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Clopidogrel and Gemfibrozil Strongly Inhibit the CYP2C8-Dependent Formation of 3-Hydroxydesloratadine and Increase Desloratadine Exposure In Humans

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ABSTRACT

A recent in vitro study suggested that CYP2C8 is essential in the metabolism of desloratadine, an H1 receptor antagonist. If the proposed biotransformation mechanism takes place in vivo in humans, desloratadine could serve as a selective CYP2C8 probe substrate in drug-drug interaction studies. Glucuronide metabolites of clopidogrel and gemfibrozil act as time-dependent inhibitors of CYP2C8, but they have not been compared clinically. We conducted a randomized crossover study in 11 healthy subjects to characterize the involvement of CYP2C8 in desloratadine metabolism and to compare the CYP2C8 inhibitory strength of clopidogrel (300 and 75 mg on two following days) with that of gemfibrozil (600 mg BID for 5 days). Compared with placebo (control), clopidogrel increased the area under the plasma concentration-time curve (AUC0–71 h) to 52% (P = 5 × 10⁻⁵) and 6% (P = 2 × 10⁻⁵), respectively. Moreover, the 3-hydroxydesloratadine:desloratadine AUC0–71 h ratios were 21% (P = 7 × 10⁻⁹) and 1.7% (P = 8 × 10⁻¹¹) of control during the clopidogrel and gemfibrozil phases. Our results confirm that CYP2C8 plays a critical role in the formation of 3-hydroxydesloratadine in humans, making desloratadine a potential CYP2C8 probe substrate. Furthermore, the findings corroborate the previous estimates that clinically relevant doses of clopidogrel cause strong CYP2C8 inhibition, whereas those of gemfibrozil almost completely inactivate the enzyme in humans.

Introduction

Desloratadine is a potent, nonsedating H1 histamine receptor antagonist indicated for symptomatic relief of urticaria and allergic rhinitis (https://www.accessdata.fda.gov/drugsatfda_docs/label/2014/021165s017,021300v014,021312v015,021563s003bl.pdf; Henz, 2001). It has linear pharmacokinetics in the 5- to 20-mg dose range, is 82%–87% bound to plasma proteins, and has a terminal half-life (t1/2) of about 21–27 hours (Henz, 2001; Murdoch et al., 2003; Molimard et al., 2004). It is extensively metabolized, and its major metabolite is 3-hydroxydesloratadine, which is further glucuronidated to 3-hydroxydesloratadine-O-glucuronide (Molimard et al., 2004). A recent in vitro study suggested a central role for cytochrome P450 2C8 (CYP2C8) in the formation of the 3-hydroxy metabolite, involving initial N-glucuronidation of desloratadine followed by CYP2C8-dependent oxidation of the N-glucuronide and subsequent rapid deconjugation (Fig. 1) (Kazmi et al., 2015a). The proposed CYP2C8-selective metabolism and the benign safety profile could make desloratadine a practical CYP2C8 probe substrate in drug-drug interaction (DDI) studies (Prenner et al., 2006).

CYP2C8 is involved in the bioactivation and biodeactivation of various drugs, and its activity is essential for the formation of the active metabolites of several drugs, including clopidogrel, an antithrombotic agent. The acyl-β-D-glucuronide metabolite of the antiplatelet agent clopidogrel and the 1-O-glucuronide metabolite of the fibrin acid derivative gemfibrozil act as mechanism-based inhibitors of CYP2C8 (Ogilvie et al., 2006; Tornio et al., 2014; Backman et al., 2016). Accordingly, in humans, clopidogrel and gemfibrozil have significantly increased the exposure to several CYP2C8 substrates such as cerivastatin, repaglinide, pioglitazone, montelukast, and dasabuvir (Backman et al., 2002, 2016; Niemi et al., 2003; Jaakkola et al., 2005; Karonen et al., 2010; Tornio et al., 2014; Itkonen et al., 2016, 2018, 2019). Recently, both clopidogrel and gemfibrozil were suggested as model CYP2C8 inhibitors for DDI studies in the FDA Draft Guidance for Clinical Drug Interaction Studies (https://www.fda.gov/downloads/drugs/guidances/ucm292362.pdf). The CYP2C8-inhibitory strength of clopidogrel, however, has not been directly compared with that of gemfibrozil in a clinical setting.

