

Correlation of Cytochrome P450 (CYP) 1A2 Activity Using Caffeine Phenotyping and Olanzapine Disposition in Healthy Volunteers

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Olanzapine has previously been shown to have predominant metabolism by cytochrome (CYP) P450 1A2. Caffeine has been shown to provide an accurate phenotypic probe for measuring CYP1A2 activity. The purpose of this study is to determine if a significant correlation exists between olanzapine disposition and caffeine metabolic ratios. Subjects were phenotyped for CYP1A2 activity with caffeine probe methodology. After 200-mg caffeine administration, blood (4 h), saliva (6 and 10 h), and urine (8 h total) were collected for high-performance liquid chromatography (HPLC) analysis of caffeine and its metabolites. CYP1A2 activity was measured as plasma (PMR_{4h}), saliva (SMR_{6h} and SMR_{10h}), and three urinary metabolic (UMR_{18h}, UMR_{28h}, and UMR_{38h}) ratios. Each of the 14 healthy nonsmokers (13 male) received a single 10 mg olanzapine dose after which blood was collected for HPLC determination of olanzapine concentrations at predose and at 0.5, 1, 1.5, 2, 2.5, 3, 4, 5, 6, 8, 10, 12, 24, 48, 72, 96, and 120 h postdose. Olanzapine pharmacokinetic parameters in this study were similar to those previously published. All caffeine metabolic ratios (PMR_{4h}, SMR_{6h}, SMR_{10h}, UMR_{18h}, and UMR_{28h}) significantly correlated with each other ($p < 0.001$) except for UMR_{38h}, which did not correlate. A significant correlation ($p < 0.05$) was also found between olanzapine clearance and PMR_{4h} ($r = 0.701$), SMR_{6h} ($r = 0.644$), SMR_{10h} ($r = 0.701$), UMR_{18h} ($r = 0.745$), and UMR_{28h} ($r = 0.710$). A negative correlation was observed between olanzapine clearance and UMR_{38h} ($r = -0.029$, $p = \text{NS}$). A significant correlation was found between olanzapine clearance and various caffeine metabolic ratios. Interpatient variability in CYP1A2 activity may explain the wide interpatient variability in olanzapine disposition. Compounds that modulate CYP1A2 activity may be expected to alter olanzapine pharmacokinetics accordingly.

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INTRODUCTION

Caffeine (1,3,7-trimethylxanthine [137X]) is one of the most commonly ingested compounds throughout the world. The metabolic pathways of caffeine are complex but involve the formation of three principal metabolites: theobromine (3,7-dimethyl xanthine [37X]), theophylline (1,3-dimethylxanthine [13X]), and paraxanthine (1,7-dimethylxanthine [17X]). Each of these metabolites are primarily formed by cytochrome (CYP) P450 1A2 (Carrillo *et al*, 2000). Conversion to the 17X metabolite comprises about 80% of the metabolic pathway for caffeine. The metabolic pathways to

the 37X and 13X metabolites are also influenced by CYP2E1 (Sinues *et al*, 1999). Caffeine has become popular as a metabolic probe for CYP1A2 activity in humans (Carrillo *et al*, 2000); it is one of the major P450 cytochromes in the liver and accounts for 15% of the total P450 content.

In addition to caffeine, clinically important psychotropic medications such as olanzapine, clozapine, haloperidol, thioridazine, imipramine, clomipramine, fluvoxamine, and tacrine are completely or partially metabolized by this enzyme (Carrillo *et al*, 2000; Callaghan *et al*, 1999; Bertilsson *et al*, 1994; Spigset *et al*, 1999a; Otani and Aoshima, 2000). Thus, interindividual variability in CYP1A2 activity may alter the pharmacokinetics of these medications and influence response to therapy, including toxicities such as tardive dyskinesia with older (typical) antipsychotics (Carrillo *et al*, 2000; Kelly *et al*, 1999; Basile *et al*, 2000; Ou-Yang *et al*, 2000).

