

# Induction of Influx and Efflux Transporters and Cytochrome P450 3A4 in Primary Human Hepatocytes by Rifampin, Rifabutin, and Rifapentine

Beth Williamson,<sup>a</sup> Kelly E. Dooley,<sup>b</sup> Yuan Zhang,<sup>a</sup> David J. Back,<sup>a</sup> Andrew Owen<sup>a</sup>

Department of Molecular and Clinical Pharmacology, The University of Liverpool, Liverpool, United Kingdom<sup>a</sup>; Department of Medicine, Divisions of Clinical Pharmacology and Infectious Diseases, Johns Hopkins University School of Medicine, Baltimore, Maryland, USA<sup>b</sup>

**Rifampin is a potent inducer of cytochrome P450 (CYP) enzymes and transporters. Drug-drug interactions during tuberculosis treatment are common. Induction by rifapentine and rifabutin is understudied. Rifampin and rifabutin significantly induced CYP3A4 (80-fold and 20-fold, respectively) in primary human hepatocytes. The induction was concentration dependent. Rifapentine induced CYP3A4 in hepatocytes from 3 of 6 donors. Data were also generated for ABCB1, ABCC1, ABCC2, organic anion-transporting polypeptide 1B1 (OATP1B1), and OATP1B3. This work serves as a basis for further study of the extent to which rifamycins induce key metabolism and transporter genes.**

Tuberculosis is a major global health problem (1). Effective short-course therapy lasts for 6 months but requires rifampin, and clinically significant drug-drug interactions are common due to induction of cytochrome P450 3A4 (CYP3A4) and key drug transporters included herein (1–3). Hence, antiretroviral coadministration with tuberculosis treatment is particularly challenging. Besides CYP3A4, drug transporters can significantly alter the absorption and distribution of drugs. Many compounds used in human immunodeficiency virus treatment, particularly the protease inhibitors and nucleoside reverse transcriptase inhibitors, are transported by proteins such as ABCB1 (3), ABCC2 (4), organic anion-transporting polypeptide 1B1 (OATP1B1), and OATP1B3 (5, 6). The genes encoding these proteins are influential in the safety, efficacy, and disposition of many drugs. For example, the induction of *ABCB1* by rifampin decreases the area under the curve (AUC) of efavirenz by 22% (2). Rifabutin is considered a less-potent inducer and is often used in place of rifampin for patients receiving antiretroviral drugs for human immunodeficiency virus to reduce the risk of drug interactions (7–9). The substitution of rifapentine for rifampin may reduce the treatment duration required for cure, but the induction potential of rifapentine is comparatively understudied (2, 10). The sterilizing activity of rifapentine is dose dependent in an established mouse model of tuberculosis, with eradication possible in 3 months or less when high-dose rifapentine is substituted for rifampin in a multidrug treatment regimen (11, 12). However, dose increases resulted in less-than-dose-proportional increases in rifapentine exposures (10, 13). In addition, the mean area under the concentration-time curve of oral midazolam, a CYP3A4 probe, decreased by 75% when coadministered with rifampin, compared to 92% when coadministered with rifapentine, each given at 10 mg/kg of body weight daily (14). We evaluated the *in vitro* induction of CYP3A4 and transporters by rifampin, rifabutin, and rifapentine in primary human hepatocyte samples from six donors. Other studies have previously investigated the induction of CYP activity by rifampin, rifabutin, and rifapentine (9) and the mRNA expression of drug transporters induced by rifampin (3, 6), but no studies have comprehensively compared the mRNA induction of CYPs

and transporters in primary human hepatocytes with all 3 compounds in parallel.

Cryopreserved hepatocyte recovery medium (CHRM medium), Williams' E medium, plating and supplement medium, cryopreserved human hepatocytes, plating cocktail, maintenance cocktail, gene expression assays, and 96-well collagen-coated plates were purchased from Life Technologies (Paisley, Scotland, United Kingdom). All other chemicals were purchased from Sigma-Aldrich (Poole, Dorset, United Kingdom), unless otherwise indicated.