Glucuronide metabolites have generally been regarded as relatively stable and physiologically inactive compounds, but owing to the reactivity of acyl glucuronides as well as their ability to act as perpetrators in drug interactions, these presumptions have been challenged by Ogilvie et al. (2006), Regan et al. (2010), Tornio et al. (2014), and Backman et al. (2016). Apart from the ability of CYP2C8 to reactivate glucuronides of gemfibrozil, clopidogrel, and celecoxib, leading to CYP2C8 inactivation, several
Materials and Methods

Subjects and Study Design. Twelve subjects were recruited for the study, but one of them withdrew from the study after the second phase for personal reasons. Eleven healthy nonsmoking volunteers (six women, five men; age range, 20–29 years; body mass index range, 18.6–27.3 kg/m²) participated in the study after giving written informed consent. Their health was confirmed by medical history, clinical examination, and routine laboratory tests before entering the study. All participants had normal blood platelet counts and hemoglobin values. None of the subjects used oral contraceptives or other continuous medication.

The primary objectives of the present clinical study were to examine the role of CYP2C8 in desloratadine metabolism and evaluate the applicability of desloratadine as a CYP2C8 probe substrate. Moreover, we aimed to compare head-to-head the CYP2C8 inhibitory effects of the typical clinically used doses of clopidogrel and gemfibrozil in vivo in humans.

Determination of Plasma Desloratadine and Its Metabolites. Prior to quantification by liquid chromatography–tandem mass spectrometry (LC-MS/MS), the plasma samples were purified by solid phase extraction using Strata-X Polymeric Reversed Phase, 96-Well Plate (10 mg/well; Phenomenex, Torrance, CA). In brief, the samples (120 μl) were mixed with 100 μl of internal standard solution containing desloratadine-d5 and 3-hydroxydesloratadine-d4 (5 ng/ml) in 50 mM ammonium acetate (pH 4.7), and drawn through the preconditioned extraction plate. The wells were then washed with 10% methanol and the analytes were eluted with methanol containing 1% formic acid. Finally, the sample extracts were dried using a centrifugal evaporator (GeneVac; Thermo Fisher Scientific, Waltham, MA) and reconstituted in 100 μl of mobile phase starting composition. The analytes were separated on a Kinetex C18 XB column (2.6-μm particle size, 2.1 × 100-mm internal diameter; Phenomenex) using a Nexera X2 ultra-high-performance liquid chromatography system (Shimadzu, Kyoto, Japan). The mobile phase consisted of 0.02% formic acid in 5 mM ammonium acetate (mobile...
phase A) and acetonitrile (mobile phase B), and a solvent gradient was adopted as follows for the total run time of 11 minutes: 3 minutes at 20% B; a linear increase from 20% B to 63% B over 3 minutes; 1 minute at 90% B; 4 minutes equilibration at 20% B. The flow rate and the column temperature were maintained at 300 μL/min and 30°C, and an aliquot of 2 μL was injected into the system. A Sciex 5500 Qtrap tandem mass spectrometer (Sciex, Toronto, Ontario, Canada) interfaced with an electrospray ion source was used for the mass spectrometric detection. The mass spectrometer was operated in the positive-polarity multiple-reaction mode and the Q1 and Q3 quadrupoles were set at unit mass resolution. The mass transitions (m/z) 311 to 259, 327 to 275, and 503 to 327 were selected for desloratadine, 3-hydroxydesloratadine, and 3-hydroxydesloratadine-N-glucuronide, respectively. The lower limit of quantification (LLQ) for desloratadine and 3-hydroxydesloratadine were 0.05 and 0.025 ng/ml, respectively. A signal-to-noise ratio of 3:1 was used as a limit of detection for 3-hydroxydesloratadine N-glucuronide, and the quantities were given in arbitrary units (units per milliliter) relative to the ratio of the peak heights of the 3-hydroxydesloratadine N-glucuronide to the internal standard (3-hydroxydesloratadine-d4). The day-to-day coefficients of variation (CV) were 6.0% (0.2 ng/ml), 2.6% (2.0 ng/ml), and 4.6% (20 ng/ml) for desloratadine, and 4.2% (0.1 ng/ml), 2.1% (1.0 ng/ml), and 3.5% (10 ng/ml) for 3-hydroxydesloratadine.

