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Concentrations of Gd-BOPTA in cholestatic fatty rat livers: role of transport functions through membrane proteins

Catherine M. Pastor^a*, Michael Wissmeyer^b and Philippe Millet^c

Gd-BOPTA (gadobenate dimeglumine) is a magnetic resonance (MR) contrast agent that, after i.v. administration, distributes within the extracellular space, enters rat hepatocytes through the sinusoidal transporters organic anion transporting peptides (Oatps) and is excreted unchanged into bile through the multidrug resistance-associated protein 2 (Mrp2). It is unclear how the hepatobiliary contrast agent would accumulate in cholestatic fatty livers from obese rats with bile flow impairment. Indeed, the expression of both Oatps and Mrp2 transporters is decreased in cholestatic hepatocytes. To assess this question, we measured on-line the hepatic concentrations of 153Gd-BOPTA with a gamma probe placed over perfused rat livers. During the perfusion of 153Gd-BOPTA, we obtained a similar maximal hepatic concentration in normal and fatty livers despite the decreased expression and function of membrane transporters in fatty livers. By pharmacokinetic modeling and mathematical simulations, we show how changes of transport into and out of hepatocytes modify the concentrations of 153Gd-BOPTA within hepatocytes. Mathematical simulations help to understand how each parameter (entry into hepatocytes, bile excretion, or efflux back to sinusoids) interferes with the hepatic concentrations. The hepatic concentrations of 153Gd-BOPTA within hepatocytes rely on the entry into hepatocytes through the sinusoidal membrane and on two paths of exit, the efflux back to sinusoids and the elimination into bile. Understanding how 153Gd-BOPTA accumulates in hepatocytes is then complex. However, such understanding is important to analyze liver imaging with hepatobiliary contrast agents in cholestatic fatty livers. Copyright © 2012 John Wiley & Sons, Ltd.

Keywords: MR hepatobiliary contrast agent; liver MRI; cellular transport

1. INTRODUCTION

Nonalcoholic fatty liver diseases include a large spectrum of abnormalities characterized by an increased intrahepatic triglyceride content (steatosis) with or without inflammation and fibrosis (1). The transition from fatty livers to more severe disease (or steatohepatitis) is triggered by inflammation. Cholestasis (or decreased bile flow) is an important marker of this increased severity. Cholestasis is associated with the dysregulation of proteins involved in the transport of bile acids, phospholipids, sterols and organic anions through hepatocytes (2-5). Interestingly, when the bile excretion of xenobiotics and endobiotics or their metabolites is impaired, the upregulation of transporters located in the sinusoidal membrane may favor their efflux back into sinusoidal blood, preventing a potentially harmful accumulation within hepatocytes (6). The expression of hepatic transporters during cholestasis is modified, but the consequences on the hepatic accumulation of hepatobiliary compounds are largely unknown. Interestingly, various liver imaging techniques have been developed to assess fatty livers in patients (7-9). Few experimental and clinical studies have used magnetic resonance imaging (MRI) with hepatobiliary contrast agents to characterize these livers (10-12).

We previously showed that hepatic concentrations of ¹⁵³Gd–BOPTA can be measured on-line using a gamma scintillation probe placed over rat livers (13). Gd–BOPTA, the active moiety of gadobenate dimeglumine (MultiHance®, Bracco Milan), is an MR contrast agent that distributes within the extracellular hepatic

space and enters into hepatocytes through the sinusoidal transporters organic anion transporting peptides (Oatp1a1, Oatp1a4, and Oatp1b2) (14). Gd–BOPTA is not metabolized during its transport to the canalicular membrane and is excreted unchanged into bile through Mrp2 (14,15). At liver MRI, Gd–BOPTA remains within the extracellular space in rat livers with severe cirrhosis that lack sinusoidal transporters (16).

Because cholestatic fatty livers have a decreased expression of Oatp1a4 (17,18) and Mrp2 (17,19), we hypothesized that Gd–BOPTA would differently accumulate within cholestatic hepatocytes. Moreover, cholestasis may also modify the transport of Gd–BOPTA through the transporters. The aim of the present study was then to compare the pharmacokinetics of Gd–BOPTA concentrations in normal and cholestatic rat fatty livers and to

- * Correspondence to: C. M. Pastor, Laboratoire de Physiopathologie Hépatique et Imagerie Moléculaire Hôpitaux Universitaires de Genève, Geneva, Switzerland. Email: catherine.pastor@hcuge.ch
- a C. M. Pastor Laboratoire de Physiopathologie Hépatique et Imagerie Moléculaire
- b M. Wissmeyer Service de Médecine Nucléaire
- c P. Millet
 Unité de Neurophysiologie Clinique et Neuroimagerie Hôpitaux Universitaires de Genève, Geneva, Switzerland

understand how the transport through cell membrane proteins generates the concentrations within cholestatic hepatocytes.