Rifampin, rifabutin, and rifapentine were prepared as 10 mM stocks in methanol and further diluted in hepatocyte medium to the required concentrations. Cryopreserved human hepatocytes (from 6 human donors) were thawed according to the manufacturer's instructions (15) and resuspended in Williams' E medium supplemented with plating cocktail (1  $\mu$ M dexamethasone, a 1% solution of penicillin-streptomycin, 4  $\mu$ g/ml insulin, 5% fetal bovine serum, 2 mM GlutaMAX, and 15 mM HEPES [Life Technologies, Paisley, United Kingdom]). Cell numbers and viability were assessed using trypan blue exclusion. Cells were seeded in 24-well plates precoated with collagen at a density of  $2 \times 10^5$  cells per well and were incubated for 12 h at 37°C with 5% CO<sub>2</sub> and 95% humidity. The medium was replaced with Williams' E medium supplemented with maintenance cocktail (0.1  $\mu$ M dexamethasone, a 0.5% solution of penicillin-streptomycin, 2 mM GlutaMAX, 15 mM HEPES, 6.25  $\mu$ g/ml human recombinant insulin, 6  $\mu$ g/ml human transferrin, 6  $\mu$ g/ml selenous acid, 1.25  $\mu$ g/ml bovine serum albumin, and 5.35  $\mu$ g/ml linoleic acid [Life Technologies, Paisley, United Kingdom]). Hepatocytes were incubated with rifampin, rifabutin, or rifapentine at concentrations spanning the

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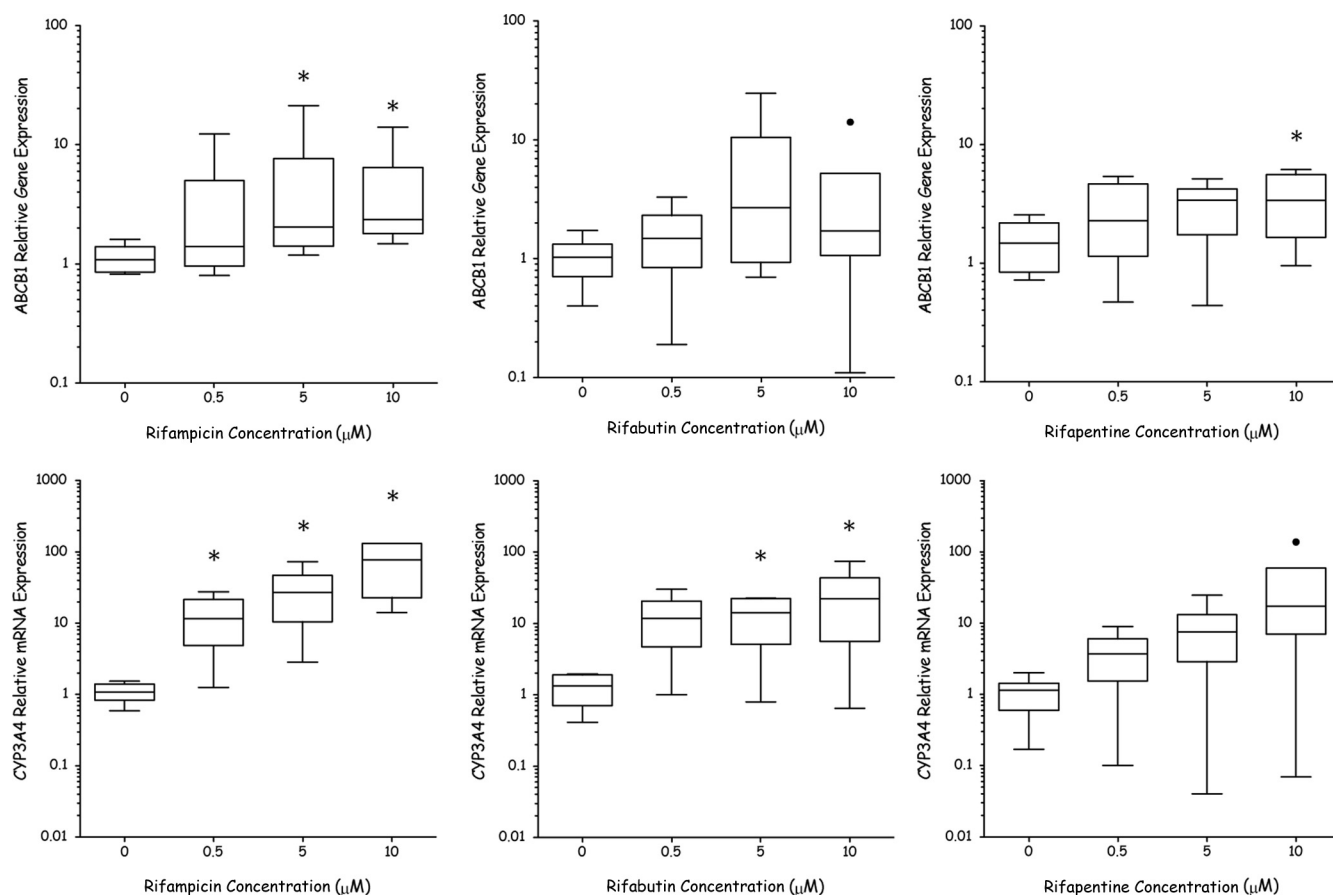
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Address correspondence to Andrew Owen, aowen@liverpool.ac.uk

