IN VITRO CHARACTERIZATION OF CYTOCHROME P450 2D6 INHIBITION BY CLASSIC HISTAMINE H1 RECEPTOR ANTAGONISTS

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ABSTRACT:

Classic antihistamines, namely diphenhydramine, chlorpheniramine, clemastine, perphenazine, hydroxyzine, and tripelennamine, share structural features with substrates and inhibitors of the polymorphic cytochrome P450 (CYP) isozyme CYP2D6. Therefore, the current study was undertaken to characterize the inhibition of CYP2D6 by these commonly used, histamine H1 receptor antagonists. Microsomal incubations were performed using bufuralol as a specific CYP2D6 substrate and microsomes derived from human cells transfected with CYP2D6 cDNA. Reaction velocities were assessed in the absence and presence of antihistamines (20 μM) at 11 substrate concentrations (1, 2.5, 5, 7.5, 10, 15, 20, 25, 50, 75, and 100 μM), as well as at three nonsaturating substrate concentrations (2.5, 5, and 20 μM) and three inhibitor concentrations (5, 20, and 50 μM). In the presence of all antihistamines, the Vmax and Km of bufuralol 1'-hydroxylation were significantly altered, compared with the uninhibited reaction (p < 0.05). Lineweaver-Burke plots suggested competitive inhibition of the reaction by diphenhydramine and mixed inhibition by all other antihistamines tested. Diphenhydramine and chlorpheniramine, with estimated Ki values of ~11 μM, were the weakest inhibitors of CYP2D6 in vitro. Whereas tripelennamine, promethazine, and hydroxyzine were similar in their inhibitory capacities (Ki ~4–6 μM), clemastine appeared to be significantly more potent, with a Ki of ~2 μM. These data demonstrate that classic histamine H1 receptor antagonists, available in over-the-counter preparations, inhibit CYP2D6 in vitro. Furthermore, the CYP2D6-inhibitory concentrations of these antihistamines are in the range of their expected hepatic blood concentrations, suggesting that, under specific circumstances, clinically relevant interactions between classic antihistamines and CYP2D6 substrates might occur.

The classic histamine H1 receptor antagonists clemastine, diphenhydramine, chlorpheniramine, tripelennamine, promethazine, and hydroxyzine were introduced into clinical practice ~50 years ago and are today among the most commonly used drugs in the world (Simons and Simons, 1994). Because these compounds have excellent overall safety records, they are found in a wide variety of over-the-counter cold and allergy treatments, as well as in sleeping aids, and they are often used in combination with other drugs (Simons and Simons, 1994). Despite their widespread use over an extended period, little is known about their pharmacokinetics, particularly their interactions with specific P4501 isozymes.

Several observations argue for an important role of the polymorphic P450 isoyme CYP2D6 in the metabolism and the interaction profiles of classic antihistamines. First, the structural criteria elaborated for the optimal binding of diphenhydramine and its analogues to P450 (Rekkka et al., 1989) are very similar to the structural characteristics of many known CYP2D6 substrates and inhibitors (de Groot et al., 1997). Second, a recent study demonstrated that, in vitro, the residual activity of bufuralol 1-hydroxylation was lowest in the presence of clemastine (5% of control activity at a concentration of 100 μM clemastine) and highest in the presence of diphenhydramine (40% of control activity at a concentration of 100 μM diphenhydramine) (Nakamura et al., 1996). The same investigators demonstrated that promethazine and chlorpheniramine inhibited CYP2D6 activity in vitro, although the type of inhibition was not determined (Nakamura et al., 1996). Third, and consistent with these in vitro data and structural considerations, a case study reported a 2-fold increase in the half-life of diphenhydramine in an elderly woman with impaired metabolism of the CYP2D6 substrate imipramine (Glassman et al., 1985). Lastly, the pharmacokinetics of classic antihistamines are characterized by intersubject variability (Chiu et al., 1979; Huang et al., 1982) similar to that described for CYP2D6 substrates; therefore, this variability might be the result of a genetically determined lack of metabolic capacity (Alvan, 1991).

The isoyme CYP2D6 has received considerable attention because of the presence of a genetic polymorphism that divides the population into individuals with high enzyme activity (extensive metabolizers) and individuals (5–10% of the population) with low enzyme activity (poor metabolizers) (Mahgoub et al., 1977). Poor metabolizers are predisposed to the accumulation of CYP2D6 substrates and drug-induced adverse effects (Lennard, 1993). Similarly, when extensive metabolizers are treated simultaneously with a substrate and a potent

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1 Abbreviation used is: P450 or CYP, cytochrome P450.

