PREDICTION OF IN VIVO DRUG-DRUG INTERACTIONS BETWEEN TOLBUTAMIDE AND VARIOUS SULFONAMIDES IN HUMANS BASED ON IN VITRO EXPERIMENTS

KANJI KOMATSU, KIYOMI ITO, YUKIKO NAKAJIMA, SHIN-ICHI KANAMITSU, SUSUMU IMAOKA, YOSHIHIKO FUNAE, CAROL E. GREEN, CHARLES A. TYSON, NORIAKI SHIMADA, AND YUICHI SUGIYAMA

Graduate School of Pharmaceutical Sciences, The University of Tokyo, Tokyo, Japan (K.K., Y.N., S.K., Y.S.); School of Pharmaceutical Sciences, Kitasato University, Tokyo, Japan (K.I.); Laboratory of Chemistry, Osaka City University Medical School, Osaka, Japan (S.I., Y.F.); Toxicology Laboratory, SRI International, Menlo Park, California (C.E.G., C.A.T.); Research and Development Division, Daiichi Pure Chemicals Co., Ltd., Osaka, Japan (N.S.)

(Received July 26, 1999; accepted January 10, 2000)

This paper is available online at http://www.dmd.org

ABSTRACT:

Drug-drug interactions between tolbutamide and sulfonamides have extensively been reported. We attempted to predict the in vivo interaction between tolbutamide and sulfonamides from the in vitro metabolic inhibition studies. The inhibition constant \( K_i \) was derived from the inhibitory effects of eight sulfonamides (sulfaphenazole, sulfadiazine, sulfamethizole, sulfisoxazole, sulfamethoxazole, sulfapyridine, sulfadimethoxine, and sulfamonemethoxine) on tolbutamide metabolism. We found that the inhibitory effect of sulfaphenazole was greatest among the eight sulfonamides examined. Furthermore, the contribution of each P450 enzyme to tolbutamide metabolism was predicted by coadministration of sulfaphenazole and sulfamethizole, the main enzyme involved was CYP2C9. The \( K_i \) values of several sulfonamides were comparable between human liver microsomes and recombinant CYP2C9. The maximum unbound plasma concentration of sulfonamides in the portal vein was calculated from literature data on the pharmacokinetics of sulfonamides. Using the \( K_i \) values obtained from in vitro inhibition studies, the degree of increase in tolbutamide area under the plasma concentration-time curve (AUC) was predicted. About 4.8- and 1.6-fold increases in tolbutamide AUC were predicted by coadministration of sulfaphenazole and sulfamethizole, respectively, which agreed well with the reported increases in humans. Furthermore, the increase in tolbutamide AUC by coadministration of sulfadiazine, sulfisoxazole, and sulfamethizole was predicted to be 1.5- to 2.6-fold, although the corresponding in vivo effects have not been reported. It is concluded that some of these sulfonamides have to be carefully coadministered with CYP2C9 substrates such as tolbutamide although coadministration of sulfaphenazole needs the greatest care.

Drug-drug interaction may cause serious side effects by raising the blood concentration of a drug whose metabolism is inhibited by coadministered drug. It is, therefore, important to predict any change in drug disposition caused by a drug-drug interaction. There are many reports of the prediction of in vivo drug disposition in humans based on animal experiments (Iwatsubo et al., 1996, 1997). However, because of species differences in metabolic enzymes, not only the metabolic activity but also metabolic pathway of drugs in animals may be different from those in humans. Therefore, predicting in vivo drug disposition in humans from animal data may sometimes lead to incorrect results. Recently, human liver samples and recombinant enzymes have become readily available, and prediction based on in vitro experiments is becoming more important (Iwatsubo et al., 1997; Ito et al., 1998a).