### Determination of Plasma Gemfibrozil and Gemfibrozil-1-O-Glucuronide.

The plasma gemfibrozil and gemfibrozil-1-O-glucuronide were purified using a simple protein precipitation with acetonitrile (1:3) containing the corresponding deuterium-labeled internal standards, gemfibrozil-d6, and gemfibrozil-1-O-glucuronide-d6 (0.5 μg/ml). The plasma extract was then further diluted with water and introduced to the Sciex 5500 Qtrap LC-MS/MS System. The mass spectrometric detection. The mass spectrometer was operated in the positive-ion mode with a m/z 503 to 327 were selected for desloratadine, 3-hydroxydesloratadine, and 3-hydroxydesloratadine-N-glucuronide, respectively. The lower limit of quantification (LLQ) for desloratadine and 3-hydroxydesloratadine were 0.05 and 0.025 ng/ml, respectively. A signal-to-noise ratio of 3:1 was used as a limit of detection for 3-hydroxydesloratadine N-glucuronide, and the quantities were given in arbitrary units (units per milliliter) relative to the ratio of the peak heights of the 3-hydroxydesloratadine N-glucuronide to the internal standard (3-hydroxydesloratadine-d4). The day-to-day coefficients of variation (CV) were 6.0% (0.2 ng/ml), 2.6% (2.0 ng/ml), and 4.6% (20 ng/ml) for desloratadine, and 4.2% (0.1 ng/ml), 2.1% (1.0 ng/ml), and 3.5% (10 ng/ml) for 3-hydroxydesloratadine.

### Pharmacokinetics.

The peak plasma concentration (Cmax, time to Cmax (tmax), area under the plasma concentration-time curve from time 0–71 hours (AUC0–71h), AUC0–∞ (when appropriate), and t1/2 were calculated for desloratadine and its metabolites by standard noncompartmental methods using Phoenix WinNonlin, version 6.4 (Certara, Princeton, NJ). In the gemfibrozil phase, the concentrations of 3-hydroxydesloratadine and desloratadine 3-hydroxy-O-glucuronide were under the LLQ at several time points after the administration of desloratadine. To calculate the AUC0–71h, the aforementioned concentration values lower than LLQ were replaced with values equal to half of the LLQ. For gemfibrozil, desloratadine, and its metabolites, Cmax, tmax, t1/2, and AUC0–12 h were calculated.

### Genotyping.

Buffy coats were prepared from 9 ml of whole-blood EDTA samples after plasma separation. Genomic DNA was extracted from the buffy coats using the Maxwell 16 LEV Blood DNA Kit on a Maxwell 16 Research automated nucleic acid extraction system (Promega, Madison, WI). The participants were genotyped for the CYP2C8*2 (rs11572103), *3 (rs10599681 and rs11572080), and *4 (rs10589330) alleles with commercially available or custom TaqMan assays with OpenArray technology on a QuantStudio 12K Flex real-time PCR system (Life Technologies/Thermo Fisher Scientific, Carlsbad, CA).

Two of the subjects had the CYP2C8*1/*3 genotype, and two had the CYP2C8*1/*4 genotype, whereas the others had the CYP2C8*1/*1 genotype. There were no apparent genotype-dependent differences in the pharmacokinetics of desloratadine or its metabolites (data not shown). Owing to the small sample size and lack of subjects with desloratadine “poor metabolizer” phenotype, UGT2B10 was not included in the genotyping panel.

### Drug-Drug Interaction Predictions.

Inhibition of CYP2C8 by clopidogrel acetyl-β-D-glucuronide and gemfibrozil 1-O-glucuronide was predicted using mechanism-based in vitro inhibition values from the literature with a mechanistic static model according to the following equation:

$$\text{Activity remaining} = \frac{k_{deg} - k_{kinact}}{K_{I} + [I]}$$

where $k_{deg}$ is the rate constant of hepatic P450 degradation in the absence of the inhibitor, $k_{kinact}$ is the maximum inactivation rate, $I$ is the uninhibited inhibitor concentration at the enzyme site, and $K_{I}$ is the inhibitor concentration needed to cause half of $k_{kinact}$. A half-life of 22 hours for CYP2C8 was used, corresponding to $k_{deg}$ of 0.000525 (Backman et al., 2009). The $k_{kinact}$ values were 2.82 1/min and 0.21 1/min, and $K_{I}$ values were 9.9 μM and 20 μM for clopidogrel acetyl-β-D-glucuronide and gemfibrozil 1-O-glucuronide, respectively (Ogilvie et al., 2006; Torino et al., 2014).

### Statistical Analysis.

On the basis of our previous drug interaction studies, twelve subjects were estimated to be adequate to detect a 30% change in the AUC of the victim drug and its main metabolite between the placebo (control) and clopidogrel and gemfibrozil phases, with a power of at least 80% (α level 5%). The results are expressed as geometric means and geometric mean ratios with geometric CV or 90% confidence intervals (CI) unless stated otherwise. Logarithmic transformation was used for pharmacokinetic variables, except $t_{1/2}$, before statistical analysis. The pharmacokinetic variables were compared by repeated-measures analysis of variance with treatment phase as a within-subjects factor, followed by pairwise comparisons with Bonferroni-corrected analysis of variance. The $t_{1/2}$ data were compared using the Wilcoxon signed rank test. Correlations between the fold-change in desloratadine AUC and the 3-hydroxydesloratadine:desloratadine AUC0–71 ratio between clopidogrel and gemfibrozil phases were quantified as Pearson’s correlation coefficients. P-values below 0.05 were considered statistically significant. Statistical analyses were performed using SPSS Statistics for Windows version 22.0 (IBM Corporation, Armonk, NY).