2. MATERIALS AND METHODS

2.1. Animals

Before liver isolation, Zucker fat (fa/fa) and Zucker lean (fa/+) rats were anesthetized with pentobarbital (50 mg kg⁻¹ i.p.). The protocol was approved by the animal welfare committee of the University of Geneva and the veterinary office and followed the guidelines for the care and use of laboratory animals.

2.2. Perfusion of Isolated Rat Livers

Livers were isolated and perfused *in situ* with a Krebs–Henseleit-bicarbonate (KHB) solution \pm contrast agents using a nonrecirculating system (Fig. 1) (13). In each experiment, the flow rate was 30 ml min⁻¹ and the bile duct was cannulated for bile sampling collection.

2.3. Quantification of Hepatic Concentrations

Gd–BOPTA was prepared by adding ¹⁵³GdCl₃ to a 0.5 M Gd–BOPTA solution (1 MBq ml⁻¹), which contained a slight excess of the ligand BOPTA (20,21). To assess the hepatic concentrations induced by the perfusion of the extracellular contrast agent Gd–DTPA (gadopentetate dimeglumine, Magnevist[®], Bayer, Zürich), Gd–DTPA was also prepared by adding ¹⁵³GdCl₃ to a 0.5 M solution of Magnevist[®]. Gd–DTPA is an extracellular contrast agent, which has the same extracellular distribution as Gd–BOPTA but does not enter into hepatocytes (13,14). Both contrast agents were then diluted in KHB solution to obtain a 200 µm concentration to be perfused in rat livers.

To quantify the on-line hepatic concentrations of ¹⁵³Gd-labeled contrast agents, a gamma-scintillation probe that measures radioactivity every 20 s was placed 1 cm above the liver (Fig. 1A) (21). To transform radioactivity count rates into contrast agent amounts, the radioactivity in the entire liver at the end of each experiment was measured (Activimeter Isomed 2000, MCD

A. Solutions of perfusion [min] B. Pharmacokinetic modeling 75 105 n 10 45 **KHB KHB** C1 153Gd-153Gd k_{12} **DTPA BOPTA** k_{45} C2 C5 k_{42} k₅₆ robe СЗ C6 **Count rates**

C. Histology of the liver

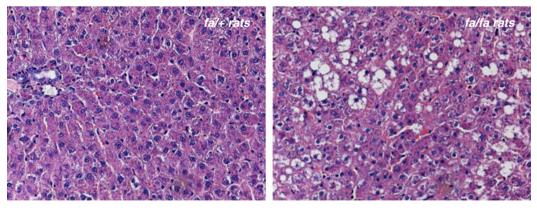


Figure 1. (A) Experimental protocol and data obtained during liver perfusion with the contrast agents 153 Gd–DTPA (200 μm during 10 min) and 153 Gd–BOPTA (200 μm during 30 min). The flow rate being 30 ml min $^{-1}$, the amount of 153 Gd–DTPA perfused was 60 μmol and that of 153 Gd–BOPTA was 180 μmol. To quantify the on-line hepatic concentrations of 153 Gd–labeled contrast agents, a gamma-scintillation probe that measures radioactivity every 20 s was placed above the liver. KHB, Krebs–Henseleit-bicarbonate solution. (B) Pharmacokinetic modeling of 153 Gd–BOPTA transport in rat perfused livers. Entry of contrast agent into the liver is modeled as a zero-order infusion rate K_{in} . Compartment C1 illustrates the dilution of 153 Gd–BOPTA at the beginning and at the end of perfusion. C2 is the extracellular space, C4 and C5 are two hepatocyte-associated spaces. C3 represents the amount of contrast agent in perfusate, and C6 in bile. First-order rate constants k_{ij} are defined with respect to i, the compartment of origin, and j, compartment of convergence. (C) Histology of the liver from obese (fa/fa) and normal (fa/+) rats is illustrated.

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Nuklear Medizintechnik, Germany) and related to the last count rates measured by the probe.

2.4. Experimental Protocol

We perfused five livers isolated from fa/+ (control Zucker) rats and five livers isolated from fa/fa (fat Zucker) rats with KHB solution (15 min, recovery period), 153 Gd–DTPA (10 min), KHB solution (rinse period, 30 min), 153 Gd–BOPTA (30 min) and KHB solution (rinse period, 30 min; Fig. 1). Both contrast agents were perfused with a 200 μ m concentrations. During each perfusion, we measured 153 Gd–BOPTA bile concentration (nmol g $^{-1}$), 153 Gd–BOPTA bile excretion rate (nmol min $^{-1}$ g $^{-1}$), 153 Gd–BOPTA concentration within the liver (nmol g $^{-1}$) and bile flow (μ l min $^{-1}$ g $^{-1}$).

2.5. Liver Tests and Histology

Alanine amino transferases, cholesterol and triglycerides in serum were determined using a serum multiple analyzer. Liver tissue was fixed, embedded in paraffin ($4\,\mu\text{M}$ thick sections), and analyzed under light microscopy after hematoxylin and eosin staining.

