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GW4064, an agonist of farnesoid X receptor (FXR), represses CYP3A4 expression in human hepatocytes by inducing small heterodimer partner (SHP) expression

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Abbreviations

CAR, constitutive androstane receptor; FXR, farnesoid X receptor; GR, glucocorticoid receptor; GW4064, 3-(2,6-Dichlorophenyl)-4-(3'-carboxy-2-chlorostilben-4-yl)oxymethyl-5-isopropylisoxazole; HH, human hepatocytes; LXR, liver X receptor; PPAR α , peroxisome proliferator-activated receptor α ; PXR, pregnane X receptor; SHP, small heterodimer partner

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ABSTRACT

Farnesoid X receptor (FXR) functions as a regulator of bile acid and lipid homeostasis, and is recognized as a promising therapeutic target for metabolic diseases. The biological function of FXR is in part mediated by small heterodimer partner (SHP); ligand-activated FXR enhances SHP expression, and SHP in turn represses activity of multiple transcription factors. The objective of this study is to investigate the effect of FXR activation on the expression of the major drug-metabolizing enzyme, cytochrome P450 (CYP) 3A4. The effects of GW4064, a synthetic agonist of FXR, on expression and activity of CYP3A4 were examined in primary human hepatocytes by using quantitative real-time PCR and S9 phenotyping. In human hepatocytes, treatment of GW4064 (1 μ M) for 48 hours resulted in 75% decrease in CYP3A4 mRNA expression and 25% decrease in CYP3A4 activity, accompanied by ~3-fold increase in SHP mRNA expression. In HepG2 cells, SHP repressed transactivation of CYP3A4 promoter by pregnane X receptor (PXR), constitutive androstane receptor (CAR), and glucocorticoid receptor (GR). Interestingly, GW4064 did not repress expression of CYP2B6, another target gene of PXR and CAR; GW4064 was shown to enhance CYP2B6 promoter activity. In conclusion, GW4064 represses CYP3A4 expression in human hepatocytes, potentially through upregulation of SHP expression and subsequent repression of CYP3A4 promoter activity. Clinically significant drug-drug interaction involving FXR agonists and CYP3A4 substrates may occur.

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Introduction

Farnesoid X receptor (FXR, NRIH4) is a ligand-activated nuclear receptor and a member of the steroid/thyroid hormone receptors. FXR is highly expressed in liver, intestine, kidney and adrenal gland (Forman et al., 1995; Parks et al., 1999; Wang et al., 1999). Endogenous bile acids are ligands for FXR, and binding of bile acids to FXR leads to regulation of genes involved in bile acid homeostasis and glucose metabolism (Wang et al., 1999; Eloranta and Kullak-Ublick, 2008). Based on the biological function of FXR as a regulator of lipid and glucose metabolism, many FXR agonists are currently being investigated as potential therapeutic agents for the treatment of metabolic disorders including hypercholesterolemia (Thomas et al., 2008; Fiorucci et al., 2012). FXR is known to cause both transactivation and transrepression of its target gene promoters (Hollman et al., 2012). For example, ligand-activated FXR transrepresses the promoter of a drug-metabolizing enzyme UGT2B7 in Caco-2 cells (Lu et al., 2005) while FXR transactivates the promoter of small heterodimer partner (SHP) by direct binding to -333/-320 of *SHP* (Goodwin et al., 2000). SHP is a nuclear receptor that lacks a DNA-binding domain, and it represses expression of genes involved in bile acid synthesis by interfering with action of other transcriptional factors (Lee et al., 2000; Wang et al., 2002).

Cytochrome P450s are the major drug-metabolizing enzymes responsible for the oxidative biotransformation of drugs, and CYP3A4 is the most prevalent among all CYP enzymes. Expression of P450 enzymes is highly inducible by endo- and xeno-biotics, mainly via actions of ligand-activated transcription factors pregnane X receptor (PXR) and constitutive androstane receptor (CAR) (Gao and Xie, 2010). PXR and CAR, upon binding to respective ligands, transactivate CYP3A4 and CYP2B6 promoters and increase their expression (Zhou, 2008). This subsequently leads to clinically significant drug-drug interactions such that the doses of CYP3A4 substrate drugs need to be adjusted to prevent a lack of drug efficacy. Additionally, corticosteroids (at physiological concentrations) enhance PXR transactivation of CYP3A4 promoter by increasing the expression of PXR and its binding partner retinoid X receptor (RXR) as well as the transcriptional activity of PXR, via glucocorticoid receptor (GR) (Pascussi et al., 2001).

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Previously, FXR was shown to transactivate CYP3A4 promoter in HepG2 cells by binding to response elements located in the distal regulatory region of *CYP3A4* (Gnerre et al., 2004). Also, in mice, GW4064 (a synthetic agonist of FXR) was shown to enhance expression of *Cyp3a11*, the murine homolog of human CYP3A4 (Gnerre et al., 2004). Together, these results suggest that FXR activators that are currently under development will likely increase CYP3A4-mediated drug metabolism in humans. However, extrapolation of the results to humans appears difficult due to minimal expression of key transcription factors involved in P450 regulation (e.g., PXR or CAR) in HepG2 cells (Hart et al., 2010; Guo et al., 2011) and well-known interspecies differences in the regulation of hepatic drug-metabolizing enzymes between humans and rodents (Martignoni et al., 2006). Taken together, the effects of FXR activation on CYP3A4 expression and the underlying regulatory mechanisms remain unclear.