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**FIG 1** Relative gene expression of cytochrome P450 isoenzyme 3A4 and ABCB1 in primary hepatocytes when incubated with rifampin (RIF), rifabutin (RBT), or rifapentine (RPT) at 0, 0.5, 5, and 10  $\mu\text{M}$ . Data were normalized to results for *GAPDH* housekeeping gene and for control primary hepatocytes (0  $\mu\text{M}$ ) using the comparative  $C_T$  method ( $C_T = 2^{-\Delta\Delta C_T}$ ). Tukey box plot represents the means and interquartile ranges (IQR) ( $n = 6$  donors completed in triplicate), with boxes showing IQR and whiskers representing  $<1.5 \times \text{IQR}$ . Data outside  $1.5 \times \text{IQR}$  are labeled as outliers ( $\bullet$ ). An asterisk indicates a significant difference from the control (0  $\mu\text{M}$ ) by paired  $t$  test or Wilcoxon signed-rank test ( $P < 0.05$ ).

therapeutic range (0.5, 5, and 10  $\mu\text{M}$ ) for 24 h; for rifampin, these concentrations have previously been shown to cause no toxicity (3). Control cells (0  $\mu\text{M}$ ) contained the same volume of methanol and hepatocyte medium. RNA was extracted using TRIzol reagent and reverse transcribed using the standard methodology recommended by Life Technologies. Gene expression analysis was conducted for *OATP1B1*, *OATP1B3*, *ABCB1*, *ABCC1*, *ABCC2*, and *CYP3A4* by real-time PCR. Gene expression was normalized to that of the housekeeping gene, glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*), and compared to that of the control (0  $\mu\text{M}$ ) using the comparative  $C_T$  method ( $C_T = 2^{-\Delta\Delta C_T}$ ;  $C_T$ , cycle number at which the fluorescence in the reaction crosses the preset arbitrary threshold;  $\Delta C_T$ , difference between the  $C_T$  target and reference;  $\Delta\Delta C_T$ , difference between the  $\Delta C_T$  of the test and the  $\Delta C_T$  of the preassigned control). The normality of the data was assessed using a Shapiro-Wilk test, and statistical analysis conducted using the paired  $t$  test or Wilcoxon signed-rank test for normally or nonnormally distributed data, respectively.