### Results

#### Effect of Clopidogrel on Parent Desloratadine.

Compared with placebo (control), clopidogrel increased desloratadine AUC0–∞ to 280% ($P = 3 \times 10^{-7}$; 90% CI 232%–338%). In addition, the Cmax of desloratadine was increased to 165% ($P = 0.0006$; 90% CI 133%–204%) and its $t_{1/2}$ prolonged from 17 to 26 hours (ratio to control 150%; $P = 0.0003$; 90% CI 130%–180%) by clopidogrel (Fig. 2; Table 1).

#### Effect of Clopidogrel on 3-Hydroxydesloratadine.

Clopidogrel decreased the 3-hydroxydesloratadine AUC0–∞ to 74% ($P = 5 \times 10^{-5}$; 90% CI 69%–80%) of control. Moreover, in the clopidogrel phase, the 3-hydroxydesloratadine:desloratadine AUC0–71 h ratio was 21% ($P = 7 \times 10^{-10}$; 90% CI 18%–24%) of that during the control phase. Compared with placebo, clopidogrel decreased 3-hydroxydesloratadine Cmax to 30% ($P = 1 \times 10^{-3}$; 90% CI 25%–37%), and prolonged the $t_{1/2}$ from 25 to 51 hours (ratio to control 200%; $P = 2 \times 10^{-5}$; 90% CI 170%–240%).

#### Effect of Clopidogrel on 3-Hydroxydesloratadine-O-Glucuronide.

During the clopidogrel phase, the Cmax of 3-hydroxydesloratadine-O-glucuronide was 71% ($P = 0.0008$; 90% CI 62%–81%) of that in the placebo phase. Furthermore, the Cmax of 3-hydroxydesloratadine-O-glucuronide was decreased to 37% ($P = 3 \times 10^{-7}$; 90% CI 31%–45%) and its $t_{1/2}$ prolonged from 19 to 35 hours (P = 2 × 10−5; 90% CI 163%–218%) by clopidogrel, compared with those in the control phase.

#### Effect of Gemfibrozil on Parent Desloratadine.

Compared with placebo (control), gemfibrozil increased desloratadine AUC0–∞ to 462% ($P = 4 \times 10^{-7}$; 90% CI 346%–616%). The Cmax of desloratadine was increased to 174% ($P = 0.0006$; 90% CI 137%–221%) of control and $t_{1/2}$ prolonged from 17 to 39 hours (ratio to control 220%; $P = 3 \times 10^{-8}$; 90% CI 200%–250%) by gemfibrozil (Fig. 2; Table 1).
The effect of clopidogrel (300 mg on day 3, followed by 75 mg on days 4 and 5) and gemfibrozil (600 mg twice daily on days 1–5) on the plasma concentrations of desloratadine (A), and its 3-hydroxy (B), and 3-hydroxy-O-glucuronide (C) metabolites. On day 3, desloratadine 5 mg was administered to 11 healthy volunteers 1 hour after the morning dose of each pretreatment. Data are presented as geometric means with 90% confidence intervals. For clarity, some error bars have been omitted. Insets depict the same data on semilogarithmic scale.
Effect of Gemfibrozil on 3-Hydroxydesloratadine. During the gemfibrozil phase, the concentrations of 3-hydroxydesloratadine were very low, and the 3-hydroxydesloratadine AUC0–71 h and 3-hydroxydesloratadine:desloratadine AUC0–71 h ratios were 6% (P = 2 × 10−8; 90% CI 4%–9%) and 1.7% (P = 8 × 10−11; 90% CI 1.3%–2.4%) of control, respectively. Gemfibrozil decreased the Cmax of 3-hydroxydesloratadine to 4% of control (P = 8 × 10−10; 90% CI 3%–5%). Owing to the extremely low 3-hydroxydesloratadine concentrations, its t1/2 could not be reliably determined in most of the subjects in the gemfibrozil phase.