In this study, we investigated the effect of FXR activation on CYP3A4 expression, using primary human hepatocytes. Our results show that GW4064 represses CYP3A4 expression, potentially through upregulating SHP expression and subsequent repression of PXR and CAR transactivation of CYP3A4 promoter.

Materials and Methods

Reagents. GW4064, midazolam, 1-hydroxymidazolam, cholic acid, rifampin and dexamethasone were purchased from Sigma-Aldrich (St. Louis, MO).

Plasmids. A luciferase vector harboring distal and proximal xenobiotic response elements of *CYP3A4* (i.e., pGL3-CYP3A4) was obtained from Dr. Hongbing Wang (Faucette et al., 2006). The luciferase vector containing 1.8-kb of CYP2B6 upstream regulatory region (i.e., pGL3-CYP2B6) and the luciferase vector containing 2.2-kb of SHP upstream regulatory region (i.e., pGL2-SHP) (Kim et al., 2003) were kindly provided by Drs. Masahiko Negishi (NIEHS) and Hueng-Sik Choi (Chonnam National University), respectively. Expression vectors for SHP (Koh et al., 2014), PXR, or CAR (Koh et al., 2012) were previously described. GR and FXR expression vectors were kindly provided by Drs. Alan McLachlan

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(University of Illinois at Chicago), and Yoon-Kwang Lee (Northeast Ohio Medical University), respectively.

Cell Culture. HepG2 cells from the American Type Culture Collection (ATCC, Manassas, VA) were cultured in complete DMEM supplemented with 10% fetal bovine serum (Gemini, Sacramento, CA), 2 mM L-glutamine, 100 U/mL penicillin, 100 µg/mL streptomycin, and 1% MEM nonessential amino acids. HEK293T cells were cultured in RPMI-1640 media supplemented with 10% fetal bovine serum (Gemini, West Sacramento, CA), 10 mM HEPES, 100 µM non-essential amino acids, 100 U/mL penicillin, and 100 µg/mL streptomycin (Life Technologies, Carlsbad, CA).

Primary Hepatocyte Culture. Freshly isolated human hepatocytes were obtained from Liver Tissue Cell Distribution System (Pittsburgh, PA). Briefly, human hepatocytes were shipped overnight in cold preservation media. Upon receipt, the media were replaced with serum-free Williams' E media (without phenol red) containing 0.1 µM dexamethasone, 100 U/mL penicillin, 100 µg/mL streptomycin, 15 mM HEPES, 2 mM L-glutamine, 5.5 µg/mL transferrin, and 5 ng/mL sodium selenite. Cells were allowed to recover from shipping for 18 hr at 37 °C in an atmosphere containing 5% CO₂, and used for experiments on the next day.

Quantitative Real-Time (qRT)-PCR. Total RNA was isolated from human hepatocytes using Trizol (Life Technologies) and used as template for the synthesis of complementary DNA using High Capacity cDNA Archive Kit (Applied Biosystems, Foster City, CA). With the cDNA as template, qRT-PCR was performed using SYBR Green reagents and the primers listed in supplemental Table S1. The fold change in mRNA levels was determined after normalizing the gene expression levels to those of GAPDH ($2^{-\Delta\Delta Ct}$ method).

Luciferase Reporter Assays. HepG2 and HEK293T cells were seeded in 24-well plates at a density of 5×10^5 or 1×10^5 cells per well, respectively. On the next day, the cells were transfected with 0.3 µg of luciferase construct, 0.1 µg of expression plasmid (or empty vector as a control) for a transcription factor, and 0.002 µg of *Renilla* expression plasmid, using Fugene HD transfection reagent (Roche Applied

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Sciences) according to the manufacturer's protocol. At 48 hr post-transfection, the transfected cells were harvested for determination of luciferase activities using Dual-Luciferase[®] Reporter Assay System (Promega, Madison, WI). Luciferase activity was normalized to the *Renilla* luciferase activity. At least two independent experiments were performed in triplicate.

