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ABSTRACT:
Muraglitazar and peliglitazar, two structural analogs differing by a methyl group, are dual peroxisome proliferator-activated receptor-α/γ activators. Both compounds were extensively metabolized in humans through acyl glucuronidation to form 1-O-β-acyl glucuronide (AG) metabolites as the major drug-related components in bile, representing at least 15% to 18% of the dose after oral administration. Peliglitazar AG was the major circulating metabolite, whereas muraglitazar AG was a very minor circulating metabolite in humans. Peliglitazar AG circulated at lower concentrations in animal species than in humans. Both compounds had a similar glucuronidation rate in UDP-glucuronic acid-fortified human liver microsomal incubations and a similar metabolism rate in human hepatocytes. Muraglitazar AG and peliglitazar AG were chemically synthesized and found to be similarly oxidized through hydroxylation and O-demethylation in NADPH-fortified human liver microsomal incubations. Peliglitazar AG had a greater stability than muraglitazar AG in incubations in buffer, rat, or human plasma (pH 7.4). Incubations of muraglitazar AG or peliglitazar AG in plasma produced more aglycon than acyl migration products compared with incubations in the buffer. These data suggested that the difference in plasma stability, not differences in intrinsic formation, direct excretion, or further oxidation of muraglitazar AG or peliglitazar AG, contributed to the observed difference in the circulation of these AG metabolites in humans. The study demonstrated the difficulty in doing risk assessment based on metabolite exposure in plasma because the more reactive muraglitazar AG would not have triggered a threshold of concern based on the recent U.S. Food and Drug Administration guidance on Metabolites in Safety Testing, whereas the more stable peliglitazar AG would have.

Introduction

Acyl glucuronidation is one of the major metabolic pathways for many carboxylic acid-containing drugs. 1-O-β-Acyl glucuronides, which are formed by UDP-glucuronosyltransferase (UGT) enzymes as the original isomer, can undergo a number of reactions including hydrolysis, rearrangement via acyl migration (Stachulski et al., 2006; Zhang et al., 2006; Xue et al., 2008), further metabolism (Kumar et al., 2002; Kochansky et al., 2005; Ogilvie et al., 2006), and reaction with nucleophiles (Sallustio and Foster, 1995; Akira et al., 2002; Bailey and Dickinson, 2003). The chemical reactivity of acyl glucuronides proceeds via two distinct pathways (Bailey and Dickinson, 2003; Stachulski et al., 2006; Corcoran et al., 2001): 1) direct displacement of the acyl residue with a nucleophile to produce an aglycon (hydrolysis product) and an acylated nucleophile; and 2) alternatively, migration of the acyl group around the sugar ring to yield 2-, 3-, and 4-acyl isomers. These isomers may undergo transient ring opening with concurrent formation of a reactive aldehyde group. Either of these pathways could lead to covalent binding to cellular proteins (Sallustio and Foster, 1995; Akira et al., 2002; Bailey and Dickinson, 2003). These pathways are effectively catalyzed at physiological pH (pH 7.4) and occur more rapidly under basic pH conditions (Xue et al., 2008). Acyl glucuronides are relatively stable under acidic conditions (pH 4–5). Careful acidification and cooling are required to stabilize acyl glucuronides in biofluids such as plasma, urine, and bile (Wang et al., 2006).

Muraglitazar and peliglitazar, oxybenzylglycine analogs (nonthiazolidinedione), are novel dual peroxisome proliferator-activated receptor-α/γ activators. The two compounds are structurally very sim-
ilar with the difference being the addition of a methyl group (Fig. 1). After oral administration to humans, both $^{14}$C-muraglitazar and $^{14}$C-peliglitazar underwent extensive conjugation, and the major portion of each radioactive dose was excreted in bile as acyl glucuronide metabolites in humans (Wang et al., 2006, 2010; Zhang et al., 2007b). The acyl glucuronide metabolites of muraglitazar and peliglitazar seemed to be stable in ex vivo plasma, urine, and bile under acidic and low temperature conditions because no significant amounts of acyl migration isomers were formed in plasma, urine, and bile of animals and humans that were acidified and stored at low temperature. In addition, the parent concentrations were not increased over time (Wang et al., 2006, 2010). The major circulating drug-related component (>90%) after oral administration in humans was muraglitazar; however, both peliglitazar and peliglitazar AG were major drug-related components (approximately equal concentrations) in humans after oral administration of peliglitazar. There was no protein co-valently bound radioactivity found in the protein pellet after extraction with organic solvents from plasma samples of humans with oral administration of either $^{14}$C-labeled compound.

The circulation of a metabolite, in general, would mainly depend on its formation, further metabolism, and direct clearance into urine and bile. For reactive metabolites such as acyl glucuronides, there could also be reactions to form various small metabolites or large molecular adducts. In this study we investigated the mechanistic response of the distributional difference among muraglitazar and peliglitazar, two structurally similar analogs, by studying formation, stability, excretion, and further metabolism of their glucuronide metabolites.

Materials and Methods

Materials. Muraglitazar, peliglitazar, and their $^{14}$C-labeled materials were synthesized at Bristol-Myers Squibb (Princeton, NJ). Their structures are shown in Fig. 1. The $^{14}$C-labeled materials had a radiochemical purity of >99%. Acetonitrile and trifluoroacetic acid were purchased from EM Scientific (Gibbstown, NJ). Pooled human liver microsomes ($n = 22$) were purchased from BD Biosciences (San Jose, CA). The human hepatocytes were acquired as freshly prepared cell suspensions from Lonza (Walkersville, MD). All other chemicals used were of reagent grade or better.

Synthesis of Peliglitazar AG and Muraglitazar AG. Muraglitazar AG and peliglitazar AG were synthesized basically following the procedure described previously (Kenny et al., 2004; Perrie et al., 2005).