Effect of Gemfibrozil on 3-Hydroxydesloratadine-O-Glucuronide. In the gemfibrozil phase, the AUC0–71 h of 3-hydroxydesloratadine-O-glucuronide was 6% (P = 7 × 10−8; 90% CI 4%–10%) and its Cmax was 4% (P = 3 × 10−8; 90% CI 2%–6%) of that in the placebo phase. In addition, the tmax of 3-hydroxydesloratadine-O-glucuronide was prolonged from 9.0 to 11.0 hours (P = 0.03). As with 3-hydroxydesloratadine, the plasma concentrations of 3-hydroxydesloratadine-O-glucuronide were too low for accurate calculation of its t1/2 during the gemfibrozil phase.

Head-to-Head Comparison of the Effects of Clopidogrel and Gemfibrozil. In general, gemfibrozil’s effect on desloratadine pharmacokinetics was stronger than clopidogrel’s (Fig. 2). By gemfibrozil, desloratadine AUC0–∞ was 165% (P = 2 × 10−5; 90% CI 142%–191%) of that by clopidogrel. Moreover, 3-hydroxydesloratadine AUC0–71 h was 12% (P = 1 × 10−5; 90% CI 9%–16%) and the 3-hydroxydesloratadine:desloratadine AUC0–71 h ratio was 8% (P = 7 × 10−10; 90% CI 7%–11%), compared with those in the clopidogrel phase. Additionally, in the gemfibrozil phase, the AUC0–71 h of 3-hydroxydesloratadine-O-glucuronide was 11% (P = 1 × 10−5; 90% CI 9%–15%) of that in the clopidogrel phase. There was a significant (Pearson two-tailed, P = 0.002; R2 = 0.80) correlation between the fold-change in desloratadine AUC0–∞ caused by clopidogrel and that by gemfibrozil (Fig. 2C).

Pharmacokinetic Variables of Clopidogrel and Gemfibrozil. The Cmax of clopidogrel and gemfibrozil, as well as their metabolites, were reached 1–3 hours after their ingestion (Table 2), i.e., around or soon after the time of desloratadine administration. There were 50- and 20-fold interindividual variations in parent clopidogrel Cmax and AUC0–12h, respectively. In addition, a 3- to 4-fold variation was observed in those of the active cis-5-thiol, carboxylic acid, and acyl-β-D-glucuronide metabolites of clopidogrel. The variation in the pharmacokinetic variables of gemfibrozil and its 1-O-glucuronide metabolite was smaller, only up to 2-fold. Consistent with previous DDI studies between clopidogrel or gemfibrozil and CYP2C8 substrates (Backman et al., 2002; Jaakkola et al., 2005; Tornio et al., 2014; Ikonen et al., 2016, 2018, 2019), no significant correlations were observed between the plasma concentrations of clopidogrel acyl-β-D-glucuronide or gemfibrozil 1-O-glucuronide and changes in desloratadine pharmacokinetics.

Discussion

Recently, CYP2C8 was suggested to have an essential role in the biotransformation of desloratadine to its major metabolite, 3-hydroxydesloratadine (Kazmi et al., 2015a). Clopidogrel and gemfibrozil are both clinically relevant CYP2C8 inhibitors, but previously their CYP2C8 inhibition potency had not been compared with each other in humans. In addition, selective, sensitive, and safe CYP2C8 probe substrates are required for DDI studies. For example, repaglinide, the CYP2C8 probe drug recommended by FDA (https://www.fda.gov/downloads/drugs/guidanceucmn/ucm292362.pdf), has also other disposition mechanisms, e.g., organic anion-transporting polypeptide 1B1 (OATP1B1)-mediated active hepatic uptake and metabolism by CYP3A4 (Tornio et al., 2012), which can complicate data interpretation. For these reasons, we conducted a clinical study in healthy volunteers to investigate the role of CYP2C8 in desloratadine metabolism, and to compare the CYP2C8 inhibitory properties of clopidogrel and gemfibrozil. Overall, both clopidogrel and gemfibrozil increased the exposure to desloratadine and radically reduced the concentrations of 3-hydroxydesloratadine.
On the basis of in vitro results, Kazmi et al. (2015a) suggested an unusual metabolic pathway, by which three distinct consecutive metabolic steps are required for the formation of 3-hydroxydesloratadine (Fig. 1). First, uridine 5'-diphospho-glucuronosyltransferase (UGT) 2B10 metabolizes desloratadine to its N-glucuronide. Subsequently, the N-glucuronide undergoes rapid oxidation to 3-hydroxydesloratadine-N-glucuronide very selectively by CYP2C8 (Kazmi et al., 2015a). Finally, the glucuronide moiety is deconjugated, most probably spontaneously, during or very shortly after the CYP2C8-mediated 3-hydroxylation (Kazmi et al., 2015a). The resulting 3-hydroxydesloratadine is further conjugated to O-glucuronide by UGT1A1, UGT1A3, and UGT2B15 (Ghosal et al., 2004). The investigators were unable to detect desloratadine-N-glucuronide, most probably owing to its instability and rapid CYP2C8-mediated oxidation (Kazmi et al., 2015a). Although desloratadine-N-glucuronide was undetectable in the experiments, desloratadine was not hydroxylated by CYP2C8 unless coincubated with UGT2B10, and thus the proposed pathway seems convincing (Kazmi et al., 2015a). Of note, desloratadine is also biotransformed to 5-hydroxy and 6-hydroxy metabolites by several P450 enzymes, namely CYP3A4, CYP2D6, and CYP2C19, in vitro (Barecki et al., 2001; McClellan and Jarvis, 2001; Kazmi et al., 2015b). However, these are only minor elimination pathways, and inhibition of these enzymes does not lead to clinically relevant changes in desloratadine pharmacokinetics in vivo (Henz, 2001; McClellan and Jarvis, 2001). Furthermore, the CYP3A4 and OATP1B1 inhibitor erythromycin did not significantly increase desloratadine exposure, suggesting lack of OATP1B1 contribution in desloratadine disposition (Henz, 2001).