In this study, in vivo drug-drug interactions involving tolbutamide metabolism in humans were predicted based on in vitro studies using human liver microsomes and recombinant cytochrome P450 (CYP) 1

1 Abbreviations used are: CYP, cytochrome P450; AUC, area under the plasma concentration-time curve; Fa, fraction absorbed from the intestinal tract; fp, unbound fraction in plasma; \( V_{max} \), maximum metabolic rate in circulating blood; \( I_u \), unbound concentration of inhibitor; \( k_{so} \), absorption rate constant; \( V_{max} \), maximum metabolic rate.

\[ K_i \]

\[ I_u \]

\[ V_{max} \]

\[ k_{so} \]

\[ V_{max} \]

Send reprint requests to: Yuichi Sugiyama, Graduate School of Pharmaceutical Sciences, University of Tokyo, 7-3-1 Hongo, Bunkyo-ku, Tokyo 113-0033, Japan. E-mail: sugiyama@seizai.i.u-tokyo.ac.jp

475
data on both drugs reported in literature (Ito et al., 1998b). In this study, we used sulfaphenazole and seven other commercially available sulfonamides (sulfadiazone, sulfamethizole, sulfisoxazole, sulfamethoxazole, sulfapyridine, sulfadimethoxine, and sulfamonomethoxine) to investigate the inhibitory effects on tolbutamide hydroxylation in a series of in vitro experiments. Furthermore, the contribution of each P450 enzyme to in vivo tolbutamide metabolism was estimated using physiological amounts of recombinant P450 enzymes. Taking the pharmacokinetic feature of each sulfonamide into consideration, we predicted the degree of increase in tolbutamide AUC caused by coadministration of sulfonamides from the in vitro experiments using microsomes obtained from human livers and CYP2C9-expressed lymphoblastoid cells.

Materials and Methods

Chemicals and Reagents. Tolbutamide, diethyl ether, sulfanomethoxine, sulfamethoxazole, magnesium chloride, hydrochloric acid, and dipotassium hydrogen phosphate were purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan). Hydroxytolbutamide was kindly provided by Daiichi Pure Chemicals, Co. Ltd. (Tokyo, Japan). NADP, glucose 6-phosphate, and glucose 6-phosphate dehydrogenase were obtained from Boehringer Mannheim (Mannheim, Germany). Sulfaphenazole, sulfisoxazole, sulfadiazine, and sulfapyridine were purchased from Sigma Chemical Co. (St. Louis, MO). Methanol of HPLC grade was purchased from Wako Pure Chemical Industries, Ltd. All other chemicals were of reagent grade.

Human Liver Microsomes and Recombinant CYP Enzymes. Human liver microsomes obtained from ten donors (six males and four females; 31–57 years old; H-19, H-35, H-36, H-38, H-50, H-51, H-56, H-57, H-66, H-67) were generous gifts among 26 different microsomes prepared from human livers stored in the human liver bank of SRI International (Menlo Park, CA). Microsomal preparations of recombinant human CYP enzymes expressed by the human B lymphoblastoid cell line, AHH-1 (recombinant microsomes) were a gift from Gentest Corp. (Woburn, MA). The level of CYP2C9 in each microsome was assayed by immuno blotting as described previously (Imaoka et al., 1996).

Metabolic Assay of Tolbutamide Hydroxylation by Human Liver Microsomes or Recombinant CYP2C9. Tolbutamide was incubated with reaction mixture (1 ml) consisting of 0.2 mg of human liver microsomal protein or 3.5 pmol of recombinant CYP2C9 and NADPH-generating system (1 mM NADP, 10 mM glucose 6-phosphate, 0.1 U/ml glucose 6-phosphate dehydrogenase, and 5 mM MgCl2) in 100 mM potassium phosphate buffer (pH 7.4). Reactions were initiated by adding 100 µM of the NADPH-generating system (preincubated for 5 min). After incubation at 37°C in a shaking water bath for 75 min (human liver microsomes) or 60 min (recombinant CYP2C9), incubations were terminated by adding 100 µl of 2 M hydrochloric acid. After terminating the reaction, 2 ml of diethyl ether and 100 µl of chloropropamide solution (internal standard; 6 pmol/ml) were added. Incubation mixtures were centrifuged for 10 min (3000 rpm). The supernatant fraction from each incubation was transferred to a 15-ml test tube and evaporated to dryness under a mild stream of N2 gas. Residues were resuspended in 100 µl of the HPLC mobile phase, and 50 µl was used for analysis. The chromatograph was fitted with a TOSOH ODS-80TM column, which was eluted with 0.05% phosphoric acid, 75% MeOH, and 250, 500, 750, and 1000 µM. The initial formation rate of hydroxytolbutamide was plotted against concentration of substrate. Km and maximum metabolic rate (Vmax) values were estimated by the nonlinear least-squares regression method “MULTI” (Yamaoka et al., 1981).