Allyl glucuronate. To a solution of $\beta$-glucuronic acid (5.0 g, 25.7 mmol) in dimethyl formamide (50 ml) at 20°C was added 1.8-diazabicyclo[5.4.0]undec-7-ene (4.3 ml, 28.2 mmol). The mixture was stirred for 15 min, after which allyl bromide (2.8 ml, 30.8 mmol) was added. The reaction was stirred overnight, after which volatile compounds were removed in vacuo at 60°C. The crude residue was then purified by flash chromatography twice on a 120-g silica gel column with a continuous gradient from 0 to 20% methanol (MeOH) in CH$_2$Cl$_2$. The isolated material was then dissolved in MeOH (5 ml) and concentrated in vacuo. CH$_2$Cl$_2$ (40 ml) was added, and the solution was cooled to 0°C for 0.5 h. The precipitated solid was filtered off and washed with ice-cold CH$_2$Cl$_2$ and then dried in vacuo to give the allyl glucuronate (3.4 g, 57% yield, $\alpha$-$\beta$-anomer ratio at 2:1).

Peliglitazar $\beta$-anomer allyl ester. To a −5°C solution of peliglitazar (5.5 g, 10.4 mmol) and PPh$_3$ (2.72 g, 10.4 mmol) in tetrahydrofuran (THF) (40 ml) and dimethyl formamide (5 ml) was added disisopropyl azodicarboxylate (2.0 ml, 10.4 mmol). After 10 min, a solution of the allyl glucuronate (1.44 g, 6.15 mmol) in THF (10 ml) and dimethyl formamide (2.5 ml) was added slowly over 10 min using a syringe pump. After the addition was complete, the reaction solution had turned dark brown. The reaction was stirred at −5°C for 2 h, and the reaction was monitored by thin-layer chromatography until completion. The formation of product(s) was monitored by HPLC-MS analysis on a Luna C18 50 × 4.6 mm column with a 0 to 100% B linear gradient in 4 min with solvent A (10% MeOH in 10 mM ammonium acetate solution) and solvent B (90% MeOH in 10 mM ammonium acetate solution at a flow rate of 4 ml/min). Volatile compounds were removed in vacuo, and the crude residue was purified by flash chromatography twice on a 120-g silica gel column with a continuous gradient of 0 to 10% ethanol in CH$_2$Cl$_2$. The resulting material contained the desired $\alpha$- and $\beta$-anomers as major products. The $\alpha$- and

![Fig. 1. Metabolite profiles in 1 h human plasma samples after single oral administration of $^{14}$C-labeled muraglitazar or peliglitazar. The samples were analyzed by HPLC system II as described under Materials and Methods.](https://example.com/fig1.png)
β-anomers with a 2 to 3:1 ratio had been partially separated by the flash chromatography procedure described above, and the mixture was further purified by preparative HPLC. The preparative HPLC conditions used a Synergi Hydro-RP 80A column (250 × 21 mm 4 μm; Phenomenex, Torrance, CA) at room temperature with a flow rate of 25 ml/min and a mobile phase of 0.05% acetic acid (pH 3.3)-acetoneitrile (50:50, v/v) of a run time of 48 min at 278 nm. A portion of 35 mg was injected on the column each time. After careful removal of acetoneitrile in vacuo at room temperature, the remaining aqueous solution was extracted with ethyl acetate (300 ml), and the organic phase was concentrated and dried in vacuo to provide the α- and β-anomer esters as white powders. The β-anomer was recovered in a 10% yield.

Peliglitazar AG. To a 0°C solution of the β-anomer allyl ester (370 mg, 0.44 mmol) in THF (2.0 ml) was added (Ph₃P)₂Pd (59.2 mg, 0.051 mmol) followed by pyridine (35.2 μL, 0.44 mmol). The mixture was stirred at 0°C for 30 min, after which volatile compounds were immediately removed in vacuo to give the crude carboxylic acid. LC-MS using the previously described Luna C18 column method indicated the formation of desired product. The crude product was purified by previously described preparative HPLC on a Synergi Hydro-RP column. After removal of acetoneitrile, the remaining solution was lyophilized to give the desired acyl glucuronide

followed by pyrrolidine (35.2 MHz, DMSO-d₆) esters as white powders. The cleavage and at 5.42 ppm (J = 8.80 Hz, 1H), 7.08 (d, J = 8.80 Hz, 1H), 3.90 (d, J = 8.80 Hz, 1H), 3.73 (s, 1H), 3.60 (d, J = 8.80 Hz, 1H), 3.28 (d, J = 8.80 Hz, 1H), 3.20–3.16 (m, 2H), 3.20–3.27 (m, 1H), 3.35 (t, J = 8.80 Hz, 1H), 4.09–4.25 (m, 4H), 4.58 (dd, J = 8.80 Hz, 1H), 5.38 (d, J = 8.80 Hz, 1H), 5.42 (d, J = 7.70 Hz, 1H), 6.94 (dd, 4H), 7.01 (d, J = 8.80 Hz, 1H), 7.08 (d, J = 8.80 Hz, 1H), 7.26 (d, J = 8.80 Hz, 1H), 7.31 (d, J = 8.80 Hz, 1H), 7.43–7.55 (m, 3H), 7.91 (d, J = 7.15 Hz, 2H). The anomeric proton of the β-anomer was at 5.38 ppm (J = 8.85 Hz) for the minor rotational conformer and at 5.42 ppm (J = 7.70 Hz) for the major conformer. The downfield chemical shift and large coupling observed are typical for the β configuration.