2.6. Simulations in the Pharmacokinetic Model

In perfused livers isolated from control fa/+ rats, the compartment model that best described the amounts of contrast agents over time in perfusate, bile and liver is illustrated in Fig. 1B (13,21). Each compartment describes the amount of contrast agent over time. Compartment 1 reflects the mixing between KHB and contrast agent solutions when the solutions of perfusion are changed and accounts for the progressive changes of concentrations before the final concentrations (200 or 0 µm) are reached. Compartment 2 (C2) describes the amount of contrast agent in the extracellular space and compartments 4 (C4) and 5 (C5) the amounts in hepatocytes from two distinct compartments. Indeed, the model distinguishes two compartments of hepatocytes C4 and C5 with different concentrations over time (13,21). The transfer between C4 and C5 is unidirectional. From both compartments, a transfer into compartment 6 (C6 or bile compartment) occurs. Compartment 3 represents the amount of contrast agent in the outflow perfusate leaving the liver. Entry of contrast agent into the system is modeled by a zero-order input rate K_{in} over a time period. In the first-order rate constant k_{ii} , i represents the compartment of origin and j the compartment of convergence. The model was implemented in the MATLAB Software (2010b). It was assumed that the rate constant k_{24} is associated with Oatp-mediated hepatic uptake of 153 Gd–BOPTA. k_{46} and k_{56} describe Mrp2-dependent canalicular excretion from both hepatocellular compartments and k_{42} is associated with the efflux of ¹⁵³Gd-BOPTA from hepatocytes back to sinusoids. Pharmacokinetic modeling of contrast agent transport was compared between fa/fa and fa/+ rats.

Additionally, we conducted mathematical simulations and evaluated how modifications of a single parameter representing $^{153}\text{Gd}\text{-BOPTA}$ entry into hepatocytes (k_{24}) , efflux back from hepatocytes to the extracellular space (k_{42}) , or bile excretion from both hepatocellular compartments $(k_{46}$ and $k_{56})$ modify the hepatic concentration of $^{153}\text{Gd}\text{-BOPTA}$. These simulations mimic the abnormalities that may be observed in cholestatic hepatocytes.

2.7. Statistics

Parameters are means \pm SD. A Mann–Whitney test compared means between fa/+ and fa/fa rats. Two-way ANOVA compared the evolution over time of parameters between the experimental groups.

3. RESULTS

3.1. Characteristics of fa/+ and fa/fa Rats

The fa/fa rats (8 week old) are overweight and have significantly higher serum triglyceride and cholesterol concentrations than fa/+ rats (Table 1). Liver weights were greater in obese rats. During liver perfusion, portal pressure was similar in both groups. Moreover, bile flow was significantly lower in fa/fa (0.6 \pm 0.1 μ l min⁻¹ g⁻¹) than fa/+ (1.3 \pm 0.1 μ l min⁻¹ g⁻¹) rats (p < 0.008, data not shown). Histology was normal in fa/+ rats but steatosis was present in fatty livers (Fig. 1C).

3.2. Concentrations of ¹⁵³Gd-BOPTA in Hepatocytes from Normal and Fatty Livers (T0-T75)

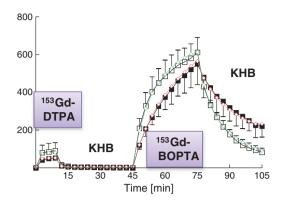
During the perfusion with the extracellular contrast agent ¹⁵³Gd–DTPA (Fig. 2A), the hepatic concentrations rapidly increased to a low steady state (<100 nmol g⁻¹). ¹⁵³Gd–DTPA was then rapidly washed out from the liver during the subsequent KHB perfusion and, as expected, no ¹⁵³Gd–DTPA was detected in bile (Fig. 3C). At time 45 min (T45), ¹⁵³Gd–BOPTA was perfused during 30 min before washing with KHB solution. ¹⁵³Gd–BOPTA hepatic concentrations were much higher than that of ¹⁵³Gd–DTPA and maximal at T75 (Fig. 2A).

To assess the concentrations of 153 Gd–BOPTA within hepatocytes, we must withdraw the concentrations induced by the extracellular distribution of the contrast agent (see Fig. 3A). We can then measure an initial hepatocellular uptake index (IHUI, mol min $^{-1}$ g $^{-1}$) from time 46 min (T46) to T49, a period with minimal 153 Gd–BOPTA excretion adequate to measure the entry of 153 Gd–BOPTA from sinusoids to hepatocytes (Fig. 3A). The IHUI was significantly decreased in fa/fa rats (27 ± 5 nmol min $^{-1}$ g $^{-1}$) in comparison to the control group (48 ± 11 nmol min $^{-1}$ g $^{-1}$; Fig. 3B, p=0.008). After this initial period, concentrations inside hepatocytes reflect both hepatocyte uptake and exit.