Correlation between Tolbutamide Hydroxylation Activity and 2C9 Content of Human Liver Microsomes. Metabolic assays for tolbutamide hydroxylation were carried out using human liver microsomes (H-19, H-35, H-36, H-38, H-50, H-51, H-56, H-57, H-66, H-67) containing various amounts of CYP2C9. The initial concentration of tolbutamide was set at 100 µM and 1 mM. We investigated the correlation between CYP2C9 content of each microsome and the metabolic activity.

Identification of the CYP Enzyme Involved in Tolbutamide Hydroxylation. Metabolic assays for tolbutamide hydroxylation were carried out using nine enzymes of P450 (1A1, 1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 2E1, and 3A4). The initial concentration of tolbutamide was set at 300 µM. Reaction mixtures were incubated for 60 min. To estimate the contribution of each enzyme in vivo tolbutamide metabolism, we used average amounts of enzyme contained in 0.2 mg of human liver microsomal protein (Shimada et al., 1994). The amounts of each enzyme used were 8.4 pmol of CYP1A1 or 1A2, 0.2 pmol of 2B6, 3.5 pmol of 2C8, 2C9, or 2C19, 1.0 pmol of 2D6, 4.4 pmol of 2E1, or 19.2 pmol of 3A4.

Inhibition Study. Inhibitory effects of eight sulfonamides on tolbutamide metabolism were investigated using H-67 microsome. The concentration of tolbutamide was set at 100 µM. The inhibition constant (Ki) was obtained by fitting the inhibition curve to the following equation using the nonlinear least-squares regression method “MULTI”:

\[
\frac{V}{V_{\text{max}}} = \frac{V_{\text{max}} \cdot C}{K_i (1 + I_u/C) + C}
\]

(1)

The inhibitory effects of three sulfonamides (sulfaphenazole, sulfadiazine, and sulfamethizole) on the tolbutamide metabolism were also studied using recombinant CYP2C9 (Lot 27). To determine the inhibition type, tolbutamide (75–1000 µM) was incubated with H-67 microsome in the presence of sulfaphenazole (0.35 µM) or sulfamethizole (50 µM).

Prediction of Increase in AUC of Tolbutamide from In Vitro Metabolic Data. In the case of competitive or noncompetitive inhibition, the ratio of intrinsic metabolic clearance (CLint) in the presence and absence of the inhibitor can be described as follows when the substrate concentration is much lower than Km:

\[
\frac{\text{CL}_{\text{int}}(\text{Inhibitor})}{\text{CL}_{\text{int}}(\text{control})} = 1/(1 + I_u/K_i)
\]

(2)

where Km is the unbound concentration of the inhibitor and Ki is the inhibition constant of the inhibitor determined from in vitro inhibition studies. To avoid false negative predictions, Iu was calculated as the sum of the maximum unbound plasma concentration in circulating blood (Iu,max × fp) and that coming from gastrointestinal absorption after oral administration (eq. 3) assuming that the unbound liver concentration equals that in plasma (Ito et al., 1998a,b). Iu = (Iu,max + k4 × Fa × Dose/Qh) × fp

(3)

where k4 is the absorption rate constant, Fa is the fraction absorbed, Dose is the amount of inhibitor administered, Qh is the hepatic blood flow rate (1610 ml/min; Bischoff et al., 1971; Dedrick, 1973; Montandon et al., 1975), and fp is the unbound fraction in plasma. Values of Iu,max after single oral administration of a typical therapeutic dose of sulfonamides were calculated from the reported concentration profiles assuming a linear pharmacokinetic. The Iu,max values for the sulfonamides were assumed to be 0.1 min-1 except for sulfaphenazole (k4 = 0.085 min-1; Vree et al., 1990a). Fa was assumed to be 1.0 except for sulfaphenazole (Fa = 0.85; Ueda et al., 1972). The AUC ratio in the presence and absence of inhibitor can be calculated from the following equation in the case of a low clearance drug such as tolbutamide, assuming that the protein binding is not altered by the inhibitor (Ito et al., 1998a,b):

\[
\text{AUC ratio} = 1/(0.8(1 + I_u/K_i) + 1 - 0.8)
\]

