

ANTITUMOR DRUG ELLIPTICINE INHIBITS THE ACTIVITIES OF RAT HEPATIC CYTOCHROMES P450

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Ellipticine is a potent antineoplastic agent, whose mode of action is considered to be based mainly on DNA intercalation and/or inhibition of topoisomerase II. Recently, we found that ellipticine also forms the cytochrome P450 (CYP)-mediated covalent DNA adducts. Here, we study the effect of ellipticine on CYP enzymes in rat hepatic microsomes, studying its binding to the enzymes and its potential to inhibit the CYP activities measured with their selective substrates. Although ellipticine was reported to be a selective and strong inhibitor of CYP1A1/2, we found that its inhibitory potential is non-specific. Ellipticine is the most potent inhibitor for CYP3A-dependent 6 β -hydroxylation of progesterone, followed by CYP1A1/2-dependent ethoxyresorufin O-deethylation and CYP2B-mediated pentoxyresorufin O-depentylation. Lower inhibition was detected for 1'-hydroxylation of bufurazol, 21-hydroxylation of progesterone and 6-hydroxylation of chlorzoxazone catalyzed by CYP2D, CYP2C and CYP2E1, respectively. Ellipticine binds to several CYPs of rat hepatic microsomes. The binding titration of ellipticine typically give reverse type I spectrum with CYPs in rat hepatic microsomes. The results indicate that inhibition of CYPs by ellipticine cannot be explained only by its differential potency to bind to individual CYPs.

INTRODUCTION

Ellipticine (5,11-dimethyl-6H-pyrido[4,3-b]carbazole) and several of its derivatives isolated from Apocyanaceae plants (i.e. *Ochrosia borbonica*, *Excavatia coccinea*) are alkaloids exhibiting significant antitumor and anti-HIV activities. Ellipticine and its more soluble derivatives (9-hydroxyellipticine, 9-hydroxy-N²-methylellipticinium, 9-chloro-N²-methylellipticinium and 9-methoxy-N²-methylellipticinium) exhibit promising results in the treatment of osteolytic breast cancer metastases, kidney sarcoma, tumors of brain and myeloblastic leukemia (for summary see¹). In order to increase the selectivity of ellipticine antitumor drugs, the attempts to link them to specific vectors able to direct these drugs towards target cells, were performed²⁻⁴. One such conjugate, a heptagastrin fragment linked to ellipticine *via* a spacer has recently been synthesized and shown to be selectively taken up and to be cytotoxic to cells expressing the cholecystokinin type B receptor⁴. The main reason for the interest in ellipticine and its derivatives for clinical purposes is their high efficiencies against several types of cancer, their rather limited toxic side effects and their complete lack of hematological toxicity⁵. However, ellipticine and most of its derivatives are mutagenic compounds (for an overview see¹).

Ellipticines are anticancer drugs, whose precise mechanisms of action have not yet been explained. It was suggested that the prevalent mechanisms of ellipticine antitumor activities are (i) intercalation into DNA (ref.^{6,7})

and (ii) inhibition of DNA topoisomerase II activity⁸⁻¹⁰. Ellipticine and 9-hydroxyellipticine also cause selective inhibition of p53 protein phosphorylation in several human cancer cell lines¹¹, and this correlated with their cytotoxic activity. Ellipticines also uncouple mitochondrial oxidative phosphorylation¹², and thereby disrupt the energy balance of cells.

Recently, we found another mechanism for the ellipticine action^{1,13-16}. We demonstrated that ellipticine covalently binds to DNA after being enzymatically activated. Cytochromes P450 (CYP) are the major enzymes catalyzing the ellipticine oxidation and its activation to more efficient metabolite(s) forming DNA adducts. Using a panel of different human recombinant CYPs - CYP3A4, CYP1A1, CYP1B1 - enzymes expressed at higher levels in tumors sensitive to ellipticine (i.e. breast cancer, renal cell cancer) than in peritumoral tissues¹⁷, were found to be the most effective CYP enzymes activating ellipticine to form covalent DNA adducts *in vitro*¹. The formation of this CYP-mediated covalent DNA adduct by ellipticine was also detected *in vivo*, in rats treated with ellipticine¹⁴. Based on these data, ellipticine might be considered a pro-drug, whose pharmacological efficiency is dependent on its enzymatic activation in target tissues^{1,13-16}.

The discrepancies in data showing oxidation of ellipticine by CYPs and the effects of the drug on these enzymes were observed. On the one hand, ellipticine was found to be a substrate of CYP (ref.^{5,16}) and, moreover, it is also an inducer of several CYPs (ref.¹⁸). On the other hand, this compound was reported to be a strong inhibitor of some

CYPs, selectively inhibiting CYP1A1/2 (ref.⁵). Therefore, the aim of the present work is to extend our knowledge on the ellipticine - CYP interaction and its inhibitory effects on the microsomal mixed function oxidase (MFO) system containing the CYP enzymes.