Olanzapine is an atypical antipsychotic agent used in the treatment of schizophrenia, bipolar disorders, and other psychotic conditions. Olanzapine is metabolized to 10- and 4'-N-glucuronide, 4'-N-desmethylolanzapine via CYP1A2, and olanzapine N-oxide by flavin monooxygenase

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(Callaghan *et al*, 1999). Conversion to the 2-hydroxymethylolanzapine by CYP2D6 is a minor pathway for olanzapine (Ring *et al*, 1995). Clozapine, like olanzapine, is metabolized by a variety of CYP isozymes including CYP1A2, CYP3A4, CYP2D6 and the flavin monooxygenase (Jann *et al*, 1993; Otani and Aoshima, 2000). Clozapine disposition was found to covary with CYP1A2 activity in 14 healthy volunteers, where the reciprocal of clozapine's area under the plasma concentration-time curve (AUC) significantly correlated with urinary caffeine *N*-3 demethylation index or its conversion to 17X metabolite ($r = 0.84$, $p < 0.0024$). Clozapine clearance was also found to correlate significantly with the other caffeine 37X and 13X indexes ($r = 0.89$, $p < 0.0013$; and $r = 0.85$, $p < 0.0023$) (Bertilsson *et al*, 1994). Olanzapine disposition was evaluated in seventeen healthy volunteers where concentrations of caffeine and 17X were measured in saliva 10 h postcaffeine dosing (Hagg *et al*, 2001). A significant correlation between caffeine activity (caffeine/17X ratio) and olanzapine clearance was not found ($r = 0.19$, $p = 0.56$). However, each of these studies used only one biological matrix to assess caffeine metabolism (such as only urine or saliva concentrations).

The purpose of this study was to examine the relation between olanzapine disposition and CYP1A2 activity as determined by caffeine phenotyping with plasma, saliva and urine concentrations of caffeine, and its 17X and 13X metabolites.

METHODS

Subjects

The Mercer University Institutional Review Board approved this study and each participant provided written informed consent. In all, 15 healthy nonsmokers (14 male, one female) participated in this study. Subjects were included in our investigation if they met the following inclusion criteria: medically healthy volunteers, men and women over 18 years of age, women of childbearing potential on medically acceptable birth control excluding oral contraceptives, hormonal implants, or depo-injections. Subjects were also excluded if they met any of the following exclusion criteria: presence of any chronic medical condition, pregnancy and/or breastfeeding, history of alcohol, tobacco or drug abuse, use of any prescription or over-the counter medications or herbal remedies metabolized by the CYP P450 1A2 system, or an inability to comply with a diet free of caffeine, ethanol, and grapefruit juice and products 48 h before the overnight stay prior to the study day. The health status of our subjects was assessed by a thorough health and medication history, physical examination, and a complete metabolic and hematologic blood profile, prior to study protocol initiation.

Study Design

Subjects were asked to comply with a diet free of caffeine, ethanol, and grapefruit juice/products for 48 h, and come to the Center for Clinical Research (CCR) for an overnight stay of approximately 12 h prior to the initiation of the study day. The investigator confirmed compliance with the protocol-specified diet via detailed subject interviews. After eating a standardized breakfast, subjects received 200 mg of

caffeine (No-Doz[®]). Previously standardized procedures were used for obtaining plasma; saliva and urine samples were established by previous investigators (Carrillo *et al*, 2000; Spigset, 1999b). Each subject had two 10 ml blood samples drawn; one sample was obtained prior to caffeine administration and the second sample 4 h postcaffeine dosing. All samples were collected in EDTA vacutainer tubes and centrifuged at 3200 g for 10 min after which plasma was harvested and stored at -80°C until it was analyzed for caffeine and paraxanthine.

Immediately after caffeine administration, subjects began to collect their urine and continued to do so over the next 8 h. All urine samples were measured individually for total volume and then aliquoted into smaller samples, which were subsequently acidified to a pH of 4.0 with 0.1 M hydrochloric acid before storage at -80°C (preservation of caffeine metabolites). At 6 and 10 h after caffeine administration subjects were asked to chew on a piece of paraffin wax film to stimulate saliva production and a 2 ml sample of saliva was collected and stored at -80°C until the time of analysis.

Each of the 14 subjects received a single 10 mg olanzapine dose. Venous blood samples were collected for high-performance liquid chromatography (HPLC) determination of olanzapine concentrations at predose and at 0.5, 1, 1.5, 2, 2.5, 3, 4, 5, 6, 8, 10, 12, 24, 48, 72, 96, and 120 h postdose. All samples were stored at -80°C until assayed. One male subject withdrew from the study for personal reasons and did not participate in olanzapine dosing.