The following resonance signals corresponded to the minor rotational conformer: 4.58 ppm (dd, 1H), 5.38 (J = 8.25 Hz, 7.31, d, J = 8.25 Hz, 2H), 7.01 (d, J = 8.25 Hz, 1H). 13C NMR (126 MHz, DMSO-d₆) δ ppm: 10.95 (s, 1C), 26.69 (s, 2C), 49.50 (s, 1C), 51.88 (br s, 1C), 56.47 (s, 1C), 67.29 (s, 1C), 72.98 (s, 1C), 73.52 (br s, 1C), 75.69 (s, 1C), 77.36 (s, 1C), 96.15 (br s, 1C), 115.10–115.47 (m, 2C), 115.65 (br s, 2C), 123.30–124.14 (m, 2C), 126.53 (s, 2C), 128.22 (s, 1C), 130.03 (s, 1C), 130.14 (s, 1C), 131.14 (s, 1C), 133.78 (s, 1C), 145.51 (s, 1C), 146.20 (s, 1C), 155.71 (s, 1C), 157.65 (br s, 1C), 158.86 (s, 1C), 159.47 (s, 1C), 169.30 (s, 1C), 169.62 (s, 1C), 172.61 (br s, 1C). The carbonyl signal at 169.30 corresponded to the minor rotational conformer.

Liver Microsomal and Hepatocyte Incubations. Incubations (250 μl) contained muraglitazar or peliglitazar (10 μM, 2.5 mM stock solution in 50:50, v/v, acetoneitrile-Tris buffer), human liver microsomes (1 mg/ml protein), UDPGA (2 mM), magnesium chloride (10 mM), and Tris buffer (100 mM, pH 7.4). Incubations were initiated by addition of a substrate at 37°C and quenched after 15 min by addition of 1 volume of acetonitrile to the incubation mixture. After centrifugation to remove the precipitated microsomal proteins, the clear supernatant (100 μl) was analyzed by LC-MS. HPLC system I consisted of a Waters 600 pump, a 717 autosampler, and a 99 photodiode array detector. The column used was a C18 YMC ODS-AQ reverse-phase column (4.6 × 150 mm, 3 μm) maintained at room temperature with a flow rate of 0.7 ml/min. The mobile phase A consisted of 5% acetonitrile in water containing 0.1% TFA and the mobile phase B consisted of 95% acetonitrile containing 0.1% TFA. A linear gradient was 10% to 100% B in 20 min and then held at 100% B for 10 min. The retention times were 12.1, 7.3, 15.4, and 8.6 min for muraglitazar, muraglitazar AG, peliglitazar, and peliglitazar AG, respectively. The HPLC was interfaced to a LQO mass spectrometer (Thermo Fisher Scientific, Waltham, MA) operated in the positive ionization mode to acquire full-scan LC-MS data with a mass scan range of 200 to 1000 Da. The percent metabolism was calculated on the basis of the ratio of peak areas of the metabolite versus metabolite plus the parent (metabolite/metabolize + parent) from the UV chromatogram (at 278 nm).

Hepatocyte incubations were performed with cells in suspension in 24-well tissue culture plates shaken at 90 rpm on an orbital shaker. The incubations were in Krebs-Henseleit buffer in a 5% CO₂/95% air atmosphere at 37°C with muraglitazar or peliglitazar (5 μM, 0.5 mM stock in 50:50 acetonitrile-potassium phosphate, v/v) and 1 × 10⁶ hepatocytes/ml. Incubation time was 1 h. The samples were quenched by addition of an equal volume of acetonitrile. The quenched samples were treated and analyzed by LC-MS with HPLC system I as described for the microsomal incubations.

Muraglitazar AG or peliglitazar AG at 20 μM was separately incubated for 15 min with human liver microsomes (2 mg/ml protein) in 1 ml of 50 mM sodium phosphate buffer with and without 1 mM NADPH. The samples were separated by HPLC system II using a Shimadzu LC-10AT system with the analytical column used for the microsomal incubation samples. The mobile phase B consisted of 50% of two solvents (A) (pH 7.4, 0.06% TFA in water) and solvent B (0.06% TFA in acetonitrile). The gradient consisted of the following steps: solvent B started at 5%, then increased linearly to 25% in 5 min, to 40% at 20 min, to 53% at 60 min, to 60% at 60 min, and to 90% at 65 min, held at 90% for 7 min, and then decreased to 5% at 75 min. The HPLC effluent was 1 ml/min. The quantities of muraglitazar and peliglitazar AGs were estimated by HPLC separation with UV detection at 278 nm. The metabolites were identified by LC-MS as described in the supplemental data.

Study Subjects, Dosing, and Sample Collection. All animal housing and care conformed to the standards recommended by the Guide for the Care and Use of Laboratory Animals (Institute of Laboratory Animal Resources, 1996). Animal rooms were maintained on a 12-h light/dark cycle. The human study was performed in accordance with the principles of the Declaration of Helsinki and its amendments, and the study protocol was approved by the Institutional Review Board and Radiation Safety Committee at the investigational site. All subjects were in good health and gave written, informed consent to participate in the study.

Male CD-1 mice (n = 5, 25–35 g), male Sprague-Dawley rats (n = 3, 250–280 g), and male cynomolgus monkeys (n = 3, 3–5 kg) were fasted for approximately 8 h before dosing. Each animal received an oral gavage dose of 30 mg/kg (300 μC/kg, 5 ml/kg) for mouse, 15 mg/kg (150 μC/kg, 5 ml/kg) for rat, and 3 mg/kg (30 μC/kg, 1 ml/kg) for monkey of [14C]muraglitazar.
dissolved in PEG−400. Human studies were conducted as described previously (Wang et al., 2006). Four healthy male subjects, aged 18 to 45 years, each received a single dose of [14C]muraglitazar (20 mg) or [14C]peliglitazar (10 mg) containing approximately 100 μCi of radioactivity as an oral solution in PEG−400 after at least an 8-h overnight fast.