The effects of rifampin, rifabutin, and rifapentine on *ABCB1* and *CYP3A4* mRNA are shown in Fig. 1. The fold changes in gene expression for *OATP1B1*, *OATP1B3*, *ABCC1*, and *ABCC2* when treated with rifampin, rifabutin, and rifapentine are shown in Table 1. Rifampin elicited significant upregulation of *ABCB1* from

5  $\mu\text{M}$  and of *CYP3A4* from 0.5  $\mu\text{M}$ . Concentration-dependent induction was observed for both genes, with the greatest induction observed at 10  $\mu\text{M}$  (80-fold [ $P = 0.03$ ] and 5-fold [ $P = 0.03$ ] for *CYP3A4* and *ABCB1*, respectively). Rifampin significantly upregulated *OATP1B1* (2-fold [ $P = 0.03$ ]) and *ABCC2* (3-fold [ $P = 0.03$ ]) at 10  $\mu\text{M}$  (similar to the results of Haenisch et al., 2011). Rifabutin elicited a 20-fold upregulation in *CYP3A4* gene expression ( $P = 0.05$ ) and a 4-fold upregulation of *OATP1B3* ( $P = 0.04$ ) at 5  $\mu\text{M}$ .

When analyzed as the average of the results for the hepatocytes from the 6 donors, rifapentine did not significantly induce the expression of *CYP3A4*, but significant induction was observed in hepatocytes from 3 of 6 donors when analyzed individually. *ABCB1* was the only gene significantly induced by rifapentine (4-fold [ $P = 0.04$ ]) at 10  $\mu\text{M}$ . The results herein suggest the hierarchy of the rifamycins' potency as *CYP3A4* inducers to be rifampin > rifabutin > rifapentine, while previous studies found that rifapentine was more potent than rifabutin (8). However, both studies agree that rifampin is the most-potent inducer of *CYP3A4*.

Consistent with all primary hepatocyte studies, great interdonor variability was observed (16). However, concentration-dependent responses were seen for most genes (including *CYP3A4*). This work highlights the extent to which rifampin induces

**TABLE 1** Fold change in gene expression of hepatic influx and efflux transporters when hepatocytes were incubated with rifampin, rifabutin, or rifapentine at various concentrations

Compound	Concn ( $\mu\text{M}$ )	Mean (range) fold change in mRNA expression compared to result for 0 $\mu\text{M}$ treatment <sup>a</sup>			
		OATP1B1	OATP1B3	ABCC1	ABCC2
Rifampin	0.5	5.19 (0.30–24.31)	7.92 (0.73–29.79)	2.15 (1.04–4.46)	0.93 (0.28–1.44)
	5.0	1.51 (0.36–3.95)	5.54 (1.02–19.89)	1.53 (0.13–8.79)	1.06 (0.77–1.83)
	10.0	1.91 (0.55–4.77)*	1.09 (0.41–2.10)	1.78 (0.17–3.70)	2.39 (0.49–7.31)*
Rifabutin	0.5	0.63 (0.11–1.69)	2.00 (0.25–5.99)	0.78 (0.35–1.91)	1.19 (0.25–4.26)
	5.0	0.94 (0.13–2.95)	3.58 (1.11–9.15)*	2.69 (0.14–8.06)	1.95 (0.22–12.01)
	10.0	0.81 (0.38–2.05)	3.06 (0.65–5.40)	1.45 (0.34–2.53)	1.89 (0.17–7.08)
Rifapentine	0.5	1.40 (0.52–3.08)	1.46 (0.08–5.29)	1.40 (0.20–4.25)	1.19 (0.72–2.50)
	5.0	0.86 (0.41–1.48)	1.07 (0.18–4.46)	1.24 (0.10–2.01)	1.52 (1.00–5.48)
	10.0	1.37 (0.32–3.12)	1.30 (0.11–6.60)	2.23 (0.10–5.08)	1.95 (0.65–3.93)

<sup>a</sup> Change in relative gene expression of hepatic influx and efflux transporters when incubated with rifampin, rifabutin, or rifapentine at 0.5, 5, or 10  $\mu\text{M}$  compared to the expression in control primary hepatocytes with no drug added. An asterisk indicates a significant difference from the control (0  $\mu\text{M}$ ) by paired *t* test or Wilcoxon signed-rank test ( $P < 0.05$ ). OATP, organic anion-transporting polypeptide; ABC, ATP-binding cassette transporter.