Analytic Methods

Olanzapine plasma concentrations were assayed by Bioanalytical Services (BAS; west Lafayette, IN) by a published HPLC method developed by the drug's manufacturer (Eli Lilly Pharmaceuticals, Indianapolis, IN) by HPLC (Catlow *et al*, 1995). Plasma, urine and salivary caffeine (137X) metabolites (17X, 13X, 5-acetylamin-6-amino-3-methyluracil (AAMU), 1-methylxanthine (1X), 1-methyl uric acid (1U), and 1,7 dimethyl uric acid (17U) were analyzed using validated HPLC methodology (El-Yazigi *et al*, 1999; Fuhr and Rost, 1994). CYP1A2 activity was measured as plasma (PMR), salivary (SMR), and urinary metabolite (UMR) to caffeine metabolic ratios were validated in previous phenotypic studies (Fuhr *et al*, 1996; Spigset *et al*, 1999; Carrillo *et al*, 2000).

Data and Statistical Analysis

Olanzapine pharmacokinetic parameters values for each subject were determined with the use of noncompartmental methods and are displayed in Table 1. Maximal plasma concentrations (C_{max}) and time to reach C_{max} (T_{max}) were determined by visual inspection of the concentration-time profiles. The elimination rate constant (λ_z) was estimated as the absolute value of the slope of a linear regression of natural logarithm of concentration vs time. Half-life was calculated as $\ln 2/\lambda_z$. The area under the plasma concentration-time curve ($\text{AUC}_{0-\infty}$) was determined by the linear trapezoidal rule with extrapolation to infinity ($\text{AUC}_{0-\infty}$) by dividing the last concentration measured by the elimination rate constant (λ_z). Oral clearance (CL/F) was determined by dividing the dose administered by $\text{AUC}_{0-\infty}$ and the

Table 1 Olanzapine Single-Dose Pharmacokinetics and Caffeine PMR_{4h} are Presented (N = 14)

Subject	AUC _{0→∞} (ng h/ml)	CL/F (l/h)	T _{1/2} (h)	C _{max} (ng/ml)	T _{max} (h)	PMR _{4h}
1	541	18.5	24	17	3	0.342
2	833	12	29	25	10	0.129
3	520	19.2	50	19	2	0.161
4	449	22.3	30	10	10	0.648
5	481	20.8	32	22	3	0.590
6	430	23.3	25	20	2	0.717
7	431	23.2	27	12	10	0.483
8	565	17.7	31	18	4	0.390
9	441	22.7	29	12	2	0.522
10	672	14.9	36	12	10	0.267
11	369	27.1	27	17	3	0.589
12	539	18.6	30	13	6	0.292
13	399	25.1	45	13	2	0.296
14	508	19.7	36	12	3	0.311
Mean	501	20	32	15	4	0.410
CI	443–582	18–23	28–36	13–19	3–7	0.312–0.508

AUC_{0→∞}, AUC from time zero to infinity; CL/F, oral clearance; T_{1/2}, half-life; C_{max}, maximum concentration; T_{max}, time to maximum concentration; V_d, apparent volume of distribution. Mean, geometric mean; CI, 95% confidence intervals.

apparent volume of distribution (V/F) was calculated as the ratio of CL/F to λ_z . The 95% confidence intervals (CIs) for the differences in means for the log-transformed pharmacokinetic parameter values were transformed back to the original scale to give 95% CIs for the geometric means.

CYP1A2 activity was determined as molar ratios (MRs) of caffeine metabolites in plasma (P) at 4 h (PMR_{4h}) calculated as 17X/137X, and saliva (S) at 6 and 10 h (SMR_{6h} and SMR_{10h}) calculated as 17X/137X. The urinary (U) MR at 8 h (UMR_{8h}) were determined in three different calculations with different metabolite concentrations: UMR_{18h} = 17X/137X; UMR_{28h} = 17X + 17U/137X; and UMR_{38h} = AAMU + 1X + 1U/17U (Fuhr *et al*, 1996; Spigset *et al*, 1999; Carrillo *et al*, 2000). Olanzapine clearance data and caffeine metabolic ratios were tested for normality with Shapiro–Wilk tests. Olanzapine clearance covariation with CYP1A2 activity PMR_{4h}, SMR_{6h}, SMR_{10h}, UMR_{18h}, UMR_{28h}, and UMR_{38h} was tested with Pearson's correlation coefficient. Statistical significance is defined as $p < 0.05$ and calculations were performed with Statistica 6.0[®].