Blood samples (terminal for mouse and rat and via the cephalic vein for monkey and direct venipuncture using Vacutainers for human) were collected at 1, 4, 12, and 24 h using K2EDTA as the anticoagulant. Plasma was prepared by centrifugation at approximately 1000g for 15 min at 4°C. Acetic acid was added to plasma to a final concentration of 5% (v/v, 0.83 M) immediately after processing. All plasma samples were frozen and stored at −20°C.

Sample preparation and analysis. Pooled plasma samples were prepared separately by mixing an equal volume (0.2−0.5 ml) of plasma sample from each subject. Portions (0.5−1 ml) of the pooled plasma samples were extracted by addition of a mixture of 1 volume of methanol and 3 volumes of acetonitrile and mixed on a Vortex mixer. The mixtures were centrifuged at 2000g for 10 min, and then the supernatants were transferred into a polypropylene centrifuge tube. The extraction was repeated two more times, and all supernatants were combined. The recovery of radioactivity in the supernatant after extraction was quantitative. The plasma protein pellets were digested in 1 M NaOH for 12 h before being neutralized with 1 M HCl and counted for radioactivity in 15 ml of Ecolite scintillation fluid. The combined supernatants were concentrated under a stream of nitrogen, the residues were then reconstituted in 0.2 to 0.5 ml of a solution of 70% of HPLC mobile phase A (0.06% TFA in water) and 30% mobile phase B (0.06% TFA in acetonitrile), vortexed, and centrifuged at 2000g for 10 min, and 100 μl of the supernatant was used for the HPLC analysis. Metabolites in plasma were analyzed by HPLC system II using a Shimadzu LC-10AT system as described in the preceding section. HPLC effluent (1 ml/min) was collected in a Deepwell LumaPlate-96 (PerkinElmer Life and Analytical Sciences, Waltham, MA) at 0.26 min. The plates were dried with a SpeedVac (Thermo Fisher Scientific) and counted for 10 min with a Top-Count analyzer (PerkinElmer Life and Analytical Sciences) to quantify radioactivity. Biotransformation profiles were prepared by plotting the resulting net counts per minute values versus time after injection. Radiochromatograms were reconstructed from the TopCount data using Microsoft Excel software.

Stability studies in incubations in buffer and plasma. A portion of 0.75 ml of 1 M phosphate buffer at pH 7.4 and 14.1 ml of human or rat plasma were mixed with 0.15 ml of 5 mg/ml 1-M phosphate buffer at pH 7.4 and 14.1 ml of water were mixed with 0.15 ml of 5 mg/ml 1-O-b-acyl glucuronide stock solution. The remaining steps were the same as those for the incubations for buffered plasma.

1-O-β-Acyl glucuronides and their isomers as well as aglycons were separated isocratically by HPLC system III using a mobile phase composed of 65:35 (v/v) acetonitrile-water containing 0.05% formic acid at a flow rate of 0.3 ml/min. The separation was performed using a Luna C18(2) analytical column (3 × 150 mm, 3 μm; Phenomenex, Torrance, CA) operated at 25°C. The injection volume was 5 μl, and the run time was 12.0 min. Under these conditions, the retention time was 6.17, 7.04, 7.55, 8.13, 8.35, 8.82, and 15.08 min, respectively, for isomer 1, isomer 2, muraglitazar AG, isomer 3, isomer 4, isomer 5, and muraglitazar. The retention time was 7.51, 8.62, 9.55, 10.25, 11.05, and 14.80 min, respectively, for isomer 1, isomer 2, peliglitazar AG, isomer 3, isomer 4, and peliglitazar. The HPLC effluent was monitored at 278 nm and analyzed by a LTQ mass spectrometer (Thermo Fisher Scientific).

Results

Table 1 and Fig. 2 show the exposures of peliglitazar and peliglitazar AG in the plasma of mouse, rat, monkey, and human. The exposure multiples of the maximum concentration (Cmax) and AUC values for the parent compound in mouse, rat, and monkey were >21. The exposure multiples of Cmax and AUC values for the major glucuronide metabolite, peliglitazar AG, were 3.8 to 6.4 in the mouse, 1 to 2 in monkey, and <0.5 in rat from the ADME studies. The projected exposure multiples of Cmax and AUC values for the metabolite were higher (all were >1) in mouse, rat, and even monkey with linearly scaled doses for the term toxicological studies. However, the exposure multiples of Cmax and AUC values for the metabolite were still <1 with the projected carcinogenicity testing doses. Overall, peliglitazar AG exposure multiples of Cmax and AUC in toxicological species were marginal even at the projected high toxicological doses.

Figure 1 shows that peliglitazar AG circulated in humans as a major metabolite, whereas muraglitazar AG had little circulation in humans in the 1-h plasma samples. Radioactivity profiles of human plasma at 4, 12, and 24 h also showed that peliglitazar AG but not muraglitazar AG was a significant circulating metabolite in humans. There were minimal to no acyl isomers detected in the plasma samples from subjects after administration of