CYP3A4-mediated metabolism and the transport of compounds compared to the induction by rifapentine, a potential alternative. Larger increases in CYP3A4 mRNA expression were observed when hepatocytes were treated with rifampin than with rifabutin and rifapentine at the same micromolar concentration. In contrast to the data herein, a study with healthy volunteers found that the AUC of midazolam was decreased 17% more when coadministered with rifapentine than with rifampin (14). However, at standard daily doses, the average rifampin, rifabutin, and rifapentine concentrations are approximately 2.3  $\mu\text{M}$ , 0.3  $\mu\text{M}$ , and 15.7  $\mu\text{M}$ , respectively (8, 17, 18). Of particular note, the plasma concentrations of rifabutin in patients are comparatively low, and this may help rationalize the limited induction by rifabutin seen clinically. Thus, there is a spectrum of levels of induction by rifampin, rifabutin, and rifapentine, but the data should be interpreted in the context of differences in plasma concentrations seen clinically. The data should also be interpreted in the context that concentrations in hepatocytes and/or gut may exceed those found in the plasma of patients. The unbound plasma concentration is often used to estimate clinical drug interactions. However, protein binding can be dependent on health status. For example, rifampin is 87 to 91% bound in healthy individuals but 84 to 88% bound in tuberculosis-infected individuals (19). Nonetheless, it should be noted that several previous investigations have shown that effects at the mRNA level are not always translated to activity (3, 7, 20).

An ATP binding cassette transporter (ABC), ABCB1, is a transmembrane efflux protein responsible for the removal of a broad range of bile acids, lipids, and xenobiotics from hepatocytes (21). Rifampin and rifapentine significantly induced ABCB1; hence, an enhanced clearance of coadministered substrates may be predicted. Also, rifampin, rifapentine, ethambutol, and isoniazid are all substrates of ABCB1 (7, 22, 23), suggesting a potential role in autoinduction and potential effects between drugs within a *Mycobacterium tuberculosis* treatment regimen. A significant upregulation of OATP1B1 mRNA was also observed with 10  $\mu\text{M}$  rifampin. Given that rifampin is in itself a substrate for OATP1B1 (24) and that polymorphisms within the *SLCO1B1* gene affect rifampin pharmacokinetics (25), these data indicate an involvement of OATP1B1 in the reported rifampin autoinduction (26).

Current clinical trials are investigating high-dose daily rifapen-

tine (Tuberculosis Trials Consortium Study 29X [27]) and high-dose rifampin (PanACEA Consortium [28]) as potential regimens to shorten tuberculosis treatment. *In vitro* results suggest that increasing the concentration of rifapentine may lead to clinically relevant drug-drug interactions mediated through ABCB1. Boeree et al. (28) found that 35 mg/kg of rifampin daily was safe and well tolerated over 14 days and that early bactericidal activity increased with increasing dose, with no apparent plateau. Trials are now being planned to assess the activity of high-dose rifampin over 8 weeks. Our data suggest that concentration-dependent induction should be considered when interpreting the results of ongoing trials of higher-dose rifampin and rifapentine, since it cannot be assumed that maximum autoinduction is achieved at standard doses. With the absence of an apparent ceiling as drug exposure increases (28), rifampin doses above those used in the clinic today may lead to significant and highly variable drug-drug interactions, which is of considerable clinical concern.

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