In the clopidogrel phase of the present study, the AUCs of 3-hydroxydesloratadine and its O-glucuronide were markedly decreased, whereas the AUC of parent desloratadine was more than doubled. The resulting reduction in 3-hydroxydesloratadine:desloratadine AUC\(_{0-71\ h}\) ratio by clopidogrel is consistent with an average 80% inhibition of CYP2C8 activity, assuming that clopidogrel had no effect on non-CYP2C8-mediated metabolism of desloratadine N-glucuronide nor on hydrolysis of either of the N-glucuronide intermediates. With respect to clopidogrel's CYP2C8 inhibition potency, this estimate is in accordance with recent findings that clopidogrel causes about 5-fold increases in exposures to repaglinide and dasabuvir, which are among the most sensitive CYP2C8 substrates identified so far (Tornio et al., 2014; Itkonen et al., 2019).

In most individuals, gemfibrozil almost abolished 3-hydroxydesloratadine and its O-glucuronide from plasma, while reducing the 3-hydroxydesloratadine:desloratadine AUC\(_{0-71\ h}\) ratio to less than 2% of that during the placebo phase (Fig. 3A; Table 1). In addition, gemfibrozil increased desloratadine exposure in all subjects. Concerning the CYP2C8-inhibitory strength of gemfibrozil, the deductions inferred from the current results are highly concordant with the estimates from previous studies with gemfibrozil and the CYP2C8 substrate repaglinide, which indicated that clinical doses of gemfibrozil (600 mg BID) decrease CYP2C8 activity by >98% (Honkalainen et al., 2012). The effects of gemfibrozil on desloratadine pharmacokinetics resembled those of clopidogrel but were significantly more prominent, which is in line with the static predictions made on the basis of the unbound plasma C\(_{\text{max}}\) of gemfibrozil and clopidogrel glucuronides, and their in vitro mechanism-based inhibition parameters reported in the literature (Fig. 3F). The AUC of desloratadine was augmented 2.8-fold (range 1.8- to 4.7-fold) and 4.6-fold (range 2.3- to 7.9-fold) by clopidogrel and gemfibrozil, respectively. The increases in desloratadine AUC caused by clopidogrel and gemfibrozil correlated with each other (Fig. 3C), and with the 3-hydroxydesloratadine:desloratadine AUC\(_{0-71\ h}\) ratio in the placebo phase that probably reflects CYP2C8 activity at baseline (Fig. 3, D and E). These findings suggest that the basis of the effects of clopidogrel and gemfibrozil is a shared CYP2C8-selective mechanism and that interindividual variation in CYP2C8 enzyme activity largely explains the variability in sensitivity of desloratadine to CYP2C8 inhibition. The present and previous reports imply that clinical doses of gemfibrozil cause stronger interactions with CYP2C8 substrates than those of clopidogrel, thus revealing more sensitivity whether CYP2C8 participates in the metabolism of the victim drug (Table 3). However, gemfibrozil also inhibits the activity of certain membrane transporters, most notably hepatic OATP1B1 (Neuvonen et al., 2006; Tornio et al., 2017), which in turn is not inhibited by clopidogrel to a clinically relevant extent (Itkonen et al., 2015; Kim et al., 2016). Therefore, interpreting the DDI mechanisms can be challenging when gemfibrozil 600 mg BID is used as a model CYP2C8 inhibitor, if the victim drug is also a substrate of OATP1B1, as repaglinide and cerivastatin are (Backman et al., 2002, 2016; Niemi et al., 2003; Shitara et al., 2003; Niemi et al., 2005; Ogilvie et al., 2006; Tornio et al., 2017). In such cases, clopidogrel could be considered as an alternative model inhibitor.