In contrast to the hepatic concentrations of ¹⁵³Gd–BOPTA shown in Fig. 2A (those obtained by the gamma probe), the accumulation and elimination concentrations in Fig. 3D and E only reflect ¹⁵³Gd–BOPTA concentrations within hepatocytes. At T75, the maximal concentrations were similar in both

Table 1. Characteristics of fa/fa and fa/+ rats					
	fa/+ rats	fa/fa rats	р		
Age (weeks)	$\textbf{8.5} \pm \textbf{0.0}$	$\textbf{8.8} \pm \textbf{0.5}$	0.59		
Body weight (g)	247 ± 10	328 ± 15	0.002		
Liver weight (g)	$\textbf{9.6} \pm \textbf{0.6}$	$\textbf{15.5} \pm \textbf{2.2}$	0.002		
Portal pressure (mmHg)	$\textbf{7.7} \pm \textbf{1.8}$	$\textbf{7.4} \pm \textbf{0.5}$	0.54		
Serum cholesterol (mmol I^{-1})	1.1 ± 0.1	$\textbf{2.2} \pm \textbf{0.2}$	0.001		
Serum triglycerides (mmol l ⁻¹)	$\textbf{0.4} \pm \textbf{0.1}$	2.3 ± 0.6	0.002		
Serum ALAT (UI I^{-1})	79 ± 10	135 ± 25	0.004		
Data are means \pm SD.					

A. Hepatic concentrations [nmol/g, measures and fits]



Fits [nmol/g] of concentrations in each compartment

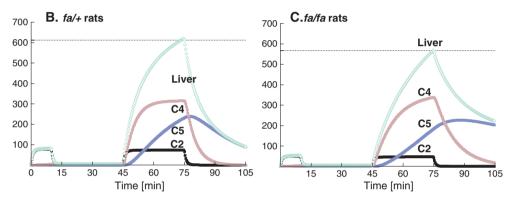


Figure 2. (A) Hepatic concentrations of contrast agents (nmol g^{-1}) during the perfusion of 200 μ m 153 Gd–DTPA (from 0 to 10 min) and 200 μ m 153 Gd–BOPTA (from 45 to 75 min). Livers were isolated and perfused from fa/+ rats (\Box and green fit, n=5) and fa/fa rats (\blacksquare and red fit, n=5). Fits obtained by pharmacokinetic modeling are shown on experimental data. (B and C) Fits of concentrations in each compartment of the liver and in the entire liver from control fa/+ rats (B) and fa/fa rats (C). Liver: total hepatic concentration; C2: extracellular compartment; C4 and C5: hepatocellular compartments.

experimental groups and a two-way ANOVA over time between both groups found no interaction (Fig. 3D, p=1.00). At T75, 153 Gd-BOPTA bile accumulation was lower in fa/fa rats while 153 Gd-BOPTA uptake was impossible to measure at this time-point, accumulation reflecting the combination of entry and exit.

3.3. 153Gd-BOPTA Elimination from Hepatocytes (T75-T105)

When ¹⁵³Gd-BOPTA perfusion was replaced by a KHB solution, ¹⁵³Gd-BOPTA concentrations within hepatocytes gradually decreased (Fig. 3E). In the absence of ¹⁵³Gd-BOPTA uptake, the elimination from hepatocytes can be measured by an initial hepatocellular elimination index (IHEI, nmol min $^{-1}$ g $^{-1}$; Fig. 3E). The IHEI was measured from T76 to T79 and was significantly decreased in fa/fa rats $(-19 \pm 5 \text{ nmol min}^{-1} \text{ g}^{-1})$ vs fa/+ rats $(-40 \pm 12 \text{ nmol min}^{-1} \text{ g}^{-1}, \ \ p < 0.008)$. Moreover, a two-way ANOVA of concentrations over time between both groups found a positive interaction (p = 0.05), the hepatic elimination of ¹⁵³Gd-BOPTA in *fa/fa* rats being significantly lower over time. Interestingly, from T95 to T105, we were able to detect the efflux from hepatocytes back to sinusoids (Fig. 3E). During this period, livers had already been perfused with a KHB solution for 20 min (total volume perfusion: 600 ml into rat livers < 16 g) and we can be sure that the radioactivity measured in outflow perfusate during these last 10 min corresponds to the efflux back of ¹⁵³Gd-BOPTA from hepatocytes to sinusoids: $64 \pm 39 \, \text{nmol}$ in fa/+ rats and

 $381\pm105\,\mathrm{nmol}$ in fa/fa rats (p<0.008). During the same period, the bile excretion was $337\pm69\,\mathrm{nmol}$ in fa/+ rats and $490\pm115\,\mathrm{nmol}$ in fa/fa rats (p=0.06). The higher amounts of $^{153}\mathrm{Gd}$ –BOPTA in perfusate and bile of fatty livers despite a decreased transport functions and/or protein expression are explained by increased residual concentrations within hepatocytes (see Discussion) (17–19). At T105, the residual hepatic $^{153}\mathrm{Gd}$ –BOPTA concentrations were $214\pm60\,\mathrm{nmol}\,\mathrm{g}^{-1}$ (fa/fa rats) and $75\pm21\,\mathrm{nmol}\,\mathrm{g}^{-1}$ (fa/+ rats p<0.008). In fa/+ rats, 13% of $^{153}\mathrm{Gd}$ –BOPTA elimination was directed to the sinusoids while 44% of $^{153}\mathrm{Gd}$ –BOPTA left hepatocytes through the sinusoidal membrane in fa/fa rats.