RESULTS

The 14 subjects (13 men and one woman) were between 20 and 28 years old (mean, 25 ± 2.9 years). A wide intersubject variability was found with olanzapine pharmacokinetic parameters shown in Table 1. No concomitant medications were taken by any of the study participants. In general, the study medications were adequately tolerated. All subjects experienced marked somnolence during the first day of the olanzapine pharmacokinetic evaluation. Sedation continued into the next day, but was much less pronounced. By the third day after olanzapine ingestion all subjects reported feeling 'normal' again.

CYP1A2 phenotype was analyzed in all 15 subjects. A wide range with three- to eight-fold differences of PMR_{4h},

SMR_{6h}, SMR_{10h}, UMR_{18h}, UMR_{28h}, and UMR_{38h} ratios were observed in the subjects. A summary of the caffeine metabolite ratios is presented in Table 2. The PMR_{4h}, SMR_{6h}, SMR_{10h}, UMR_{18h}, and UMR_{28h} ratios were normally distributed as determined by the Shapiro–Wilk test, while the UMR_{38h} ratio was not. The correlation between PMR_{4h}, SMR_{6h}, SMR_{10h}, UMR_{18h}, UMR_{28h}, and UMR_{38h} ratios are shown in Table 3. All caffeine metabolic

Table 2 Summary of Caffeine Metabolic Ratios (N = 15)

Parameter	Median (range)
PMR _{4h}	0.402 (0.129–0.717)
SMR _{6h}	0.614 (0.186–0.899)
SMR _{10h}	0.892 (0.307–1.37)
UMR _{18h}	2.21 (0.664–5.36)
UMR _{28h}	4.12 (1.24–8.48)
UMR _{38h}	14.1 (9.22–32.0)

Table 3 Correlation Matrix of Caffeine Metabolite Ratios (N = 15)

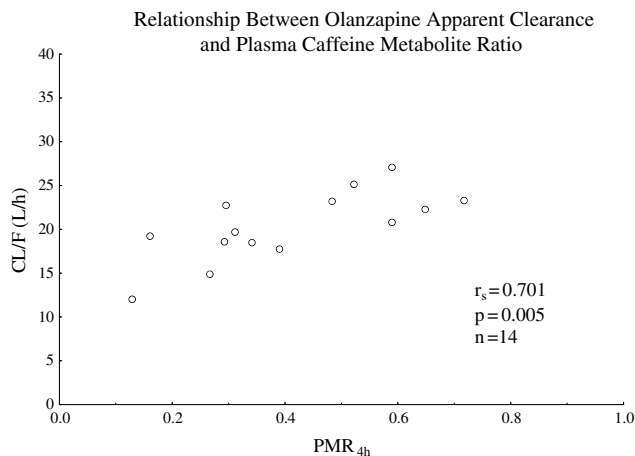
Parameter	PMR _{4h}	SMR _{6h}	SMR _{10h}	UMR _{18h}	UMR _{28h}
PMR _{4h}					
SMR _{6h}	0.871*				
SMR _{10h}	0.861*	0.800*			
UMR _{18h}	0.921*	0.882*	0.761*		
UMR _{28h}	0.925*	0.896*	0.771*	0.968*	
UMR _{38h}	(–)0.321	(–)0.232	(–)0.079	(–)0.354	(–)0.379

* $p < 0.001$, (–) = negative correlation.

Table 4 Correlation Between Olanzapine Clearance (CL/F) and Various Caffeine Metabolite Ratios (N = 14)

Parameter	Olanzapine (CL/F)
PMR _{4h}	0.701*
SMR _{6h}	0.644*
SMR _{10h}	0.701*
UMR _{18h}	0.745*
UMR _{28h}	0.710*
UMR _{38h}	(–)0.029

* $p < 0.05$; (–) = negative correlation.

**Figure 1** Relation between olanzapine apparent clearance and plasma caffeine metabolite ratio.

ratios were significantly correlated to each other except for the UMR_{3h} ratio.

The correlation between olanzapine clearance and caffeine metabolite ratios are presented in Table 4. In spite of the wide intersubject variability in PMR_{4h} and olanzapine clearance (Figure 1; Table 1), these parameters were significantly correlated ($r=0.701$, $p<0.005$) as were all caffeine metabolite ratios with olanzapine clearance except for UMR_{3h}.