Determination of CYP3A4 activity. S9 fractions were prepared as described previously (Felmlee et al., 2008). Midazolam (final concentration 3 μ M) was mixed with S9 fractions (10 μ g) in NADPH-regenerating system (5 mM isocitric acid, 0.2 unit/mL isocitric acid dehydrogenase, and 5 mM magnesium chloride in 100 mM Tris-HCl buffer, pH 7.4; 100 μ L total volume). The reactions were started by addition of NADP⁺ (10 mM) and further incubated for 1 hr. The reactions were quenched by adding cold acetonitrile (120 μ L) that contains phenytoin (0.5 μ M) as internal standard. The mixture was kept on ice for 30 min, followed by centrifugation at 16,100g for 15 min at 4 °C to obtain the supernatant. The concentrations of 1-hydroxymidazolam in the supernatants were determined by LC-MS/MS (Agilent 1200 HPLC interfaced with Agilent 6410 Triple Quadrupole tandem MS equipped with an electrospray ion source. Chromatographic separation was carried out by using a Waters XTerra MS C18 column (2.1 \times 50 mm, 3.5 μ m; Waters Corporation, Milford, MA). Mobile phase was delivered at 250 μ L/min, and the gradient was initiated at 90% A and 10% B [A, 0.1% (v/v) formic acid in water; B, acetonitrile]. The proportion of mobile phase B was increased to 90% over 1 min, held constant for 2 min, and then restored to the initial composition. The injection volume was 10 μ L. Midazolam was detected by MS/MS (341.9/324.0), and phenytoin was used as the internal standard (253.2/182.2) in positive ion mode. All data were acquired employing Agilent 6410 Quantitative Analysis version analyst data processing software.

Statistical Analysis. Each experiment with primary human hepatocytes was conducted in triplicate, and data were expressed as mean \pm standard deviation (S.D.). Student's t-test was performed for statistically analysis.

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Results

GW4064 represses CYP3A4 expression and activity. To determine whether FXR activation alters CYP3A4 expression, human hepatocytes from three different donors were treated with GW4064 (a FXR agonist, 1 μ M) or DMSO for 48 hr, and then CYP3A4 mRNA expression and enzyme activity levels were examined. GW4064 significantly repressed mRNA expression levels (Fig. 1A) and enzyme activity (Fig. 1B) of CYP3A4.

FXR is capable of causing transactivation or transrepression of its target gene promoters. To determine whether FXR directly transrepresses CYP3A4 promoter, luciferase reporter assays were performed in HepG2 cells. The cells were transfected with a CYP3A4 promoter vector, and treated with GW4064 (or vehicle control), after which luciferase activities were measured. GW4064 did not decrease CYP3A4 promoter activity (Fig. 1C), suggesting that direct FXR transrepression of CYP3A4 promoter is an unlikely mechanism for CYP3A4 repression by GW4064. On the other hand, in human hepatocytes, CYP3A4 repression by GW4064 was accompanied by increased SHP expression (Fig. 1D). Downregulation of CYP3A4 and upregulation of SHP were also observed in human hepatocytes treated with cholic acid, a natural ligand of FXR (Fig. 1E), suggesting that FXR activation leads to CYP3A4 repression. In HEK293T cells co-transfected with a luciferase vector where *luc* expression is driven by 2.2-kb of *SHP* promoter (along with FXR and RXR expression plasmids), GW4064 enhanced SHP promoter activity in a concentration-dependent manner with an estimated EC₅₀ of 75 nM.

SHP represses PXR and CAR transactivation of CYP3A4 promoter. SHP is known to repress activity of multiple transcription factors, which potentially include PXR, CAR, and GR that are involved in the regulation of CYP3A4 expression. Considering the absence of prominent PXR or CAR ligands, but the presence of dexamethasone, in the culture media for human hepatocytes, we first examined whether SHP represses regulation of CYP3A4 promoter activity via GR, by using promoter reporter assays. HepG2 cells were co-transfected with a luciferase vector where *luc* expression is driven by xenobiotic response element of *CYP3A4* that binds directly to PXR or CAR (Faucette et al., 2006), along with expression vectors of GR and/or SHP. The transfected cells were treated with vehicle control or

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dexamethasone, and luciferase activity was measured. The results showed that activation of GR (by dexamethasone treatment) led to a significant increase in CYP3A4 promoter activity (Fig. 2A, lane 2 vs. 4), consistent with the previously reported data (Pascussi et al., 2001). The enhanced CYP3A4 promoter activity, however, was abrogated by SHP (Fig. 2A, lane 4 vs. 8).

GR activation leads to increased CYP3A4 expression via enhancing both the expression and activity of PXR and CAR (Pascussi et al., 2001). To determine whether GW4064 alters the GR-mediated regulation of PXR or CAR expression, mRNA expression levels of PXR and CAR were measured in human hepatocytes treated with GW4064 (or vehicle control) by using qRT-PCR. The result showed that PXR or CAR expression did not differ between the groups (data not shown), suggesting that GW4064 does not repress GR action on PXR or CAR expression. Then, to examine whether SHP decreases PXR or CAR transactivation of CYP3A4, the promoter reporter assay was performed. The results showed that rifampin-activated PXR dramatically enhanced CYP3A4 promoter activity as expected (Fig. 2B, lane 2 and 4), and this was significantly repressed by SHP (Fig. 2B, lane 4 vs. 8). CAR transfection led to increases in CYP3A4 promoter activity (Fig. 2C, lane 1 vs. 2), consistent with the previous report that CAR is constitutively active even in the absence of ligands when transfected into immortalized cells (Baes et al., 1994). The enhanced CYP3A4 promoter activity was decreased upon co-expression of SHP (Fig. 2C, lane 4 vs. 8). Together, these results indicate that SHP represses PXR/CAR transactivation of CYP3A4 promoter.