MATERIAL AND METHODS

Preparation of microsomes and assays

Microsomes were isolated from livers of untreated rats and those of pre-treated with β -naphthoflavone (β -NF), phenobarbital (PB), ethanol (EtOH) and pregnenolon 16 α -carbonitrile (PCN) by procedures as described¹⁸. 7-Ethoxyresorufin O-deethylation (EROD, CYP1A1/2 activity), 7-pentoxoresorufin O-depentylation (PROD, CYP2B activity), progesterone 21-hydroxylation (CYP2C activity), bufuralol 1'-hydroxylation (CYP2D activity), chlorzoxazone 6-hydroxylation (CYP2E1 activity) and progesterone 6 β -hydroxylation (CYP3A activity) were carried out as described^{19,20}. The activity of NADPH:CYP reductase was measured using cytochrome c as substrate (i.e. as NADPH:cytochrome c reductase)¹⁹.

Spectral measurements

The binding of ellipticine to CYPs in microsomes was monitored by difference spectroscopy¹⁹. The concentra-

tion of microsomal protein was adjusted to 1.0 mg per ml of 0.1 M potassium phosphate buffer, pH 7.4. Ellipticine (0.5-10 μ l of methanolic stock solution) was directly added to the sample cuvette containing the enzyme system. The same volume of the solvent was added to the reference cuvette. Absorption spectra were recorded at ambient temperature between 350 and 500 nm. The difference in absorbance between the wavelength maximum and minimum was plotted vs the ellipticine concentration, which was analyzed by nonlinear regression methods with Origin 6.0 software. Two equations were compared statistically to determine the best fit: $\Delta A = B_{\max} S / (K_s + S)$ and $\Delta A = B_{\max} S^n / (K_s^n + S^n)$, where S represents substrate concentration, K_s is the apparent spectral dissociation constant, B_{\max} is the maximal binding, and n is a measure of the cooperativity¹⁹.

RESULTS AND DISCUSSION

Ellipticine acts as an inhibitor of cytochromes P450

Although ellipticine was reported to be a selective and strong inhibitor of CYP1A1/2, we found that its inhibitory potential is non-specific. Control, β -NF-, PB-, ethanol- and PCN-induced microsomal CYP activities are inhibited by ellipticine. A degree of CYP inhibition by the compound was quantified. Ellipticine is the most potent inhibitor

Table 1. Inhibition of reactions catalyzed by individual cytochromes P450 by ellipticine

specific reaction	CYP	microsomes - pre-treatment	IC ₅₀ ellipticine μ M	substrate concentration μ M	inhibition constant K _i μ M	inhibition type
7-ethoxyresorufine O-dealkylation	CYP1A1 CYP1A2	β -NF	0.11	2	0.038 \pm 0.010	NC
7-pentoxoresorufine O-dealkylation	CYP2B	PB	0.11	2	0.050 \pm 0.020	mixed C/NC
progesterone 21-hydroxylation	CYP2C	control	2	50	*)	
bufuralol 1'-hydroxylation	CYP2D	control	0.5	50	0.24 \pm 0.03	mixed C/NC
chlorzoxazone 6-hydroxylation	CYP2E1	EtOH	140	50	*)	
progesterone 6 β -hydroxylation	CYP3A	PCN	0.5	100	0.021 \pm 0.01	mixed C/NC

IC₅₀ concentration of ellipticine causing 50 % inhibition

C competitive inhibition

NC non-competitive inhibition

* for higher values of IC₅₀ the inhibition constant was not determined

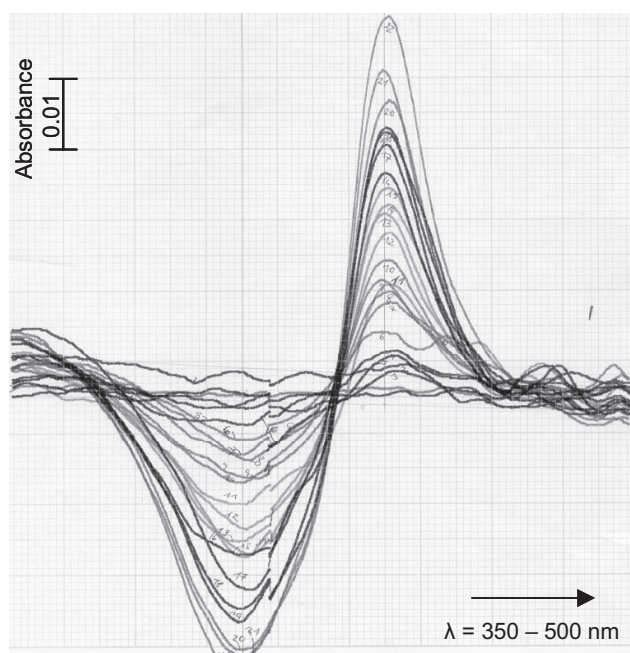


Fig. 1. Ellipticine exhibits a reverse type I binding spectrum with rat hepatic microsomal CYPs (herein with PCN microsomes)