Our current results imply that the metabolism of desloratadine to its 3-hydroxy metabolite is highly dependent on CYP2C8 activity in humans in vivo. Although gemfibrozil and clopidogrel increased the AUC of desloratadine less than they had increased the AUCs of repaglinide and dasabuvir (Table 3), the 3-hydroxydesloratadine:desloratadine ratio seemed to be highly sensitive to CYP2C8 inhibition. These observations corroborate the previous in vitro findings of Kazmi et al. (2015a) and suggest the possibility of using desloratadine as a CYP2C8 probe substrate. However, an important caveat is that the formation of 3-hydroxydesloratadine is also dependent on UGT2B10 activity (Kazmi et al., 2015b) and can probably be altered by inhibition and induction of UGT2B10. Of note, neither clopidogrel nor gemfibrozil is

### Table 2
Pharmacokinetic variables of clopidogrel, clopidogrel active cis-5-thiol metabolite, clopidogrel carboxylic acid, clopidogrel acyl-β-D-glucuronide, gemfibrozil, and gemfibrozil 1-O-glucuronide in 11 healthy volunteers after clopidogrel 300 mg or gemfibrozil 600 mg (fifth dose of twice daily dosing) ingested 1 hour before desloratadine 5 mg.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Clopidogrel or Gemfibrozil Phase</th>
</tr>
</thead>
<tbody>
<tr>
<td>C(_{\text{max}}) (ng/ml)</td>
<td>2.88 (0.48–23.2)</td>
</tr>
<tr>
<td>t(_{\text{max}}) (h)</td>
<td>1.5 (0.92–2.0)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>3.4 (2.6–10)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (ng/ml)</td>
<td>23.2 (11.0–41.2)</td>
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<tr>
<td>t(_{\text{max}}) (h)</td>
<td>0.92 (0.92–1.5)</td>
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<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>1.8 (1.3–3.9)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (ng/ml)</td>
<td>34.6 (16.3–66.0)</td>
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<tr>
<td>t(_{\text{max}}) (h)</td>
<td>9850 (7020–15,800)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>1.5 (0.92–1.5)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (gmol/l)</td>
<td>4.2 (3.1–8.0)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>35,400 (38,900–92,100)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (mg/l)</td>
<td>3980 (1750–7390)</td>
</tr>
<tr>
<td>t(_{\text{max}}) (h)</td>
<td>2.0 (1.5–3.0)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>3.3 (2.7–4.0)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (gmol/l)</td>
<td>14,400 (10,000–35,800)</td>
</tr>
<tr>
<td>t(_{\text{max}}) (h)</td>
<td>18.1 (14.3–29.1)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>2.0 (1.5–4.0)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (gmol/l)</td>
<td>2.3 (1.7–2.7)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>757 (56.5–130)</td>
</tr>
<tr>
<td>C(_{\text{max}}) (gmol/l)</td>
<td>14.5 (10.2–20.3)</td>
</tr>
<tr>
<td>t(_{\text{max}}) (h)</td>
<td>3.0 (1.5–6.0)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (ng × h/ml)</td>
<td>2.5 (1.7–3.4)</td>
</tr>
<tr>
<td>AUC(_{0-\text{last}}) (gmol × h/l)</td>
<td>8.00 (6.2–147)</td>
</tr>
</tbody>
</table>

Data are given as median with range.
Fig. 3. The individual change in 3-hydroxydesloratadine:desloratadine area under the plasma concentration time curve (AUC\textsubscript{0-71h}) ratio (A) and desloratadine AUC\textsubscript{0-\textinfty} (B) in the clopidogrel and gemfibrozil phase in comparison with placebo phase value. The correlation of desloratadine AUC\textsubscript{0-\textinfty} ratio between gemfibrozil to control and clopidogrel to control (C), and the correlation of 3-hydroxydesloratadine:desloratadine AUC\textsubscript{0-71h} ratio in the placebo phase with desloratadine AUC\textsubscript{0-\textinfty} clopidogrel to control ratio (D), and desloratadine AUC\textsubscript{0-\textinfty} gemfibrozil to control ratio (E), where the lines depict the linear regression with 95% CI. Predicted CYP2C8 inhibitory effect of gemfibrozil and clopidogrel using a static prediction equation whose basis was in vitro mechanism-based inhibition parameters of their glucuronide metabolites reported in the literature and their concentrations at the enzyme site (F). See Materials and Methods for details of the prediction equation.