GW4064 enhances CYP2B6 promoter activity. PXR and CAR enhance expression of many hepatic drug-metabolizing enzymes including CYP2B6 (Faucette et al., 2006). To determine whether the SHP-mediated repression of PXR and CAR action alters expression of other target genes, we examined expression of CYP2B6 in human hepatocytes treated with GW4064. Interestingly, CYP2B6 mRNA levels were significantly increased (rather than decreased) in two of three different batches of hepatocytes (Fig. 3A). To examine whether GW4064 can directly activate CYP2B6 promoter, promoter reporter assays were performed. HepG2 cells were transfected with a luciferase vector where *luc* expression is driven by 1.8-kb upstream regulatory region of *CYP2B6*, treated with vehicle control or GW4064, and luciferase

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activity was measured. The results showed that GW4064 treatment led to ~3-fold increase in CYP2B6 promoter activity (Fig. 3B).

To determine whether the enhanced CYP2B6 promoter activity is through FXR-transactivation of CYP2B6 promoter, promoter reporter assays were performed in HEK293T cells co-transfected with a CYP2B6-promoter driven luciferase vector and expression vector for FXR and RXR. The cells were treated with GW4064 (or vehicle control), and luciferase activity was measured. A luciferase vector harboring 2.2-kb *SHP* promoter was included as a control. Results showed that GW4064 enhanced CYP2B6 promoter activity ~2.5-fold (Fig. 3C, lane 1 and 4). This induction was of a similar magnitude as the increase in *SHP* promoter activity by GW4064 (Fig 3C, lane 5 and 6). Collectively, these data indicate that the enhanced CYP2B6 expression could be due to increased CYP2B6 promoter activity by FXR activation.

Discussion

In this study, we examined the effects of FXR activation on expression of CYP3A4 by using freshly isolated primary human hepatocytes. Our results showed that GW4064 significantly decreases the expression of the major drug-metabolizing enzyme CYP3A4 in human hepatocytes, and that this is in part attributable to increased *SHP* expression upon FXR activation. We found that a different inducer of *SHP*, all-trans retinoic acid (Koh et al., 2014), also repressed CYP3A4 expression in human hepatocytes (data not shown; a manuscript in preparation), further supporting the important role of *SHP* in CYP3A4 regulation. CYP3A4 expression is regulated by nuclear receptors including PXR and CAR. While PXR and CAR are activated to the fullest extent upon binding of respective ligands to the receptors, both transcription factors appear activated at basal levels in human hepatocytes (Zamule et al., 2008; Hariparsad et al., 2009). This could be in part due to the presence of corticosteroids in culture media that are known to enhance the expression as well as transcriptional activity of PXR and CAR, via glucocorticoid receptor (GR) (Pascussi et al., 2000; Pascussi et al., 2001). Our results from luciferase reporter assays showed that *SHP* represses actions of GR as well as those of PXR and CAR on CYP3A4

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promoter, suggesting that repression of these transcriptional activators on CYP3A4 promoter by SHP may potentially trigger CYP3A4 repression by GW4064. Interestingly, such repressive action of SHP on PXR/CAR target gene promoter was not shown for CYP2B6, in part because GW4064 activates CYP2B6 promoter. The model for differential regulation of CYP3A4 and CYP2B6 by GW4064 is shown in Fig. 4.

In addition to PXR and CAR, multiple other transcription factors including peroxisome proliferator-activated receptor (PPAR) α and liver X receptor (LXR) upregulate CYP3A4 expression (Drocourt et al., 2002; Duniec-Dmuchowski et al., 2007; Thomas et al., 2013). However, potential involvement of PPAR α and LXR in CYP3A4 repression by GW4064 appears unlikely. Our results from human hepatocytes indicate that GW4064 does not alter the mRNA expression of the representative target genes of these nuclear receptors, including MDR3 (Ghonem et al., 2014), FASN, and SREBF1 (Wagner et al., 2003) (data not shown). Inflammation is also known to alter CYP3A4 expression; CYP3A4 expression is down-regulated in the state of inflammation (Aitken et al., 2006). In GW4064-treated human hepatocytes, there were no increases in the expression levels of inflammatory marker TNF α (data not shown) and no significant changes in the gross morphology of hepatocytes (that are indicative of drug toxicity or inflammation). Together, these findings suggest that PPAR α , LXR, or inflammation plays minimal roles, if any, in CYP3A4 repression by GW4064.