<table>
<thead>
<tr>
<th>TABLE 1</th>
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<tbody>
<tr>
<td>Estimated exposures of peliglitazar and its acyl glucuronide in mice, rats, monkeys, and humans after a single oral dose of peliglitazar</td>
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<table>
<thead>
<tr>
<th>Study</th>
<th>Analyte</th>
<th>Species</th>
<th>Doses (mg/kg)</th>
<th>Conc. 't at 1 h (μM)</th>
<th>Conc. Exposure Multiples</th>
<th>AUC Exposure Multiples (μM h)</th>
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<tbody>
<tr>
<td>Radiolabeled ADME</td>
<td>Peliglitazar</td>
<td>Mouse</td>
<td>30</td>
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<td>4.51</td>
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<td></td>
<td></td>
<td>Monkey</td>
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<td>3.64</td>
<td>24</td>
<td>15.5</td>
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<tr>
<td>Radiolabeled ADME</td>
<td>Peliglitazar AG</td>
<td>Mouse</td>
<td>30</td>
<td>1.18</td>
<td>6.4</td>
<td>4.72</td>
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<tr>
<td></td>
<td></td>
<td>Rat</td>
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<td>0.4</td>
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<tr>
<td></td>
<td></td>
<td>Human&lt;sup&gt;a&lt;/sup&gt;</td>
<td>10</td>
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<td>1</td>
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<tr>
<td>Long-term toxicity studies&lt;sup&gt;b&lt;/sup&gt;</td>
<td>Peliglitazar AG</td>
<td>Mouse</td>
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<td>9.8</td>
<td>54</td>
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<td></td>
<td></td>
<td>Rat</td>
<td>300</td>
<td>1.4</td>
<td>7.8</td>
<td>4.2</td>
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<tr>
<td>Carcinogen studies&lt;sup&gt;c&lt;/sup&gt;</td>
<td>Peliglitazar AG</td>
<td>Mouse</td>
<td>25</td>
<td>0.28</td>
<td>1.6</td>
<td>1.21</td>
</tr>
</tbody>
</table>

ADME, absorption, distribution, metabolism, and excretion.

<sup>a</sup> Exposure multiples were calculated by dividing Cmax or AUC values in animals by that in humans.

<sup>b</sup> Data are milligrams per subject.

<sup>c</sup> Concentration is estimated from the relative distribution of the parent or acyl glucuronide metabolites in the plasma, the total concentration of radioactivity, and the specific activity of the administered drug; AUC<sub>Exposure</sub> is estimated from the concentration in plasma at limited time points (1, 4, and 12 h) using the trapezoidal rule.

<sup>d</sup> Doses are the highest dose from the 6-month rat, 3-month mouse, and 1-year monkey toxicology studies.

<sup>e</sup> Concentration and AUC values are scaled linearly with regard to dose values in ADME studies.

<sup>f</sup> Highest projected doses for the carcinogenicity studies.
In addition, the plasma protein pellets after extraction with the organic solvent contained no detectable levels of radioactivity. Together, the low levels of isomerization and protein covalent binding suggest ex vivo stabilization of the plasma samples by addition of acetic acid and low levels of total plasma radioactivity. Overall, muraglitazar AG represented approximately 4% of the muraglitazar AUC_0–24h, whereas...
peliglitazar AG represented 24% of the peliglitazar AUC0-24h. With concentrations of peliglitazar AG of 102, 48.8, and 21.3 ng/ml at 1, 4, and 12 h postdose, the estimated half-life of the peliglitazar AG was approximately 3 h based on the limited data points (Wang et al., 2010). Estimation of half-life for muraglitazar AG was not possible. Although both muraglitazar and peliglitazar underwent extensive conjugation metabolism to form AGs as the major metabolite, only peliglitazar AG was the major circulating component after oral administration of the 14C-labeled drug. A similar distributional difference was observed in monkeys as well as in rats after oral administration of 14C-labeled muraglitazar or peliglitazar (data not shown).

Table 2 shows the glucuronidation rates of muraglitazar and peliglitazar in human liver microsomes and hepatocytes and biliary/urinary excretion of acylglucuronide metabolites in humans after oral administration of 14C-labeled compounds. At clinically relevant concentrations, 5 μM in hepatocytes and 10 μM in human liver microsomes (Zhang et al., 2006; Wang et al., 2010), muraglitazar and peliglitazar showed similar glucuronidation rates in incubations with both UDPGA-fortified human liver microsomes and human hepatocytes. In humans, muraglitazar and peliglitazar AGs showed similar elimination profiles with at least 15 to 16% of dose excreted in the bile over the 3- to 8-h postdose collections and 0.2 to 0.8% of the dose in the urine over 0- to 168-h collections after oral administration of 14C-labeled muraglitazar or peliglitazar (data not shown).

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Table 3 shows the stability of muraglitazar AG and peliglitazar AG in incubation in buffer or human plasma. Figure 3 shows distinct concentration-time profiles of muraglitazar AG and peliglitazar AG degradation in the buffer and rat or human plasma samples. Muraglitazar AG degraded in a phosphate buffer or human plasma at approximately 7 to 8 times faster than peliglitazar AG by hydrolysis to aglycons and to a lesser extent by acyl migration to positional isomers. Muraglitazar and peliglitazar AGs degraded approximately 3 to 5 times faster in human plasma than in the buffer (Table 3). Both glucuronides had degradation that was faster in rat plasma than in buffer but was different from that in human plasma (Fig. 3). Good separation of the positional isomers of muraglitazar AG and peliglitazar AG and their degradation products was achieved by simple reverse-phase HPLC using an isocratic elution as described by Xue et al. (2008). The hydrolysis rates for formation of muraglitazar and peliglitazar were 5 to 10 times faster in plasma than in buffer (Figs. 4 and 5). In the incubation with muraglitazar AG in human plasma, the acyl migration isomers 3 and 4 were formed and then quickly hydrolyzed to form the aglycon (Fig. 4B); however, both the isomers and the aglycon were formed slowly in the buffer. In addition, the acyl isomers degraded to aglycon at a slower rate than that in plasma (Fig. 4A). Similar degradation profiles were observed with peliglitazar AG but at a much slower rate than with muraglitazar AG (Fig. 5).