DISCUSSION

Similar to previous pharmacokinetic studies conducted by others, our study showed a large intersubject variability in olanzapine disposition (Callaghan *et al*, 1999). Nonetheless, olanzapine pharmacokinetic parameters are similar between the investigations.

Olanzapine has previously been shown to be metabolized by CYP1A2. The other major olanzapine metabolic pathway involves the conversion to olanzapine 10-*N*-glucuronide via the uridine diphosphate glucuronyltransferase (UDPGT) enzyme system (Ring *et al*, 1995). To a minor extent, CYP2D6 and other metabolic enzymes convert olanzapine to its remaining metabolites such as 2-hydroxyolanzapine, olanzapine *N*-oxide, and olanzapine 4'-*N*-glucuronide (Callahan *et al*, 1999). Thus, CYP1A2 appears to play a major role in olanzapine disposition. For example, smoking—a well-known inducer of CYP1A2—contributed to a 40% higher olanzapine clearance among smokers compared to nonsmokers (Callahan *et al*, 1999). Since many psychiatric patients smoke, they should be cautioned that smoking cessation may result in elevated plasma concentrations of olanzapine; increased adverse effects, such as sedation, may result.

Gender is another factor that affects olanzapine disposition. Large population studies have found that women have a 25% lower olanzapine clearance compared to men (Callahan *et al*, 1999). A retrospective investigation examined differential olanzapine plasma concentrations by gender with a fixed-dose regimen (Kelly *et al*, 1999). Women volunteers in this study were found to have significantly higher plasma concentrations of olanzapine than men at five (5) weeks, even with a fixed dose administration. It was proposed that these differences in olanzapine pharmacokinetics were in part because of gender differences in CYP1A2 metabolism, with women having less CYP1A2 activity than men (Kelly *et al*, 1999).

Caffeine metabolism is complex but its biotransformation profile has been clearly determined. In addition to caffeine's metabolism previously described, conversion from paraxanthine (17X) to 1U via the 1X metabolite is mediated by CYP1A2 and xanthine oxidase. Paraxanthine (17X) is converted to an intermediate metabolite by CYP1A2, then to AFMU by *N*-acetyltransferase type 2 enzyme (NAT2), and finally to AAMU (Sinues *et al*, 1999). Paraxanthine is also converted to its 17U metabolite by demethylation and influenced by CYP1A2 and CYP2A6 (El-Yagiki *et al*, 1999). Several concentration ratios of caffeine and its metabolites in plasma, saliva, and urine have been proposed to assess CYP1A2 activity. Whereas plasma and saliva paraxanthine to caffeine ratios have been shown to correlate closely to intrinsic clearance of caffeine 3-demethylation and CYP1A2

content (Fuhr *et al*, 1996), optimal sampling for phenotype analysis is suggested to be 3–6 h after caffeine intake for serum and 6–10 h after caffeine intake for saliva (Spigset *et al*, 1999). The time periods in which plasma and saliva were collected for this study were comparable with those of Spigset, and the caffeine metabolite ratios are very similar to those of previous studies (Table 2) (Fuhr *et al*, 1996; Spigset *et al*, 1999).

Unlike plasma and saliva samples, adequate urine collection interval for the measurement of CYP1A2 activity has not been established. Previous studies used urine samples collected from 4 to 5 h (Chung *et al*, 2000; Lampe *et al*, 2000; Sinues *et al*, 1999) or 0 to 24 h (Bertilsson *et al*, 1994; Sinues *et al*, 1999) after caffeine intake to assess CYP1A2 activity. We collected urine from 0 to 8 h in this study to minimize the effects of urine flow rate on excretion of different metabolites and also to make the sample collection easier. The results that we obtained are different from those of previous studies; however, a direct comparison of results from our study and others is difficult because of different urine collection schedules.

The correlation matrix for caffeine metabolic ratios shown in Table 3 displays significant relations between plasma, saliva and urinary ratios except for UMR_{3h}. These findings are consistent with previous results that plasma ratios significantly correlated with saliva ratios at 3 and 6 h postdose (Fuhr *et al*, 1996). The urinary metabolite ratio UMR_{10-8h} and UMR_{20-8h} in this study significantly correlated with plasma and saliva ratios, and olanzapine apparent clearance, indicating that these ratios can also be used as a reliable marker for CYP1A2 activity in addition to plasma and saliva ratios. A possible explanation for the lack of significant correlation for UMR_{3h} ratio and the other caffeine metabolite ratios is that the metabolic pathways for these metabolites (AAMU and 1U) are also influenced by other enzymes (NAT2 and xanthine oxidase), thereby increasing the variability of metabolite concentrations.