Previous studies using HepG2 cells have shown that GW4064 *increases* (not decreases as shown in our study) CYP3A4 expression (Gnerre et al., 2004); GW4064 (at 1 μ M) increased CYP3A4 mRNA levels by 1.8-fold in HepG2 cells, potentially by FXR binding to \sim 7.8-kb upstream region of *CYP3A4* and transactivating the promoter (Gnerre et al., 2004). Also, GW4064 enhanced expression of *Cyp3a11* (a CYP3A4 homolog gene) in mice by \sim 2-fold (Gnerre et al., 2004). While the reason for discrepancy between the previous and our study is unclear, it appears possible that in HepG2 cells, the expression levels of key hepatic transcription factors including PXR and CAR may have been low for SHP to exhibit any repressive action. Also, significant interspecies differences in the regulation of drug-metabolizing enzymes expression hamper the extrapolation of animal data to humans (Lu and Li, 2001; Vignati et al., 2004). Considering that primary human hepatocytes have served as a gold standard tool to study

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regulation of hepatic drug-metabolizing enzymes, it appears likely that GW4064 decreases CYP3A4-mediated drug metabolism in humans through SHP-mediated repression of CYP3A4 expression.

FXR agonists including GW4064 are currently under development as potential therapeutic agents for metabolic diseases such as obesity, type 2 diabetes, hypertriglyceridaemia, atherosclerosis, and non-alcoholic steatohepatitis (reviewed in (Thomas et al., 2008)). As CYP3A4 repression by GW4064 is potentially triggered by increased expression of SHP and that SHP induction is a class effect of FXR agonists, other FXR agonists are also expected to repress CYP3A4 expression, potentially causing drug-drug interactions with CYP3A4 substrates. It remains difficult, however, to quantitatively predict the clinical outcome of these interactions based only on the results from human hepatocytes. This is in part due to the long degradation half-lives of CYP3A4 protein (i.e., 36-50 hr (Fromm et al., 1996)) such that CYP3A4 protein levels in human hepatocytes do not reach the steady state after the typical time of drug treatment (e.g., 48 hr) passes. In accordance, CYP3A4 mRNA expression decreased >50% in GW4064-treated hepatocytes in this study while the decrease in catalytic activity of CYP3A4 (as determined in S9 fraction of hepatocytes) was only 10-30%. The clinical consequences of FXR agonists repressing CYP3A4 expression, thus, remain to be examined.

In conclusion, we showed that GW4064 represses CYP3A4 expression, potentially through upregulating SHP expression and subsequent repression of PXR and CAR transactivation of CYP3A4 promoter. This suggests that drug-drug interactions may occur clinically between CYP3A4 substrates and FXR agonists that are currently under development for the treatment of metabolic diseases.

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Authorship Contributions

Participated in research design: Zhang, Pan, and Jeong

Conducted experiments: Zhang, Pan, and Pan

Performed data analysis: Zhang, Pan, and Jeong

Wrote or contributed to the writing of the manuscript: Zhang, Pan, and Jeong

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Footnotes

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Figure Legends

Fig. 1. GW4064 represses CYP3A4 expression. Primary human hepatocytes (HH) from three different donors were treated with vehicle (DMSO) or GW4064 (1 μ M) for 48 hr in triplicate. RNAs were collected and mRNA levels of CYP3A4 (A) and SHP (D) were determined by using qRT-PCR. (B) CYP3A4 activity was measured in S9 fractions from the GW4064- or vehicle-treated hepatocytes using midazolam. Data shown are production rates of hydroxymidazolam in pmol/hr/mg protein. (C) HepG2 cells were co-transfected with pGL3-CYP3A4 and *Renilla* luciferase vector. On the next day, the cells were treated with vehicle (DMSO) or GW4064 (1 μ M) for 24 hr, and dual luciferase assays were performed. (E) Primary human hepatocytes were treated with vehicle (DMSO) or cholic acid (50 μ M) for 48 hr in triplicate. RNAs were collected to measure CYP3A4 and SHP mRNA expression by qRT-PCR. (F) HEK293T cells were co-transfected with pGL2-SHP and *Renilla* luciferase vectors, along with FXR and RXR expression vectors. On the next day, the cells were treated with vehicle (DMSO) or GW4064 (0.1, 1, 10, 100, 1000, or 10,000 nM) for 24 hr, and dual luciferase assays were performed. Data shown are mean \pm standard deviation. *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. vehicle-treated group; NS, statistically not significant.

Fig. 2. SHP represses PXR and CAR transactivation of CYP3A4 promoter. (A) HepG2 cells were co-transfected with pGL3-CYP3A4 and *Renilla* luciferase vector, along with expression vectors for GR or SHP. On the next day, the cells were treated with vehicle (DMSO) or dexamethasone (1 μ M) for 24 hr, and dual luciferase assays were performed. (B) HepG2 cells were co-transfected with pGL3-CYP3A4 and *Renilla* luciferase vector, along with expression vectors for PXR or SHP. On the next day, the cells were treated with vehicle (DMSO) or rifampin (5 μ M) for 24 hr, and dual luciferase assays were performed. (C) HepG2 cells were co-transfected with pGL3-CYP3A4 and *Renilla* luciferase vector, along with

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expression vectors for CAR or SHP. The cell lysates were collected after 24 hr, and dual luciferase assays were performed. **, $p < 0.01$; ***, $p < 0.001$.