3.4. Pharmacokinetic Modeling of ¹⁵³Gd-BOPTA Transport in Perfused Rat Livers

To better understand the transport of ¹⁵³Gd–BOPTA in chole-static hepatocytes, we performed pharmacokinetic modeling of ¹⁵³Gd–DTPA and ¹⁵³Gd–BOPTA transport over time using the compartmental model illustrated in Fig. 1B and described in previous publications (20,21). The model successfully fits the experimental data for both *fa/fa* and *fa/+* rats (Fig. 2A) with three compartments in the liver: an extracellular compartment (C2, depicted by ¹⁵³Gd–DTPA) and two hepatocellular compartments (C4 and C5). ¹⁵³Gd–BOPTA from C2 can enter into C4 but not into C5, ¹⁵³Gd–BOPTA in C5 being provided by C4 without possible return from C5 to C4.

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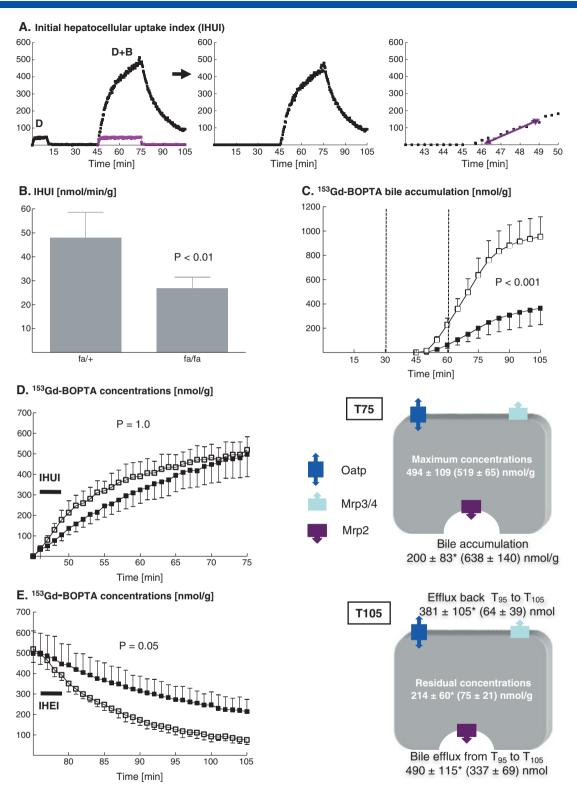


Figure 3. 153 Gd-BOPTA concentrations within hepatocytes rely on uptake and exit functions. (A) Uptake is measured by the initial hepatocellular uptake index (IHUI in nmol min $^{-1}$ g $^{-1}$). For illustration, the concentrations of 153 Gd-DTPA (D) and 153 Gd-BOPTA (B) in a single liver are shown. The extracellular distribution of 153 Gd-DTPA being similar to that of 153 Gd-BOPTA, we can obtain the exact 153 Gd-BOPTA concentrations over time within hepatocytes. Because few contrast agents are excreted into bile 5 min after the start of the perfusion, the initial slope from time 46 min (T46) to T49 measures the IHUI. B. The IHUI is shown in both experimental groups: control fa/+ and fatty fa/fa rats. (C) 153 Gd-BOPTA bile accumulation (nmol g $^{-1}$) in fa/+ (\square) and fa/fa (\blacksquare) rats. 153 Gd-BOPTA accumulation (D) in hepatocytes from T45 to T75 and 153 Gd-BOPTA elimination from hepatocytes (E) in fa/+ (\square) and fa/fa (\blacksquare) rats. At T75, the maximal concentrations within hepatocytes and bile excretion are shown. At T105, the residual concentrations within hepatocytes are shown. Efflux back (in nmol) from hepatocytes to sinusoids is compared to bile excretion (nmol) from T95 to T105. Results from fa/+ rats are between brackets. Uptake transporters are organic anion transporting polypeptides (Oatps) and exit transporters to bile are multiple resistance-associated proteins 2 (Mrp2). Transporters that return 153 Gd-BOPTA back to sinusoids are either Mrp3/4 or Oatps. * p < 0.05.

The concentrations of ¹⁵³Gd–BOPTA in C2 were lower in *fa/fa* than *fa/+* rats (Fig. 2A–C). Because the same concentration of Gd–BOPTA was perfused in both groups, only a larger extracellular distribution volume can explain the decreased hepatic concentrations measured by the gamma probe. In *fa/+* rats, the accumulation in C4 rapidly increased until a plateau is reached by T60 (Fig. 2B). When ¹⁵³Gd–BOPTA solution was switched to a KHB solution, ¹⁵³Gd–BOPTA was rapidly eliminated from C4. In C5, ¹⁵³Gd–BOPTA had a linear accumulation and its maximal accumulation remained lower than that observed in C4. Moreover, the concentrations in C5 remained high during the rinse period with KHB.