Previously, clozapine clearance was shown to correlate significantly with caffeine disposition, indicating a significant relation between clozapine clearance and CYP1A2 activity (Bertilsson *et al*, 1994). Clozapine clearance was found to be significantly correlated with the 17X/137X ($r=0.89$, $p<0.0013$) and the 37X/137X ratio ($r=0.85$, $p<0.0023$). Very high CYP1A2 activity as indicated by a urinary caffeine metabolic ratio of 17.9 (usual range 1.9–1.47) was reported in one patient; excessive CYP1A2 activity in this patient was associated with a subtherapeutic clozapine plasma concentration of 661 nmol/l (minimum threshold 1150 nmol/l = 350 ng/ml), despite a drug dose of 850 mg/day (Ozdemir *et al*, 2001a). Fluvoxamine 25 mg/day, a well-known CYP1A2 inhibitor, was added to the patient's regimen and a subsequent clozapine plasma concentration of 1600 nmol/l was reached and the patient's clinical status improved. A repeat urinary caffeine ratio was conducted and it was reduced to 9.1 during fluvoxamine coadministration indicating a reduction in CYP1A2 activity with the addition of this agent. Like clozapine, a minimum therapeutic threshold for olanzapine plasma concentrations was identified and found to be 9.3 ng/ml (Perry *et al*, 1997, 2001). It is possible that a very high CYP1A2 activity could reduce olanzapine concentrations and influence a patient's response to the drug. Conversely, patients with low CYP1A2 activity could experience adverse side effects at low to moderate

olanzapine doses because of increased olanzapine exposure. Caffeine phenotyping—a relatively inexpensive and available analytical procedure—could provide an explanation for these clinical situations.

In a recent study, olanzapine clearance did not significantly correlate with CYP1A2 activity when SMR's_{10h} were compared (Hagg *et al*, 2001). Subjects in that study were also phenotyped for CYP2D6 status with 12 patients identified as EM's and 5 as PM's; however, this differentiation had no impact upon CYP1A2 activity and olanzapine clearance. Our findings, where SMR_{6h} and SMR_{10h} caffeine ratios correlated with olanzapine clearance, are not in agreement with these results. In fact, all other caffeine metabolic ratios except UMR_{3h} in our study were found to correlate significantly with olanzapine clearance (Table 4). A possible explanation for disparity between the studies is unclear and warrants further investigation. It was reported that the *N*-desmethyl-olanzapine metabolic ratio significantly correlated with olanzapine clearance ($r^2 = 0.35$, $p < 0.0002$) in large population studies (Callahan *et al*, 1999). This major metabolic pathway for olanzapine is mediated by CYP1A2. Although our study does not extend any ground-breaking new knowledge regarding olanzapine metabolism and CYP1A2, it does confirm the use of caffeine as a phenotypic probe for olanzapine disposition and explains the large intersubject variability found with olanzapine metabolism. Similar results with clozapine and desmethylclozapine metabolite plasma concentrations were found in schizophrenic patients at steady-state conditions (Ozdemir *et al*, 2001b). A significant correlation was reported between clozapine and desmethylclozapine ($r = 0.86$, $p < 0.01$). A significant but negative correlation between desmethylclozapine plasma concentrations and urinary caffeine metabolic ratios were found ($r = (-) 0.76$, $p < 0.01$), indicating that its formation is influenced by CYP1A2.

Limitations to our study include the fact that only one female participated; thus gender influences on olanzapine disposition could not be evaluated. Also, our study did not include phenotyping for CYP2D6 status among the study subjects. Since this is a minor metabolic pathway for olanzapine, the contribution of this information would be minimal and difficult to justify additional study procedures.

CONCLUSIONS

Our study indicates that olanzapine disposition is significantly correlated with CYP1A2 activity as measured by caffeine PMR_{4h}, SMR_{6h}, SMR_{10h}, UMR_{18h}, and UMR_{28h} metabolic ratios. Further studies are needed to compare the influence of gender and smoking upon olanzapine clearance and caffeine metabolic ratios.

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