Fig. 3. GW4064 upregulates CYP2B6 expression via FXR. (A) Primary human hepatocytes from three different donors were treated with vehicle (DMSO) or GW4064 (1 μ M) for 48 hr in triplicate. RNA was collected and CYP2B6 mRNA expression level was measured by using qRT-PCR. (B) HepG2 cells were co-transfected with pGL3-CYP2B6 and *Renilla* vector, treated with vehicle or GW4064 (1 μ M), and dual luciferase assay was performed. (C) HEK293T cells were co-transfected with pGL3-CYP2B6 (or pGL2-SHP) and *Renilla* vector, along with FXR and RXR expression vectors. On the next day, the cells were treated with vehicle (DMSO) or GW4064 (1 μ M). The cell lysates were collected after 24 hr, and dual luciferase assays were performed. **, $p < 0.01$; ***, $p < 0.001$.

Fig. 4. Differential regulation of CYP2B6 and CYP3A4 by GW4064. FXR activation by GW4064 leads to decreased CYP3A4 expression, potentially through SHP upregulation and subsequent repression of PXR and CAR transactivation of CYP3A4 promoter. On the other hand, GW4064 enhances CYP2B6 promoter activity.

Fig. 1

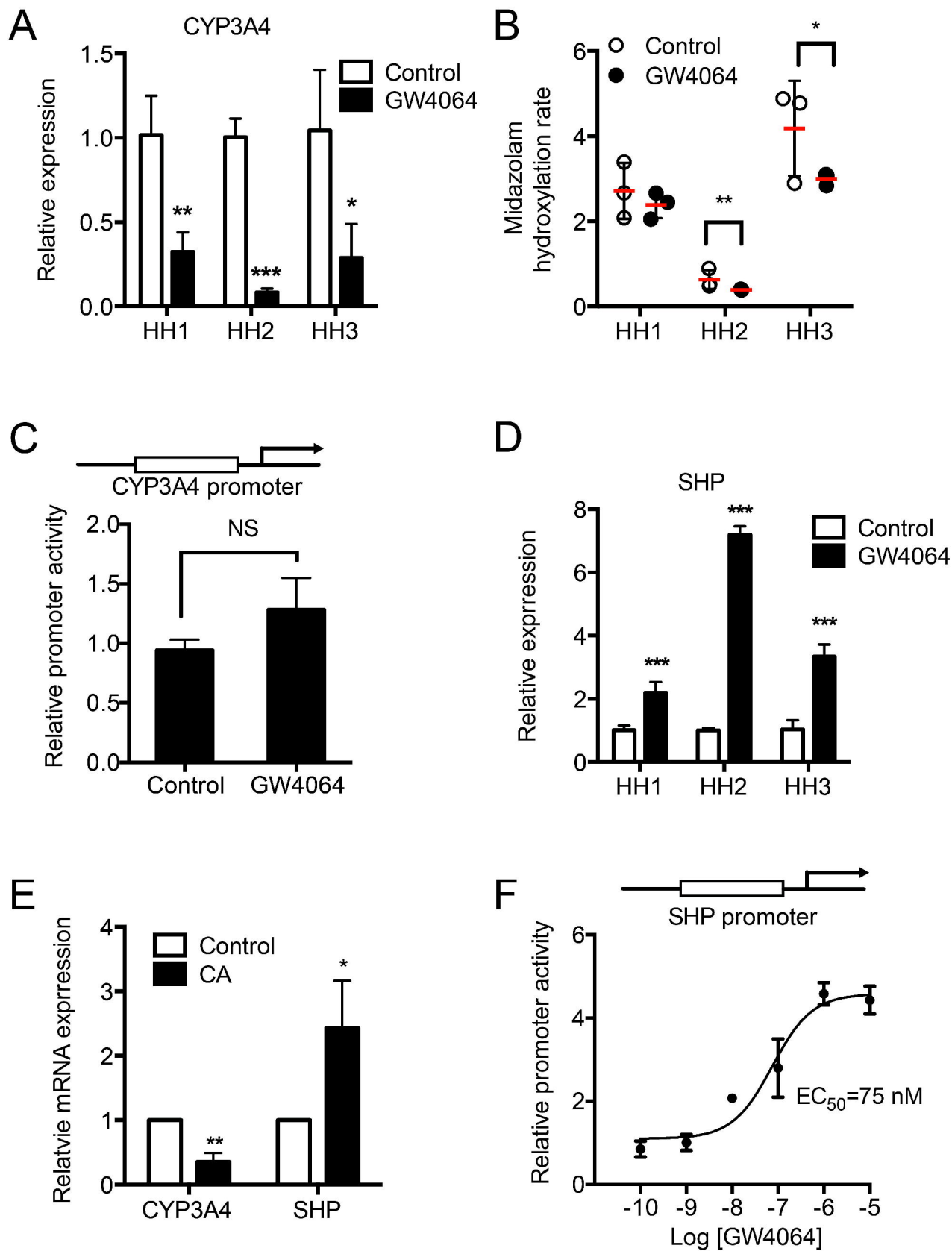


Fig. 2

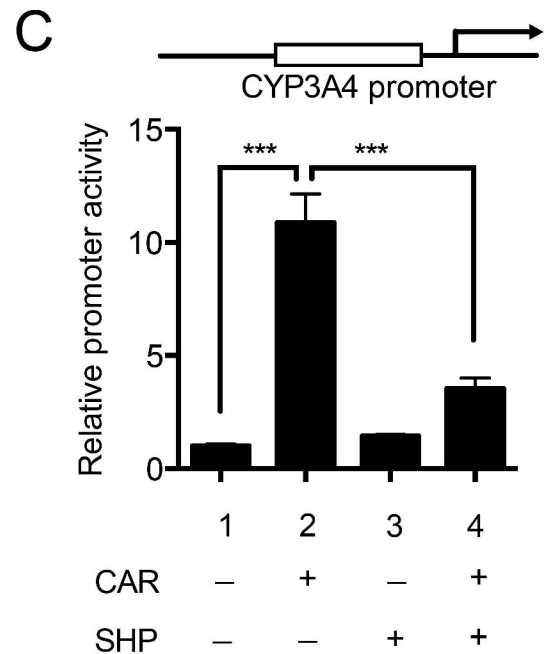
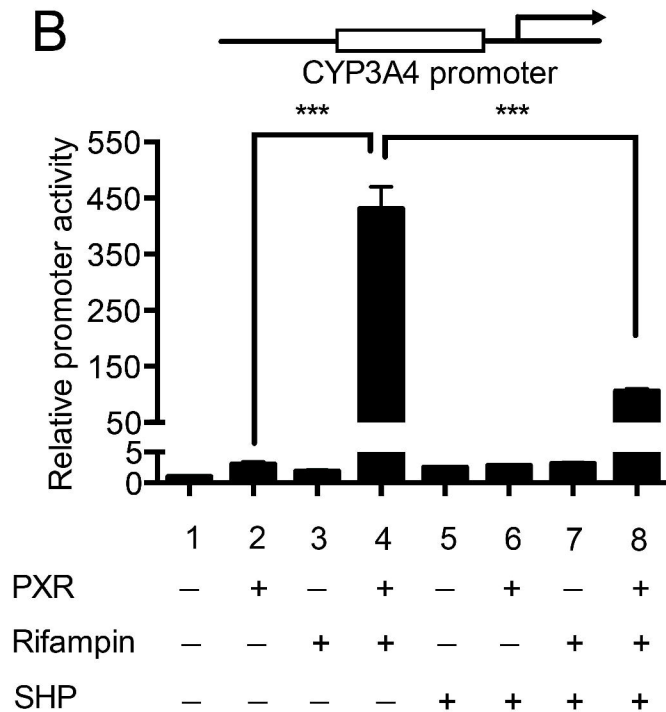
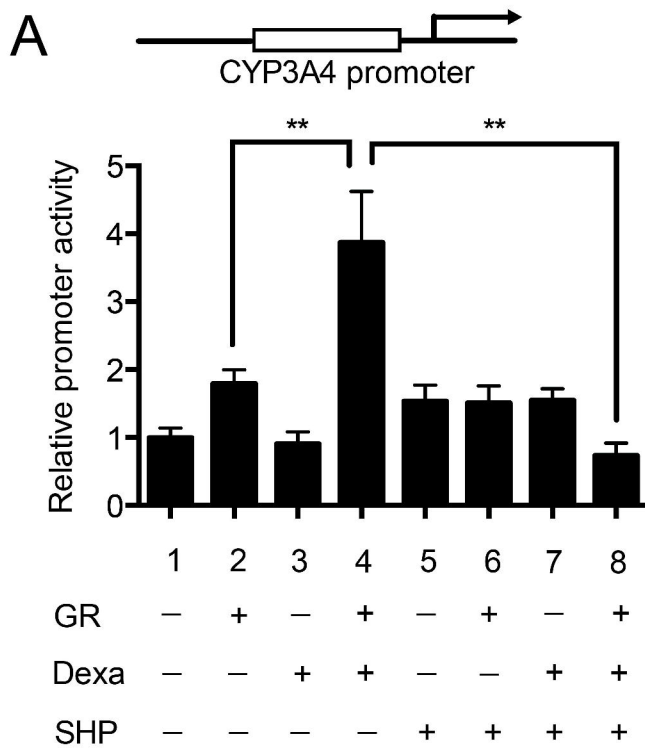
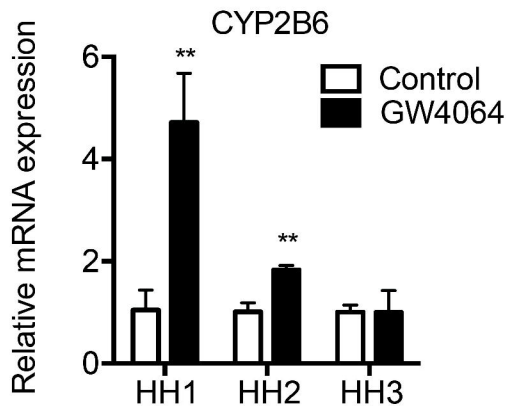
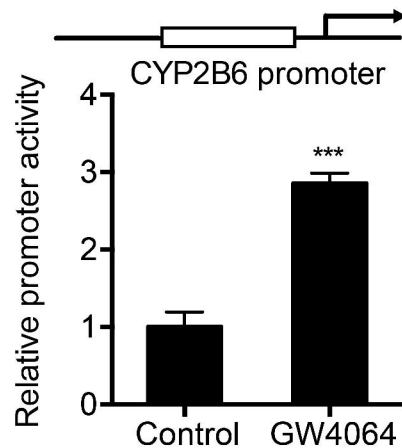


