thus, sustained plasma concentrations may be achieved by the enterohepatic circulation of the prodrug, minimizing exposure of the intestine to active drug. On the other hand, it is also possible that enterohepatic cycling leading to secondary peaks in plasma concentration could result in toxicity.

A limited number of studies involving enterohepatic recirculation have considered pharmacodynamic outcomes. The slow elimination phase associated with the enterohepatic cycling of furosemide acyl glucuronide, for instance, coincides with a pharmacodynamic rebound phase of urine retention.^[281] These investigators suggested that accumulation of furosemide and its acyl glucuronide in patients with end-stage renal failure results

from infinite hepatic cycling and that dose increments are required to maintain a required level of diuresis. A pharmacodynamic model associated with enterohepatic recycling has been described.^[308]

Most clinical studies for quantification of the biliary excretion of drugs have been performed in patients following cholecystectomy or other biliary tract surgery. Such studies should be interpreted with caution, as the patients generally have had some kind of hepatobiliary disease and may have experienced extrahepatic cholestasis. There is a possibility that their disease may have influenced some component of the enterohepatic recirculation of the drugs. Patients with cholestatic liver disease show a lower biliary excretion of drugs, increasing the risk of toxicity.^[258,260] Although there are a number of studies on the kinetics of biliary excretion of drugs,^[5-7,33] the results reflect the small number of patients in many studies, the methods of bile sampling and the influence of hepatobiliary diseases. Future research should obtain more information about biliary excretion of xenobiotics in humans based on improved methods of bile sampling and specific drug analyses.

A number of studies have used a T-tube inserted in the common bile duct of patients to study biliary pharmacokinetics. Most of these studies have been performed in post-surgical patients with T-tube drainage, and it should be recognised that this method of bile collection is not ideal because bile flow and composition are often severely altered during the period of study, not all bile is collected and enterohepatic circulation is partially interrupted.^[118] Some examples of these studies are as follows. Serra et al.^[309] studied the excretion of free and conjugated BSP in 19 adult patients with choledocholithiasis. Westphal et al.^[124] studied the hepatobiliary extraction profile of cefixime, a dianionic cephalosporin antibacterial, in 10 patients after cholecystectomy. They suggested that cefixime biliary clearance was nonlinear, mostly in its initial phase, which is consistent with a concentrative uptake and intracellular protein binding. The biliary pharmacokinetics of sulbactam and

ampicillin have also been reported for 19 patients with normal liver function after surgery of the biliary tract.^[116] The biliary excretion of chlorthalidone^[310] and methadone and its metabolites^[311] was low. In contrast, the biliary excretion of the glucuronide/sulfate conjugates of hydroxylated derivatives of tolfenamic acid was found to be significant.^[134] High concentrations of ceftazidime in the bile have been reported, but with a low biliary excretion in terms of the dose used.^[128] Biliary excretion of unchanged temafloxacin and temafloxacin glucuronide measured by T-tube only accounted for approximately 2.2 and 1% of the administered dose, respectively.^[131]

The effect of antimicrobials on female hormones remains controversial, especially in terms of the effects on the enterohepatic recycling of oral contraceptive steroids. Substantial differences have been observed in plasma estradiol levels in patients treated with ampicillin 250mg every 8 hours in the 48 hours before delivery (figure 12).^[312] It is also suggested that there is probably a group of susceptible women who have lower plasma concentrations of oral contraceptive hormones and ex-

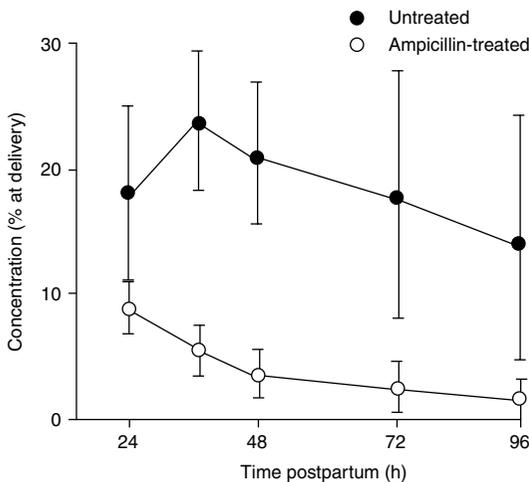


Fig. 12. Postpartum plasma total estradiol concentrations (as a percentage of concentrations at delivery) in patients treated with ampicillin for 48 hours prior to delivery and untreated patients (reproduced from Buchan and Klopper,^[312] with permission).

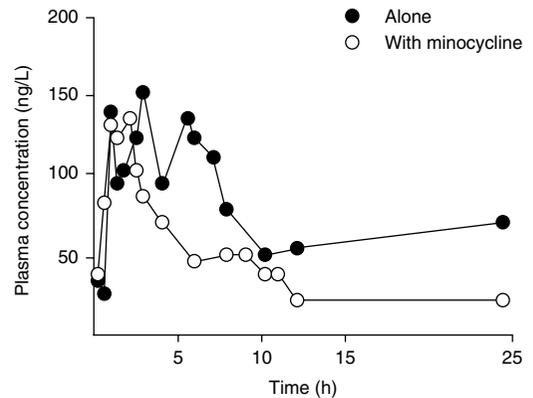


Fig. 13. Plasma concentration-time profiles of ethinylestradiol on day 15 of a pill cycle on two occasions in the same individual taking a triphasic contraceptive pill with and without minocycline 100mg twice daily.^[313]

perience breakthrough bleeding or become pregnant when given broad-spectrum antibacterials. This may relate to interruption of the enterohepatic recirculation of ethinylestradiol (figure 13).^[313,314] Weisberg^[315] suggests that while the effectiveness of oral contraceptives may be impaired by concomitant treatment with antimicrobials, there are no scientific data to support the anecdotal evidence that concomitant use reduces contraceptive efficacy in the majority of women. It is thus apparent that a number of drugs currently in clinical use have the potential to interact to affect not only the liver plasma membrane transporter activities and hepatic bile acid uptake, but also enterohepatic recirculation.

Also indirectly affecting enterohepatic recirculation are a multitude of drug interactions that affect the production of polar metabolites by CYP, conjugation, transporters, intestinal transit time and bioavailability, amongst others. Associated with such drug interactions is interindividual variability in the inhibition and induction of enzymes,^[316] with varying degrees of clinical significance as a consequence.

7. Conclusions

Enterohepatic recirculation is dependent on a range of factors associated with gastrointestinal

absorption, hepatic uptake, metabolism and biliary excretion. Each of these processes is dependent on species, age, disease, drug, genetics, physiology and coadministered drugs. An effect on any one of these processes may interrupt or enhance the enterohepatic cycle. Many of the determinants affecting these processes are still being defined. Compartmental and noncompartmental models originally developed to analyse the pharmacokinetics of enterohepatic recycling in rats are only partly applicable to humans. Models that account for irregular biliary emptying are more realistic in clinical pharmacokinetics; however, based on drug plasma concentration data, reliable parameter estimation may be possible for one or two recirculations only.

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Correspondence and offprints: Dr *Michael S. Roberts*, Department of Medicine, University of Queensland, Princess Alexandra Hospital, Brisbane, QLD 4102, Australia.
E-mail: mroberts@medicine.pa.uq.edu.au