The oxidative metabolism of muraglitazar AG and peliglitazar AG was investigated in human liver microsomes in the presence of NADPH. Multiple hydroxylated metabolites and the O-demethylated metabolite of muraglitazar AG were observed in the incubation starting with muraglitazar and peliglitazar AGs, and metabolite identification is shown in the supplemental data. The glucuronide oxidation rates were estimated to be 22 and 24 pmol/(min · mg), respectively, for muraglitazar AG and peliglitazar AG.

### Discussion

Fig. 6 illustrates the potential factors that would be expected to affect the circulating levels of an acyl glucuronide metabolite include

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**TABLE 2**

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<thead>
<tr>
<th>In Vitro Glucuronidation Rates&lt;sup&gt;a&lt;/sup&gt;</th>
<th>Glucuronide Excretion&lt;sup&gt;b&lt;/sup&gt;</th>
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<tbody>
<tr>
<td>HLM</td>
<td>Hepatocytes&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>---</td>
<td>---</td>
</tr>
<tr>
<td>Muraglitazar</td>
<td>33 pmol/(min · mg protein)</td>
</tr>
<tr>
<td>Peliglitazar</td>
<td>32 pmol/(min · mg protein)</td>
</tr>
</tbody>
</table>

<sup>a</sup>The substrate concentration was 5 and 10 μM, respectively, for hepatocyte and microsomal incubations.

<sup>b</sup>Excretion of muraglitazar AG and peliglitazar AG in humans after oral administration of 14C-labeled compounds (20 mg/subject, 100 μCi). Urinary elimination represented 2.13 and 1.24% of dose and biliary elimination represented 39.9 and 24.4% of dose, respectively, for muraglitazar and peliglitazar.

<sup>c</sup>The value was the average of formation rates within the linear ranges (1/2t₁/₂ was calculated from initial rates of glucuronide disappearance (1/2t₁/₂ = 0.693/slope)). The value was the average of formation rates within the linear ranges (<3 h).

**TABLE 3**

<table>
<thead>
<tr>
<th>Compounds</th>
<th>Products</th>
<th>Buffer</th>
<th>Human Plasma</th>
</tr>
</thead>
<tbody>
<tr>
<td>Disappearance of glucuronide (t₁/₂, h)&lt;sup&gt;a&lt;/sup&gt;</td>
<td>Muraglitazar AG</td>
<td>N.A.</td>
<td>1.23</td>
</tr>
<tr>
<td>Peliglitazar AG</td>
<td>N.A.</td>
<td>10.1</td>
<td>2.3</td>
</tr>
<tr>
<td>Aglycon formation rates [μg/(ml · h)]&lt;sup&gt;b&lt;/sup&gt;</td>
<td>Muraglitazar AG</td>
<td>3.5</td>
<td>16.7</td>
</tr>
<tr>
<td>Peliglitazar AG</td>
<td>0.5</td>
<td>5.9</td>
<td></td>
</tr>
</tbody>
</table>

<sup>a</sup>T₁/₂ was calculated from initial rates of glucuronide disappearance (t₁/₂ = 0.693/slope).

<sup>b</sup>The buffer was 50 mM sodium phosphate, pH 7.4. Plasma samples were buffered with 50 mM phosphate, pH 7.4.
formation rate, biliary and renal excretion, further metabolism, tissue distribution, and reactions with small and large molecules. The disproportional exposures to pelaglitazar AG across species and the striking differences in exposures to AGs of pelaglitazar and muraglitazar in humans lead us to evaluate factors that may influence AG exposures for these two compounds. Intrinsic formation in vitro and direct excretion for acyl glucuronide metabolites in vivo were found to be similar for muraglitazar and pelaglitazar. Incubations of muraglitazar and pelaglitazar in UDPGA-fortified human liver microsomes or in hepatocytes generated the acyl glucuronides as the major metabolites at similar formation rates. Examination of the acyl glucuronide excretion profile revealed that the acyl glucuronide was excreted as the major metabolite in the bile of humans and to the same extent. Urinary excretion was a minor clearance pathway for both compounds. There was no tissue accumulation or other major differences in tissue distribution patterns in rats after oral administration of [14C]muraglitazar or [14C]pelaglitazar (data not shown). These observations do not explain the in vivo data, which showed that muraglitazar AG had a much lower exposure than pelaglitazar AG. Thus, the difference in chemical stability or further metabolism of their glucuronide metabolites may have contributed to their very different circulation profiles in humans.

In incubations in pH 7.4 buffer at 37°C, muraglitazar AG degraded >7 to 8 times faster than pelaglitazar AG (Fig. 3; Table 3). In incubations of muraglitazar and pelaglitazar AGs in human plasma, the oxidation of both muraglitazar AG and pelaglitazar AG was similarly slow and cannot be used to explain their differences in the circulation.