Fig. 3

A



B



C

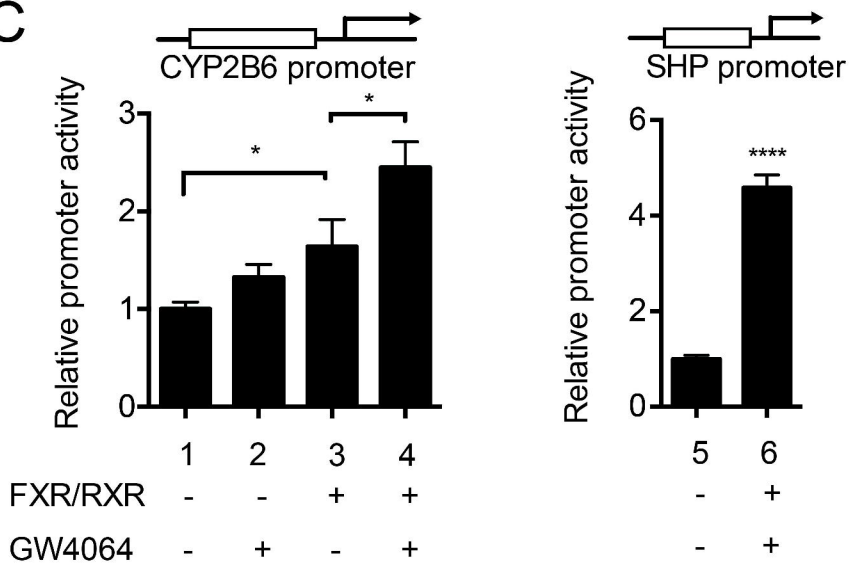
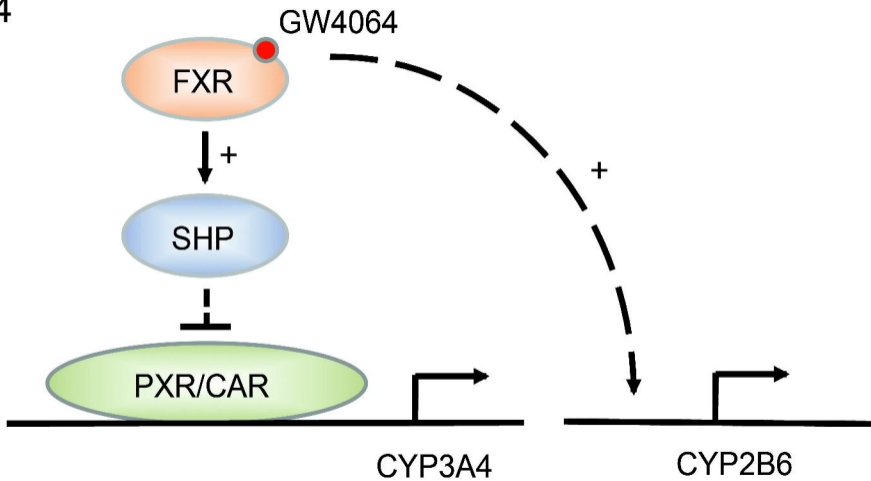


Fig. 4



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GW4064, an agonist of farnesoid X receptor (FXR), represses CYP3A4 expression in human hepatocytes by inducing small heterodimer partner (SHP) expression

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Drug Metabolism and Disposition

Table. S1. Sequences of oligonucleotides used in qRT-PCR

Gene	Forward primer	Reverse primer
<i>SHP (NROB2)</i>	TCAAGTCCATTCGACCAGC	AAGAAGGCCAGCGATGTCAA
<i>PXR</i>	GATGATCATGTCCGACGAG	GACTTCAGAGTAATGGCGA
<i>MDR3</i>	AATTTATCCTGCCAATCGGA	GCATCAGCAGCAAACAAAAA
<i>FASN</i>	ACAGGGACAACCTGGAGTTC	TGTGGTCCCACCTTGATGAGT
<i>SREBF1</i>	TTCTCACCTCCAGCTCTGT	GGTCCTGGAAGTCAGTCCAT
<i>TNFα</i>	GCCAGAGGGCTGATTAGAGA	TCAGCCTCTTCTCCTTCCTG