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inhibitor of CYP2D6, substrate accumulation occurs (Brosen et al., 1987).

The objectives of the present study were to substantiate and extend previous work by determining the type of inhibition of in vitro CYP2D6 activity produced by the classic histamine H1 receptor antagonists diphenhydramine, chlorpheniramine, promethazine, and clemastine. Furthermore, we intended to determine the type and extent of CYP2D6 inhibition produced by two additional, structurally related, classic antihistamines, namely triprolamine and hydroxyzine.

Materials and Methods

Chemicals. Clemastine fumarate, chlorpheniramine maleate, promethazine hydrochloride, triprolamine hydrochloride, diphenhydramine hydrochloride, hydroxyzine dihydrochloride, NADP, glucose-6-phosphate, and glucose-6-phosphate dehydrogenase were obtained from Sigma Chemical Co. (St. Louis, MO). Magnesium chloride was purchased from Anachemia Science (Ville St-Pierre, Montreal, Quebec, Canada), and (-)-bufuralol hydrochloride and 1’-hydroxybufuralol were purchased from Gentest Corp. (Woburn, MA). Other chemicals were obtained from the usual commercial sources and were of analytical grade.

Recombinant Human CYP2D6 Isozymes. Human microsomes expressing CYP2D6 protein was purchased from Gentest. This protein was derived from a human AHH-1-TK+/− cell line transfected with cDNA encoding human CYP2D6-Val374.

Microsomal Incubations. Incubations (final volume, 250 µl) contained substrate (0–350 µM bufuralol in 100 mM potassium phosphate buffer, pH 7.4), 10 µl of microsomes (0.4 mg/ml), 100 µM of a NADP Regenerating System (7.75 mM of NADP, 7.75 mM of glucose-6-phosphate, and 36 units of glucose-6-phosphate dehydrogenase in 3.0 ml of 100 mM potassium phosphate buffer, pH 7.4), and potassium phosphate buffer (100 mM, pH 7.4). Incubation mixtures containing microsomes, substrate, and buffer were preincubated at 37°C for 15 min before the addition of the NADP Regenerating System. As a result, the incubation was incubated at 37°C for 30 min in 37°C in a Dubnoff incubator (Precision Scientific, Chicago, IL), and the reaction was stopped by the addition of 25 µl of perchloric acid (69–72%, by volume). Proteins were sedimented by centrifugation. All experiments were performed in duplicate.

Clemastine fumarate, chlorpheniramine maleate, promethazine hydrochloride, triprolamine hydrochloride, diphenhydramine hydrochloride, and hydroxyzine dihydrochloride were dissolved in potassium phosphate buffer (100 mM, pH 7.4). To characterize the extent and type of inhibition of CYP2D6 by the antihistamines, incubation mixtures contained microsomes, buffer, cofactor, 11 concentrations of bufuralol (1–100 µM), and a fixed concentration (20 µM) of each antihistamine. In addition, for graphical determination of Ki values, incubations were performed with three nonsaturating bufuralol concentrations (2.5, 5, and 20 µM) and three inhibitor concentrations (5, 20, and 50 µM). Bufuralol and its hydroxylated metabolite were analyzed by HPLC as described previously (Kronbach et al., 1987).

Data Analysis. Reaction velocities were expressed in units of picomoles of 1’-hydroxybufuralol formed per picomole of CYP2D6 per minute. Velocity data for bufuralol 1’-hydroxylation in the absence and presence of antihistamines were estimated by derivative-free, iterative, nonlinear, least-squares regression (Fig60; Bioskop, Ferguson, MO). Comparisons of the rates of formation of 1’-hydroxybufuralol at various inhibitor concentrations were accomplished graphically by using the Dixon method. In addition, Ki values were estimated by nonlinear, least-squares regression, using equations for competitive [V = Vmax × S/(Km + [I]Ki)], noncompetitive [V = Vmax × S/(1 + [I]Ki)] + Km, or mixed [V = Vmax × S/(1 + [I]Ki) + Km] (1 + [I]Ki) inhibition. Data are expressed as means and 95% confidence intervals.