The evolution of 153 Gd-BOPTA concentrations was different in fa/fa rats (Fig. 2C). In fatty livers, the accumulation in C4 was more gradual and, from T75, the elimination was slower than in fa/+ rats. At T105, the concentrations in C5 were high and corresponded to the increased residual hepatic concentrations.

From the model, rate constants were calculated (Table 2). The hepatic uptake of 153 Gd-BOPTA through Oatps (k_{24}) was lower in fa/fa rats without reaching statistical significance. The bile excretions from C4 (k_{46}) and C5 (k_{56}) were significantly lower in fa/fa rats. Exchange from C4 to C5 (k_{45}) was unidirectional and low in both groups. Efflux from hepatocytes back to sinusoids (k_{42}) was also significantly decreased in fatty livers.

3.5. Pharmacokinetic Modeling and Mathematical Simulations

To understand why the maximal hepatic concentration (at T75) was similar in fa/fa and fa/+ rats, we performed mathematical simulations decreasing a single parameter k_{24} (entry into hepatocytes), k_{46} (bile excretion from C4), k_{56} (bile excretion from C5) or k_{42} (efflux from C4 to sinusoids), while the three others remained unchanged (Fig. 4). With these simulations, we understand how changing a single parameter interferes with hepatic concentrations. Each parameter was modified from the value obtained in fa/+ rats to 0 (10% change at each step). The values calculated in fa/fa livers were found between these extreme values. We showed that, by decreasing k_{24} , the maximal hepatic concentration at T75 decreased until the extracellular ¹⁵³Gd-BOPTA concentration was reached for $k_{24} = 0$. When k_{46} , k_{56} , or k_{42} were decreased, the maximal hepatic accumulation at T75 increased, and the increase was high with k_{42} set to 0, emphasizing the importance of efflux from hepatocytes to control the intracellular concentrations of ¹⁵³Gd-BOPTA.

To better understand the hepatic residual concentration, we also performed simulations at T105. The residual hepatic accumulation of 153 Gd–BOPTA was moderately modified by the decrease in k_{46} and k_{42} but decreasing k_{56} greatly increased the residual concentration, C5 being the only hepatocellular compartment to retain 153 Gd–BOPTA at the end of the protocol.

Finally, to understand the repartition of the contrast agent concentrations in C2, C4, and C5 over time in cholestatic livers, we modified each parameter while the three other parameters remain similar to that calculated in fa/+ rats (Fig. 5). When k_{24} was set at 0.768 (value close to that of fa/fa rats), while k_{46} , k_{56} , and k_{42} remained similar to fa/+ rats, the hepatic concentration decreased. Modifications of k_{46} , k_{56} or k_{42} increased the hepatic concentration. Interestingly, setting k_{56} at 0.013 increased 153 Gd-BOPTA concentrations in C5 over those observed in C4.

4. DISCUSSION

4.1. Relevance of Gd-BOPTA Concentrations within Hepatocytes

The transport of the organic anion Gd–BOPTA across hepatocytes relies on the expression and function of proteins located on the sinusoidal and canalicular membranes. It is relatively easy to measure a single transport parameter in *in vivo* or *in vitro* experimental studies but it is less frequent to measure simultaneously all the parameters (hepatocyte uptake, bile excretion, efflux back to perfusate) as well as the consequences of all parameters on the intracellular concentrations. The hepatic concentrations of Gd–BOPTA depend on the balance between uptake via sinusoidal transporters, biliary excretion and efflux back to sinusoids (either through bidirectional sinusoidal transporters or through sinusoidal Mrp transporters; Fig. 3).

In the experimental model of isolated perfused livers, we evaluate the extracellular space (C2) by measuring Gd–DTPA concentrations. The exact concentrations of Gd–BOPTA within hepatocytes are available (C4 and C5) and the IHUI measures the initial uptake function before the start of bile excretion. Bile excretion of Gd–BOPTA is available during the entire protocol while the efflux back from hepatocytes to sinusoids is evidenced at the end of the study (Fig. 3). During the protocol, the hepatic perfusate flow is constant in all experiments and set to 30 ml min⁻¹, thus the decreased IHUI of fatty livers is not related to changes of hepatic perfusion. In our experimental model, liver isolation precludes interferences with extrahepatic organs. The composition of solutions is well controlled and does not include binding

Table 2. Hepatic pharmacokinetics by compartmental analysis				
Rate constants (min ⁻¹)	fa/+ rats	fa/fa rats	р	
k ₁₂	1.19 ± 0.13	1.33 ± 0.36	0.42	
k ₂₃	9.60 ± 4.52	9.27 ± 1.44	1.00	
k ₂₄	0.96 ± 0.39	0.79 ± 0.17	0.55	
k ₄₂	$\textbf{0.11} \pm \textbf{0.02}$	0.05 ± 0.01	0.008	
k ₄₅	$\textbf{0.05} \pm \textbf{0.02}$	0.03 ± 0.01	0.10	
k ₄₆	0.065 ± 0.011	0.025 ± 0.007	0.008	
k ₅₆	0.045 ± 0.008	0.013 ± 0.006	0.008	

The rate constant k_{24} is associated with 153 Gd-BOPTA uptake. k_{46} and k_{56} describe bile excretion in C6 from the hepatic compartments C4 and C5. k_{42} is associated with the efflux of 153 Gd-BOPTA from hepatocytes back to sinusoids.