In incubations in pH 7.4 buffer at 37°C, muraglitazar AG degraded >7 to 8 times faster than pelaglitazar AG (Fig. 3; Table 3). In incubations of muraglitazar and pelaglitazar AGs in human plasma,
migration isomers were formed to lesser amounts, and these incubations generated aglycons as the dominant products (Figs. 4B and 5B). In addition, the acyl migration isomers quickly degraded to the aglycon in buffer. For comparison with incubations in buffer (Figs. 4A and 5A), the aglycon was formed 3 to 5 times more slowly than in human plasma. In addition, more acyl isomers were formed and then slowly degraded to the aglycon in the buffer. The greater formation of aglycon in plasma may be due to hydrolysis catalyzed by β-glucuronidases and esterases or an unknown hydrolyase. The differences in the plasma stability of muraglitazar and peliglitazar AGs could be governed by the steric hindrance of nearby groups to the sugar aglycon C–O–C = O bond for hydrolysis or migration. The additional methyl group that is near the sugar aglycon bond in peliglitazar compared with muraglitazar might provide the steric hindrance to prevent hydrolysis of peliglitazar AG. This hypothesis of a steric effect on hydrolysis may be tested with model compounds in the future. Differences in the chemical stability of structurally related acyl glucuronides have been well documented (Spahn-Langguth and Benet, 1992; Bailey and Dickinson, 2003). Even stereo isomers (R- versus S-) of acyl glucuronides have been shown to have a 2-fold difference in reactivity (Fenselau, 1994; Akira et al., 2000; Hasegawa et al., 2001; Mortensen et al., 2001). Compared with literature first-order degradation half-lives in a pH 7.4 aqueous buffer of AGs of tolmetin (0.26 h), probenecid (0.4 h), diclofenac acid (0.51 h), R-naproxen (0.92 h), salicylic acid (1.3 h), S-naproxen (1.8 h), ibuprofen (3.3 h), bilirubin (4.4 h), mefenamic acid (16.5 h), gemfibrozil (44 h), and valproic acid (79 h) (Ebner et al., 1999; Shipkova et al., 2003), and dabigatran (1 h) (Ebner et al., 2010), muraglitazar AG (1.23 h) and peliglitazar AG (10.1 h) showed moderate aqueous buffer stability. However, reactivity evaluation in buffer was not sufficient because both muraglitazar and peliglitazar AGs showed approximately 4-fold additional instability in plasma compared with buffer. This additional degradation of muraglitazar AG and peliglitazar AG is consistent with a catalytic hydrolysis by an unknown hydrolyase for acyl glucuronides in plasma samples, which could be the reason for the inconsistent correlation between buffer degradation stability and plasma exposures of various acyl glucuronides (Ebner et al., 2010). Although the reactivity in plasma may not completely correlate with chemical stability either, as demonstrated in this study, the real exposure to an acyl glucuronide metabolite in animals or humans is a dynamic process and difficult to assess, especially when one is dealing with a metabolite that had a different degradation rate in plasma samples of different species. This study provides an excellent example to show that muraglitazar glucuronidation as a major metabolic pathway resulted in a minor circulating metabolite due to chemical degradation in humans.

Another potential difference in disposition of AGs that could lead to different circulating levels is further metabolism. Multiple oxidation metabolites of muraglitazar and peliglitazar AGs were identified (as described in the supplemental data). The extent of oxidative metabolism was similar for both muraglitazar AG and peliglitazar AG. Therefore, the elimination of muraglitazar and peliglitazar AGs may partially depend on oxidation of the AGs themselves, but that does not appear to explain the differential exposure seen in humans. Therefore, although other factors such as differential transport of AGs out of the hepatocytes into blood may also play a role, the chemical stability differences certainly would be expected to contribute to the observed exposure differences between muraglitazar AG and peliglitazar AG.

Acknowledgments

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Authorship Contributions

Participated in research design: D. Zhang, Raghavan, Wang, Xue, Li, Humphreys, and Cheng.

Conducted experiments: D. Zhang, Raghavan, Wang, and Obermeier.


Performed data analysis: D. Zhang, Raghavan, Wang, Obermeier, Ramanathan, Li, Yang, and Humphreys.

Wrote or contributed to the writing of the manuscript: D. Zhang, Ramanathan, Yang, and Humphreys.

References


Plasma stability-dependent circulation of acyl glucuronide metabolites in humans. How circulating metabolite profiles of muraglitazar and peliglitazar can lead to misleading risk assessment

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Pharmaceutical Candidate Optimization (DZ, NR, LW, YX, MO, WL, RR, ZY, WGH), Discovery Chemistry (SC, ST, HZ, PTC). Bristol-Myers Squibb Research and Development, Princeton, NJ 08543

Supplemental data: Oxidation of glucuronide metabolites of muraglitazar and peliglitazar in human liver microsomes

Muraglitazar AG or peliglitazar AG at 20 μM was separately incubated for 15 min with human liver microsomes (2 mg/mL protein) in 1 mL of 50 mM sodium phosphate buffer with and without 1 mM NADPH. The reaction was quenched by adding one volume of ice-cold acetonitrile containing 2% acetic acid following by centrifugation at 2000xg for 10 min. An aliquot of 100 μL was injected to LC/UV/MS/MS analysis. The samples were analyzed by HPLC system II using a Shimadzu LC-10AT system (Shimadzu Scientific Instruments, Kyoto, Japan) and LC/MS analyses were performed using the analytical column used for the microsomal incubation samples. The mobile phase consisted of two solvents: A) 0.06% TFA in water and B) 0.06% TFA in acetonitrile. The gradient consisted of the following steps: Solvent B started at 5%, then linearly increased to 25% at 5 min, to 40% at 20 min, to 53% at 60 min, to 60% at 63 min, to 90% at 65 min, held at 90% for 7 min, and then decreased to 5% at 75 min. HPLC effluent (1 mL/min) was monitored at 278 nm. The HPLC eluent was partially diverted to a LTQ mass spectrometer (ThermoFisher, San Jose, CA). Full LC/MS (Scan range of 200-1000 Da) and MS/MS spectra were collected.
Figures 1S and 2S show oxidative metabolite profiles of incubations of muraglitazar AG and peliglitazar AG in human liver microsomes in the presence of NADPH. Multiple hydroxylated metabolites and O-demethylated metabolite of muraglitazar and peliglitazar AGs were observed in the incubation starting with muraglitazar AG or peliglitazar AG. After incubations in human liver microsomes, the major components were still the starting materials (muraglitazar and peliglitazar AGs) and the incubations also led to hydrolysis of muraglitazar and peliglitazar AGs (Figures 1S and 2S). Table 1S shows mass spectrometric characterization of these oxidative metabolites. Metabolites M1, M2, and M3 had a molecular ion of m/z 709 from muraglitazar AG and m/z 723 from peliglitazar AG, and respective fragmentation ions at m/z 533 and 547 (loss of 176), which are consistent with hydroxylation products of muraglitazar AG and peliglitazar AG. Although these metabolites could be acyl migration isomers of one metabolite, more likely they are different metabolites with the hydroxyl group at different sites based retention times. Metabolite M4 had a molecular ion of m/z 679 from muraglitazar AG and m/z 693 from peliglitazar AG, and respective fragmentation ions at m/z 503 and 517 (loss of 176), which are consistent with demethylation products of muraglitazar AG and peliglitazar AG. Metabolites M5, M6, and M7 had a molecular ion of m/z 533 from muraglitazar AG and m/z 547 from peliglitazar AG, which are consistent with hydroxylated products of muraglitazar and peliglitazar. Metabolite M8 had a molecular ion of m/z 503 from muraglitazar AG and m/z 517 from peliglitazar AG, which are consistent with demethylated products of muraglitazar and peliglitazar. Therefore, in addition to these oxidative metabolites of glucuronide, multiple hydroxylated metabolites and an O-demethylated metabolite of muraglitazar were also observed in the incubation.
Without NADPH, no oxidative metabolites of muraglitazar, peliglitazar, muraglitazar AG, or peliglitazar AG were observed. In the incubations without NADPH, glucuronide isomers of muraglitazar or peliglitazar (e.g. the peak after the starting material peaks in Figures 1S and 2S) were observed. The muraglitazar or peliglitazar oxidative metabolites could be formed from muraglitazar or peliglitazar that resulted from hydrolysis of muraglitazar AG or peliglitazar AG or from hydrolysis of oxidized muraglitazar AG or peliglitazar AG. Very similar degradation profiles were observed for the incubation of muraglitazar AG or peliglitazar AG in the human liver microsomes (Figures 1S and 2S).