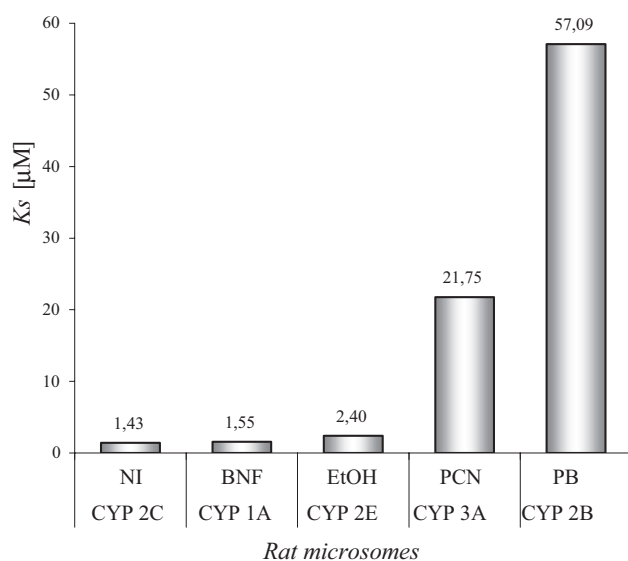


Fig. 2. Apparent dissociation constant (K_s) values reflecting the affinity of ellipticine to microsomal CYPs. NI, uninduced; BNF, β -naphthoflavone; EtOH, ethanol; PCN, pregnenolone 16 α -carbonitrile; PB, phenobarbital.

for CYP3A-dependent 6 β -hydroxylation of progesterone ($K_i=0.021 \mu\text{M}$), followed by CYP1A1/2-dependent EROD activity ($K_i=0.038 \mu\text{M}$) and CYP2B-mediated PROD activity ($K_i=0.05 \mu\text{M}$). Lower, but measurable, inhibition was detected for 1'-hydroxylation of bufurazol, 21-hydroxylation of progesterone and 6-hydroxylation of chlorzoxazone catalyzed by CYP2D, CYP2C and CYP2E1, respectively. Ellipticine acts as a non-competitive or mixed-type inhibitor of these enzymes (Table 1).

It should be noted that the interpretation of the results from the inhibition studies is sometimes difficult, because the inhibitor may be more effective with one substrate than another. Therefore, to explain the mechanism of inhibition of CYPs by ellipticine, the binding of ellipticine to microsomal CYPs was examined.

Binding of ellipticine to CYPs in microsomes

Ellipticine binds to microsomal CYPs with affinity greater than most of other compounds known to interact with these enzymes. It binds to CYPs of microsomes isolated from livers of uninduced rats and from those of rats pre-treated with β -NF (enriched with CYP1A1/2), PB (enriched with CYP2B1/2), PCN (enriched with CYP3A1/2) and ethanol (enriched with CYP2E1). The binding titration of ellipticine typically give reverse type I spectrum (λ_{max} 430 nm) with CYPs in all microsomes tested in the study, measured at 37 $^\circ$ C (see Fig. 1 for PCN microsomes). This finding indicates that ellipticine interacts both with Fe of the heme prosthetic group of CYP and with the amino acid residues present in the active centre of the enzyme. The magnitude of the difference of the absorbance maximum and minimum for both compounds is assumed to represent the extent of their binding to the catalytic site. The values of the spectral dissociation constants (K_s) for the complexes of ellipticine with CYPs in microsomes, reflecting the affinity of ellipticine to these CYPs are shown in Fig. 2.

As shown from Table 1 and Figure 2, the degree of the CYP inhibition by ellipticine does not correlate with the potency of its binding to the individual CYP enzymes. These results indicate that degrees of inhibition by ellipticine might be explained not only by its differential potency to bind to individual CYP, but also by its effect on another member of the MFO system, e.g. NADPH:CYP reductase. Even though the inhibition of reduction of a non-physiological substrate of NADPH:CYP reductase, cytochrome *c*, was not detected in the presence of ellipticine up to concentrations of 0.5 mM (results not shown), the inhibition of this enzyme using the CYP enzyme as a substrate cannot be excluded. Therefore, the question of the exact mechanism responsible for CYP inhibition by ellipticine remains still to be answered.

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