(4)

where 0.8 is the fraction of tolbutamide eliminated by CYP2C9 metabolism (Miller et al., 1990).
Results

Enzyme Kinetics of Tolbutamide Hydroxylation by Human Liver Microsomes. Tolbutamide metabolism to hydroxytolbutamide was studied using four human liver microsomes that contained different amounts of CYP2C9. The hydroxylation by all microsomes followed Michaelis-Menten kinetics. Figure 1 shows typical Eadie-Hofstee plots for two microsomal samples. $K_m$ and $V_{max}$ values obtained by nonlinear least-squares regression method are summarized in Table 1.

Interindividual Difference in Tolbutamide Hydroxylation by Human Liver Microsomes. Good correlations were observed between CYP2C9 content and tolbutamide hydroxylating activity of ten human liver microsomes (Fig. 2). The correlation coefficient was $r = 0.955$ and $r = 0.904$ when tolbutamide concentration was set at 1 mM and 100 μM, respectively.

Contribution of Each CYP Enzyme to Tolbutamide Hydroxylation. The CYP enzymes expressed in human B lymphoblastoid cells did not metabolize tolbutamide except for CYP2C8, 2C9, and 2C19 (Fig. 3). The tolbutamide hydroxylation activity of CYP2C8 was less than one-third that of CYP2C9. When the same amounts of CYP2C enzymes were used for the assay, CYP2C19 had an even weaker activity, only about one-fifth that of CYP2C9. We also used the physiological amount of CYP2C19 (0.27 pmol; average amount of ten human liver microsomes used in our assay) for this assay, but 0.27 pmol of CYP2C19 did not produce hydroxytolbutamide more than the detection limit. From these data, it was shown that the major enzyme for tolbutamide hydroxylation is CYP2C9.

Enzyme Kinetics of Tolbutamide Hydroxylation by Recombinant CYP2C9. Tolbutamide hydroxylation by two lots of recombinant CYP2C9 that was expressed by human B lymphoblastoid cells followed Michaelis-Menten kinetics (Fig. 4). $K_m$ and $V_{max}$ values obtained by the nonlinear least-squares regression method were approximately 130 to 190 μM and 8.3 to 9.1 pmol/min/pmol enzyme, respectively, which were comparable with those obtained by human liver microsomes (Table 1).

Inhibitory Effects of Sulfonamides. All of the eight sulfonamides showed inhibitory effects on tolbutamide hydroxylation in both human liver microsomes (Fig. 5) and recombinant CYP2C9 (Fig. 6). The extent of inhibition, however, differed among the sulfonamides. The most potent inhibitor was sulfaphenazole whose $K_i$ value in human liver microsomes was about 0.3 μM, which was much smaller than that of other sulfonamides (Table 2). Among the drugs investigated, sulfadiazine and sulfamethizole were relatively potent inhibitors with $K_i$ values of about 50 μM. The $K_i$ values of sulfaphenazole and these two drugs in the recombinant CYP2C9 were comparable with those in human liver microsomes (Table 3).

Inhibition Type of Sulfonamides. The slope of the Eadie-Hofstee plot for tolbutamide hydroxylation by human liver microsomes was decreased by both sulfaphenazole and sulfamethizole with no substantial change in $x$-intercept (Fig. 7). This showed that inhibition type of sulfonamides was competitive. Furthermore, tolbutamide hydroxylation activities of human liver microsome (H-67) after a 10- and 40-min preincubation with 3 μM sulfaphenazole were equal to the control value without preincubation.

Prediction of AUC Increase. $I_u$ (maximum unbound concentration of inhibitor in the portal vein) was calculated from literature data on an average dose in clinical practice, plasma unbound fraction, maximum systemic concentration, absorption rate constant, and fraction absorbed of each sulfonamide (Table 4). The calculated degrees of increase in tolbutamide AUC caused by coadministration of sulfonamides are summarized in Table 5. The most potent inhibitor was sulfaphenazole, which was estimated to increase tolbutamide AUC about 5-fold. Tolbutamide AUC was estimated to be increased around or more than 2-fold by three sulfonamides that have relatively small $K_i$ values, i.e., sulfamethizole, sulfadiazine, and sulfisoxazole. AUC increase by other sulfonamides was predicted to be less than 1.5-fold.
cause tolbutamide is metabolized by a single pathway (Thomas and Ikeda, 1966), its inhibition will have serious effects.