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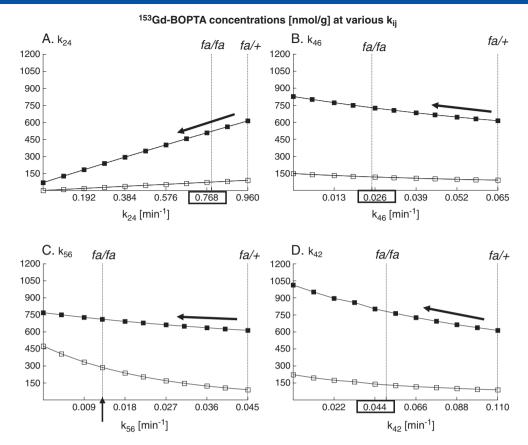


Figure 4. Mathematical simulations of 153 Gd-BOPTA concentrations (nmol g $^{-1}$) at various k_{ij} according to the pharmacokinetic model presented in Fig. 1B A single parameter representing 153 Gd-BOPTA entry into hepatocytes (A, k_{24}), bile excretion from both hepatocellular compartments (B, k_{46} and C, k_{56}), and efflux back from hepatocytes to the extracellular space (D, k_{42}), was modified from the value obtained in control fa/+ rats to 0 (10% change at each step on the x-axis). The consequences of changing a single parameter on 153 Gd-BOPTA hepatic concentrations (nmol g $^{-1}$) were calculated. Hepatic accumulations were calculated at T75 (\blacksquare) and T105 (\square). Decreasing k_{24} (153 Gd-BOPTA uptake into hepatocytes) lowers hepatic concentrations. Decreasing k_{42} (efflux through the sinusoidal membrane), k_{46} or k_{56} (efflux through the canalicular membrane) augments them.

proteins. Because the transport of Gd–BOPTA through transporters is related to the concentration gradients of the contrast agent between sinusoids, hepatocytes and bile, we administered Gd–BOPTA at steady concentrations rather than as a bolus injection similarly to the clinical situation.

Understanding the function of transporters is important to evidence how Gd–BOPTA accumulates in hepatocytes. Cholestatic fatty livers from *fa/fa* rats have a decreased expression of Oatp1a4 (17,18) and Mrp2 (17,19). Interestingly, Mrp3 and Mrp4 are increased in another experimental model of fatty livers (6). In addition to the expression of transporters, other function regulations were described. Phosphorylation of Mrp2 can retrieve the transporter from the canalicular membrane and trap Gd–BOPTA within rat hepatocytes (14,22,23). Entry of Gd–BOPTA into hepatocytes is also impaired by drug–drug interaction when Gd–BOPTA is perfused together with bromosulfophthalein (24).

Thus, it is important to assess the function of transporters in injured hepatocytes. We show that fatty hepatocytes have a decreased uptake function as measured by the decreased uptake index (IHUI). The elimination function of Gd–BOPTA from hepatocytes is also impaired, as shown by the decrease in the IHEI, k_{46} , and k_{56} . Gd–BOPTA may also exit from hepatocytes by efflux back to sinusoids and this function is also impaired in fatty livers (decreased k_{42}). However, the amounts of Gd–BOPTA that exit from hepatocytes during the period T95–T105 are higher in fatty than control livers because the residual concentrations

are higher during this period. The amounts (A) of Gd-BOPTA transported through the sinusoidal membrane back to hepatic vessels are defined by the equation: A (nmol min⁻¹ g⁻¹) = k_{42} (min⁻¹) × C (nmol g⁻¹), where C is the concentration in hepatocytes. Thus, Gd-BOPTA amount transported back to sinusoids in fatty hepatocytes can be higher that in normal hepatocytes even though the transport function through Oatps is decreased. We previously showed, that the efflux back to sinusoids and bile excretion in normal hepatocytes are proportional to the concentrations inside livers (13). This point emphasizes the importance of concentrations within hepatocytes in the regulation of transport proteins. Of note, we do not distinguish the mechanisms that induce a decreased transport of Gd-BOPTA through membrane proteins in fatty hepatocytes: either decreased expression of transporters and/or dysregulation of protein function.