Figure 3S shows oxidation pathways as well as hydrolysis and isomerization of muraglitazar AG or peliglitazar AG in the *in vitro* incubations. In the incubations in the presence of NADPH, the oxidized glucuronides could have only been formed by oxidation of the acyl glucuronides since UDPGA was not present in the incubations. There are several reports in the literature on the oxidative metabolism of AGs including diclofenac (Kumar et al., 2002), MRL-C (Kochansky et al., 2007), bilirubin (Crawford et al., 1992), valproic acid (Tang et al., 1996), and gemfibrozil (Ogilvie et al., 2006). In humans, in addition to muraglitazar AG and peliglitazar AG, oxidized (hydroxylated and O-demethylated) acyl glucuronides were also metabolites of muraglitazar and peliglitazar following oral administration (Wang et al., 2006; Wang et al., 2010).

References:


**Supplement Figures**

**Figure 1S.** UV and ion chromatograms of oxidative metabolites of muraglitazar AG formed in HLM incubations in the presence of NADPH.

**Figure 2S.** UV and ion chromatograms of oxidative metabolites of peliglitazar AG formed in HLM incubations in the presence of NADPH.

**Figure 3S.** Hydrolysis, isomerization and metabolic pathways of muraglitazar AG and peliglitazar AG.
Table 1S. LC/MS/MS characterization of oxidative metabolites of muraglitazar AG and peliglitazar AG in human liver microsomes in the presence of NADPH

<table>
<thead>
<tr>
<th>Metabolite</th>
<th>Muraglitazar AG</th>
<th>Peliglitazar AG</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>T_R (min)</td>
<td>MH⁺ m/z</td>
</tr>
<tr>
<td>M1</td>
<td>25.25</td>
<td>709</td>
</tr>
<tr>
<td>M2</td>
<td>27.15</td>
<td>709</td>
</tr>
<tr>
<td>M3</td>
<td>29.12</td>
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<td>M4</td>
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<td>M7</td>
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<td>M8</td>
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<td>503</td>
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<tr>
<td>Aglycone</td>
<td>60.35</td>
<td>517</td>
</tr>
<tr>
<td>Starting material</td>
<td>41.38</td>
<td>693</td>
</tr>
</tbody>
</table>
Fig 1S

Channel A
UV: 278
Mura_glu
HLM_NADPH

Muraglitazar glucuronide

Hydroxylated muraglitazar glucuronides

O-Demethylated muraglitazar glucuronide

NL: 7.96E2
m/z = 532.50-533.50
ITMS + c ESI Full ms2
709.00@20.00
Mura_glu_HLM_NADPH

NL: 4.24E3
m/z = 502.50-503.50
ITMS + c ESI Full ms2
679.00@20.00
Mura_glu_HLM_NADPH
Fig 2S

Channel A
UV: 278 nm
Peli_glu
HLM_NADPH

Hydroxylated peliglitazar glucuronides

- NL: 1.44E2
- m/z= 546.50-547.50
- ITMS + c ESI Full ms2
- 723.00@20.00
- peli_glu_HLM_NADPH

O-Demethylated peliglitazar glucuronide

- NL: 2.12E3
- m/z= 516.50-517.50
- ITMS + c ESI Full ms2
- 693.00@20.00
- peli_glu_HLM_NADPH
Fig 3S

2,3,4-O-glucuronide

Acyl migration

1-O-β-glucuronide

\[ \text{UGT/UDPGA Hydrolysis} \]

muraglitazar or peliglitazar

\[ \text{P450 Hydrolysis} \]

M1, M2, M3 (hydroxy glucuronide)

\[ \text{P450} \]

M5, M6, M7 (hydroxy metabolite)

M4 (O-demethyl glucuronide)

\[ \text{Hydrolysis} \]

M8 (O-demethyl metabolite)

\[ R = H \text{ for muraglitazar and } R = \text{CH}_3 \text{ for peliglitazar} \]