Tolbutamide hydroxylation in human liver microsomes followed Michaelis-Menten kinetics (Fig. 1). $K_m$ and $V_{max}$ values obtained in the present study (Table 1) were consistent with those reported by Doecke et al. (1991; $K_m = 85.6 \mu M$) and by Miners et al. (1988; $K_m = 120 \mu M$, $V_{max} = 0.273 \text{ nmol/min/mg}$).

In metabolic studies using ten human liver microsomes containing various amounts of CYP2C9, significant correlation was observed between CYP2C9 content and initial velocity of tolbutamide hydroxylation (Fig. 2). This suggests that hydroxylation of tolbutamide is mainly mediated by CYP2C9, which supports the previous reports (Knodell et al., 1987; Miners et al., 1988; Back and Orme, 1989; Brian et al., 1989; Veronese et al., 1990a, 1993). Furthermore, estimation of the contribution of each enzyme using recombinant CYP enzymes also showed that CYP2C9 is a major enzyme in tolbutamide hydroxylation (Fig. 3). Figure 3 shows that tolbutamide is slightly metabolized also by the other CYP2C enzymes. Some groups have reported that CYP2C8 is also involved in hydroxylation of tolbutamide (Relling et al., 1990; Srivastava et al., 1991; Veronese et al., 1993). However, the average level of CYP2C8 in human liver microsomes is reported to be less than one-third that of CYP2C9 (Imaoka et al., 1996), suggesting little contribution of CYP2C8 in tolbutamide hydroxylation. Nevertheless, the correlation between tolbutamide hydroxylation activity and CYP2C9 content does not intercept at the origin (Fig. 2), suggesting that part of tolbutamide hydroxylation (e.g., about 25% of total hydroxylation of 100 $\mu M$ tolbutamide by microsomes with average content of CYP2C9, i.e., about 15 pmol/mg protein) may be mediated by some other enzymes. It cannot be ruled out that such enzymes as CYP2A6, CYP3A5, or CYP4A9/11 etc., the recombinant systems of which have not been investigated in Fig. 3, may be partly involved in tolbutamide hydroxylation.

The $K_m$ and $V_{max}$ values (per picomole of CYP2C9) for tolbutamide hydroxylation by recombinant CYP2C9 were comparable with those obtained using human liver microsomes (Table 1). This finding indicates that recombinant CYP2C9 can be used as an alternative to human liver microsomes in prediction of in vivo drug interactions of tolbutamide.

The inhibition study using human liver microsomes showed that sulfonamides had various $K_i$ values for tolbutamide hydroxylation (Table 2). Sulfaphenazole was the most potent inhibitor among all the sulfonamides examined. From the Dixon plot analysis using human liver microsomes, the $K_i$ values of sulfaphenazole, sulfamethizole, and sulfamethoxazole for tolbutamide hydroxylation are reported to be 0.3, 3.5, and 254 $\mu M$, respectively (Back et al., 1988), which are consistent with our results. The $K_i$ values of sulfonamides, except for sulfaphenazole, showed that they are less potent inhibitors. This experimental result could explain why there are fewer reports of elevation of tolbutamide concentration or hypoglycemia due to these other sulfonamides. In addition, tolbutamide hydroxylation by recombinant CYP2C9 was also inhibited by sulfonamides (Table 3). $K_i$ values obtained in this study were comparable with those obtained in the liver microsomal study.

The reason why only sulfaphenazole has such a potent inhibitory effect is unknown. One hypothesis is that sulfaphenazole could inhibit tolbutamide hydroxylation by a mechanism-based inhibition (Ito et al., 1998b). However, metabolic assay was performed after a 10- and 40-min preincubation of human liver microsomes with sulfaphenazole and the inhibitory effect exhibited no significant differences, suggesting that the mechanism-based inhibition is not involved in the inhibition of CYP2C9 by sulfaphenazole. Also, an Eadie-Hofstee plot analysis using a fixed concentration of inhibitor showed changes in

---

**Fig. 3.** The contribution of each P450 enzyme in tolbutamide hydroxylation.

Reaction mixture contained 8.4 pmol of recombinant CYP1A1 or 1A2; 0.2 pmol of 2B6; 3.5 pmol of 2C8, 2C9, or 2C19; 1.0 pmol of 2D6; 4.4 pmol of 2E1; or 19.2 pmol of 3A4. Each recombinant enzyme was incubated with tolbutamide (300 $\mu M$) for 60 min. Each point and vertical bar represents the mean ± S.E.