4.2. Hepatobiliary Contrast Agents and Liver Imaging

The clinical importance of our findings is provided by recent studies that investigated the hepatic accumulation of Gd–BOPTA and the other closely related compound Gd–EOB–DTPA (Primovist[®], Bayer, Germany). In multinodular focal fatty infiltrations, Marin *et al.* (10) found a decreased signal intensities during the vascular and equilibrium phases following the injection of Gd–BOPTA, a finding similar to the decreased concentrations of the contrast agent in C2 from *falfa* livers. However, in contrast to our results,

Figure 5. Concentrations of 153 Gd-DTPA and 153 Gd-BOPTA (nmol g $^{-1}$): effect of changing a single parameter. The pharmacokinetics over time of 153 Gd-BOPTA concentrations in fa/+ rats is shown in the entire liver (green), compartment 2 (C2 or extracellular compartment, black), C4 (hepatocytes, brown), and C5 (hepatocytes, purple) (A). Pharmacokinetics was calculated in fa/+ rats with k_{24} set to 0.768 (B), fa/+ rats with k_{46} = 0.026 (C), fa/+ rats with k_{24} = 0.013 (D), and fa/+ rats with k_{42} = 0.044 (E). These figures illustrate how changing a single k_{ij} (from the value measured in fa/+ rats to that measured in fa/fa rats) modifies the distribution of 153 Gd-BOPTA in the entire liver, C3, and C4. The black horizontal line shows the maximal concentration obtained in fa/+ livers and the green horizontal one, the maximal concentration obtained with a single k_{ij} change.

Gd–BOPTA did not enter human fatty hepatocytes and the lesions were hypointense in comparison to surrounding normal parenchyma. In rats with nonalcoholic steatohepatitis, the maximal intensity following a bolus of Gd–EOB–DTPA was similar to normal rats while the elimination was delayed (25). In this experimental model, the expression of Oatps and Mrp2 was similar in normal and fatty livers. However, the function of the transporters could not be determined.

4.3. Pharmacokinetic Modeling of Gd-BOPTA Transport

In previous studies, we described the pharmacokinetics of Gd–DTPA and Gd–BOPTA transport by compartmental analysis in normal rats (21,26). Such modeling is a valuable tool to understand the kinetic of drug concentrations in various compartments over time. The best model includes six compartments and eight rate

constants. We show that this model perfectly fits the experimental data in both normal and fatty livers. The extracellular distribution volume is higher in fatty liver, and the evolution of Gd-BOPTA accumulation is different in both hepatocellular compartments C4 and C5. In C4 of normal livers, the initial accumulation rate is high until a plateau is reached. Gd-BOPTA is rapidly excreted from this compartment during the rinse period. Accumulation in C5 is more linear and late to disappear. In fatty livers, the accumulation in C4 is more gradual without reaching a plateau and the emptying is slower. Chandra et al. (27) also found two hepatocellular compartments in isolated rat livers perfused with 5-(and 6)carboxy-2',7' dichlorofluorescein, which uses the same Oatps/ Mrp2 pathway as Gd-BOPTA. The signification of two cellular compartments is puzzling but we can speculate that Gd-BOPTA accumulation in C4 is close to the canalicular membrane because its hepatic elimination is rapid.

4.4. Mathematical Simulations

Pharmacokinetic modeling with mathematical simulations has already been performed to better understand the hepatobiliary transport of drugs through the Oatps/Mrp2 pathway. Thus, in healthy volunteers, a pharmacokinetic model was developed based on blood, urine, and bile concentration-time profiles after the injection of ^{99m}Tc-mebrofenin (28). In this model, mathematical simulations were also performed to evaluate how decreasing the transport through OATPs, MRP2, or both OATPs and MRP2, would influence the hepatic accumulation (28). Interestingly, liver imaging with ^{99m}Tc-mebrofenin also represents the accumulation of tracers (29,30).

Our simulations show that hepatic concentrations over time are the result of four biological processes: hepatocyte uptake through Oatps (k_{24}) , bile excretion through Mrp2 (two parameters: k_{46} and k_{56}) and efflux back to sinusoids (k_{42}) . The most important parameters are those located on the sinusoidal membrane, a finding in line with the evidence that a compound that does not enter into hepatocytes will not be excreted in bile. Finally, to understand the repartition of the contrast agent over time in C2, C4 and C5 in cholestatic livers, we modified each parameter while the three other parameters remain similar to that calculated in normal livers rats. When k_{24} was set at 0.768 (value close to that of fatty livers), while k_{46} , k_{56} and k_{42} remained similar to normal livers, the hepatic accumulation decreased. Decreasing k_{46} , k_{56} or k_{42} increased the hepatic accumulation.

In summary, we show that cholestatic hepatocytes have similar maximal concentrations of Gd–BOPTA than normal cells despite a decreased function of Oatps and Mrp2. The study emphasizes the importance of concentrations of organic anions within hepatocytes for the transport function through membrane proteins. Although complex, such understanding is important to analyze liver imaging with hepatobiliary contrast agents in cholestatic fatty livers. Moreover, we describe an index of hepatocellular uptake function (IHUI) that may score the severity of fatty livers in clinical liver MRI associated with the injection of hepatobiliary contrast agents.

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