Results and Discussion

Estimated Vmax, Km, and Ki values for bufuralol 1’-hydroxylation in vitro are summarized in table 1. Compared with the uninhibited reaction (control), addition of all antihistamines resulted in significant changes of Vmax and Km, with no overlap of the 95% confidence intervals (all p < 0.05, compared with control). Lineweaver-Burke plots (not shown) suggested that diphenhydramine is a competitive inhibitor and that chlorpheniramine, clemastine, triprolamine, promethazine, and hydroxyzine are mixed inhibitors of CYP2D6 in vitro (table 1). Fitting of velocity data by derivative-free, iterative, nonlinear, least-squares regression, assuming competitive or mixed inhibition, revealed that diphenhydramine and chlorpheniramine, with estimated Ki values of ~11 µM, were significantly weaker inhibitors of CYP2D6 in vitro than were the other antihistamines tested (both p < 0.05). Triprolamine, promethazine, and hydroxyzine were similar in their inhibitory capacities (Ki = 4–6 µM) and were significantly more potent inhibitors than diphenhydramine and chlorpheniramine (p < 0.05). Of all antihistamines tested, clemastine appeared to be the most potent in vivo inhibitor, with a Ki of ~2 µM (p < 0.05, compared with all other antihistamines). Ki values obtained by regression analysis corresponded to the respective Ki values obtained from Dixon plots. These results appear to be clinically relevant, because all antihistamine drugs tested undergo extensive first-pass metabolism, resulting in hepatic blood concentrations that are expected to lie in the range of the determined Ki values (table 2).

Our results are in agreement with previous reports of in vitro CYP2D6 inhibition by various antihistamines. Ki values reported for diphenhydramine ranged from 0.124 to 2.5 µM, depending on the type of in vitro system and substrate used (Fonne-Pfister and Meyer, 1988; Hiroi et al., 1995). Furthermore, Ki values for promethazine and chlorpheniramine were graphically estimated to be 13 and 20 µM, respectively, and clemastine as well as diphenhydramine inhibited CYP2D6-mediated bufuralol 1’-hydroxylation in vitro (Nakamura et al., 1996). The data presented by us extend previous work by including a whole series of the most commonly used classic antihistamines, by assessing not only the extent but also the type of inhibition, and by using a more objective approach of nonlinear regression analysis.

These in vitro data, suggesting that classic antihistamines may indeed be rather potent inhibitors of CYP2D6, are somewhat incon-
and, despite an estimated accumulation factor of 4–9 for chlorpheniramine (Yasuda et al., 1995), could occur. However, multiple-dose data are limited to a few subjects in which the enzyme, one might speculate that autoinhibition and indirectly for chlorpheniramine (Yasuda et al., 1995), and they inhibit the same enzyme, one might speculate that autoinhibition could occur. However, multiple-dose data are limited to a few subjects and, despite an estimated accumulation factor of 4–9 for chlorpheniramine, its elimination half-life was similar after single (two subjects) and multiple (two different subjects) doses (Huang et al., 1982).

The presence of aromatic rings and alkyl substituents renders classic antihistamines very lipophilic and allows these molecules to readily traverse membranes. Thus, after oral administration, antihistamine drugs undergo extensive first-pass elimination, resulting in relatively low plasma concentrations (table 2) (Tonn et al., 1996). However, hepatic blood concentrations of classic antihistamines are 7–42-fold higher than the respective plasma concentrations, based on data on radio labelled drug distribution in animals or findings from autopsy specimens (table 2). Thus, expected human hepatic blood concentrations in the range of the determined CYP2D6-inhibitory concentrations for all classic antihistamine drugs tested. In support of this hypothesis, we recently reported that chronic administration of diphenhydramine significantly prolonged the hemodynamic effects of metoprolol in healthy volunteers with high CYP2D6 activity, compared with those with low CYP2D6 activity (Bouayad et al., 1997).

In summary, six classic antihistamines, namely diphenhydramine, chlorpheniramine, clemastine, promethazine, trimepzenamine, and hydroxyzine, inhibit CYP2D6-mediated bufuralol 1'-hydroxylation in vitro. Because these compounds accumulate in the liver and K, values are in the range of expected antihistamine concentrations in hepatic blood, classic H1 receptor antagonists may cause clinically relevant drug-drug interactions with cardiovascular, antidepressant, and antipsychotic CYP2D6 substrates. Thus, well-controlled studies of the interactions of classic antihistamines with such substrates must be performed with individuals with high and low CYP2D6 activities.

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References