\textbf{A}\textsuperscript{max,u,clop} unbound peak plasma concentration of clopidogrel acyl-\beta-D-glucuronide after 300 mg clopidogrel in the current study (0.83 \textmu M with an estimated fraction unbound of 10%); \textbf{C}_{\text{max,u,gemfi}} unbound peak plasma concentration of gemfibrozil 1-O-glucuronide after 600 mg gemfibrozil in the current study (3.9 \textmu M with an estimated fraction unbound of 11.5%).
known to inhibit UGT2B10. Moreover, in some individuals, the $t_{1/2}$ and drug exposure of desloratadine are very high, resulting in a "poor metabolizer" phenotype, which has a population frequency of 17% in African-Americans and 8% in Native American but is less common in Caucasians and Hispanics, with a frequency of about 2% (Molimard et al., 2004; Prenner et al., 2006; Ramanathan et al., 2007). The "poor metabolizer" phenotype and almost complete CYP2C8 inactivation by gemfibrozil result in surprisingly similar exposures to desloratadine and its 3-hydroxy metabolite (Prenner et al., 2006). However, the most probable explanation for this phenotype is that common UGT2B10 variants prevent the expression of functional UGT2B10 (Berg et al., 2010a, 2010b; Chen et al., 2010; Murphy et al., 2014; Fowler et al., 2015). Another possibility for this phenomenon is variability in CYP2C8, even though nonfunctional alleles in CYP2C8 are rare in all major populations (Zhou et al., 2017). In any case, individuals with desloratadine "poor metabolizer" phenotype are probably less sensitive to changes in desloratadine pharmacokinetics by CYP2C8 inhibitors. Consequently, desloratadine might be suboptimal as a CYP2C8 probe substrate in, for example, African-Americans. From another perspective, the good tolerability of desloratadine and its 3-hydroxy metabolite (Prenner et al., 2006). However, the most probable explanation for the effects of the CYP2C8 inhibitors gemfibrozil and clopidogrel on the pharmacokinetics of desloratadine is inhibition of CYP2C8-mediated 3-hydroxylation of a glucuronide metabolite of desloratadine, our results provide clinical evidence supporting the findings of several in vitro studies that CYP2C8 has the ability to oxidize glucuronide metabolites in humans (Backman et al., 2016). On the basis of X-ray crystallographic studies, the large CYP2C8 active site cavity is capable of binding substrates diverse in structure (Schoch et al., 2008). As demonstrated in molecular docking simulations, there is a hydrophilic region in the active site that can accommodate the glucuronide moiety of certain metabolites, including clopidogrel acyl-β-D-glucuronide and gemfibrozil-1-O-glucuronide (Baer et al., 2009; Tornio et al., 2014; Backman et al., 2016), in contrast to most other P450 enzymes. Accordingly, molecular modeling combined with in vitro experiments provide useful means to predict CYP2C8-involved DDI potential of drugs and their metabolites.

In conclusion, our clinical study revealed two DDIs with interesting characteristics and implications. First, in the clopidogrel-desloratadine and gemfibrozil-desloratadine interactions, both the perpetrator and victim compounds are glucuronide metabolites, highlighting the need to consider the properties of xenobiotic glucuronides with respect to evaluation of the risk of pharmacokinetic DDIs. Second, despite its complex metabolism, desloratadine could serve as a safe CYP2C8 probe substrate in clinical DDI studies owing to the CYP2C8-dependent 3-hydroxydesloratadine formation. It could be especially useful if the perpetrator also inhibits CYP3A4 or OATP1B1, because it differs from other recommended CYP2C8 probe substrates that are also dependent on CYP3A4 and OATP1B1. However, the suggested sequential metabolism and obligatory role of UGT2B10 prior to CYP2C8-mediated oxidation of desloratadine (Kazmi et al., 2015a) may complicate its DDI interpretation. Therefore, the applicability of desloratadine in populations with a high prevalence...
of desloratadine “poor metabolizer” phenotype requires further studies. Last, our study confirms that gemfibrozil is a stronger CYP2C8 inhibitor than clopidogrel, but they both can be used as model CYP2C8 inhibitors, as long as their DDI profile is taken into account in the study design and interpretation.

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Participated in research design: Ikkonen, Tornio, Neuvonen, P. J. Neuvonen, Niemi, Backman.
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Performed data analysis: Ikkonen, Tornio, Neuvonen, P. J. Neuvonen, Niemi, Backman.
Wrote or contributed to the writing of the manuscript: Ikkonen, Tornio, Neuvonen, P. J. Neuvonen, Niemi, Backman.

References

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