**Fig. 4.** Tolbutamide hydroxylation by recombinant CYP2C9.

Reaction mixture contained 3.5 pmol of recombinant CYP2C9, and was incubated for 60 min. Each point and vertical bar represents the mean ± S.E. (a) and (b), Lot 2; (c) and (d), Lot 27.
Kₘ values with minor changes in Vₘₐₓ values (Fig. 7), indicating a competitive inhibition at least at this concentration of inhibitor. The Kᵢ values for sulfaphenazole and sulfamethizole estimated from Fig. 7 (0.22 and 39 μM, respectively) were not so different from those obtained from Fig. 5, assuming a competitive inhibition (0.31 and 53 μM, respectively; Table 5), suggesting that a competitive inhibition also takes place at other concentrations of inhibitor.

The AUC of tolbutamide (500 mg p.o.) was predicted to increase about 5-fold by coadministration of sulfaphenazole 500 mg, which is equal to the reported increase (Veronese et al., 1990b) (Table 5). Furthermore, the AUC of tolbutamide (750 mg p.o.) is reported to increase by 1.6 times when sulfamethizole (1 g) is coadministered (Lumholtz et al., 1975), which was also predicted well by our method. Our prediction is based on avoiding false negatives, so that the portal

<table>
<thead>
<tr>
<th>TABLE 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>The Kᵢ values for the inhibition of tolbutamide metabolism by various sulfonamides in human liver microsomes</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>Sulfaphenazole</td>
</tr>
<tr>
<td>Sulfadiazine</td>
</tr>
<tr>
<td>Sulfamethizole</td>
</tr>
<tr>
<td>Sulfisoxazole</td>
</tr>
<tr>
<td>Sulfamethoxazole</td>
</tr>
<tr>
<td>Sulfapyridine</td>
</tr>
<tr>
<td>Sulfadimethoxine</td>
</tr>
<tr>
<td>Sulfamonomethoxine</td>
</tr>
</tbody>
</table>

Numbers in parentheses represent the S.E.s of the mean values.

FIG. 5. Inhibitory effects of sulfonamides on tolbutamide metabolism by human liver microsome (H-67).

Reaction mixture contained 100 μM tolbutamide. Kᵢ values for the sulfonamides are summarized in Table 2. Each point and vertical bar represents the mean ± S.E. Curves represent the fitting lines (eq. 1). a, sulfaphenazole (n = 6); b, sulfadiazine (n = 6); c, sulfamethizole (n = 4); d, sulfisoxazole (n = 3); e, sulfamethoxazole (n = 4); f, sulfapyridine (n = 6); g, sulfadimethoxine (n = 6); h, sulfamonomethoxine (n = 3).

FIG. 6. Inhibitory effects of sulfonamides on tolbutamide metabolism by recombinant CYP2C9 (Lot 27).

Reaction mixture contained 100 μM tolbutamide. Kᵢ values for the sulfonamides are summarized in Table 3. Each point and vertical bar represents the mean ± S.E. Curves represent the fitting lines (eq. 1). a, sulfaphenazole; b, sulfadiazine; c, sulfamethizole.
Concentrations in the clinical situation, especially in the case of repeated administration of higher doses of the sulfonamides, may be higher than estimated in this study, which may result in greater degrees of in vivo interactions.

In this study, we systematically evaluated the inhibitory effects of sulfonamides on tolbutamide metabolism mediated by CYP2C9. This evaluation may also be applied to other drugs that are metabolized by CYP2C9. Therefore, attention should be paid in coadministration of sulfonamides with relatively small $K_i$ values (sulfaphenazole, sulfadiazine, sulfamethizole, and sulfisoxazole) and CYP2C9 substrates with narrow therapeutic ranges such as phenytoin (an antiepileptic) and warfarin (an anticoagulant).

### References


