Differential Inhibition of Cytochromes P450 3A4 and 3A5 by the Newly Synthesized Coumarin Derivatives 7-Coumarin Propargyl Ether and 7-(4-Trifluoromethyl)coumarin Propargyl Ether

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Abstract:

The abilities of 7-coumarin propargyl ether (CPE) and 7-(4-trifluoromethyl)coumarin propargyl ether (TFCPE) to act as mechanism-based inactivators of P450 3A4 and 3A5 in the reconstituted system have been investigated using 7-benzyloxy-4-(trifluoromethyl)coumarin (BFC) and testosterone as probes. CPE inhibited the BFC O-debenzylation activity of P450 3A4 in a time-, concentration-, and NADPH-dependent manner characteristic of a mechanism-based inactivator with a half-maximal inactivation ($k_{inact}$) of 0.04 min$^{-1}$, and a $t_{1/2}$ of 13.9 min. Similarly, TFCPE inhibited the BFC O-debenzylation activity of P450 3A4 in a time-, concentration-, and NADPH-dependent manner with a $K_i$ of 14 μM, a $k_{inact}$ of 0.04 min$^{-1}$, and a $t_{1/2}$ of 16.5 min. Parallel losses of P450 3A4 enzymatic activity and heme were observed with both compounds as measured by high-performance liquid chromatography and reduced CO spectra. Interestingly, neither compound inhibited the BFC O-debenzylation activity of P450 3A5. Reactive intermediates of CPE and TFCPE formed by P450 3A4 were trapped with glutathione, and the resulting adducts were identified using tandem mass spectral analysis. Metabolism studies using TFCPE resulted in the identification of a single metabolite that is formed by P450 3A4 but not by P450 3A5 and that may play a role in the mechanism-based inactivation.

The cytochrome P450-dependent mixed-function oxidases are the main enzymes involved in the NADPH-dependent metabolism of various xenobiotics to more polar products that can be more readily excreted. In humans, the P450 3A family of enzymes is involved in the metabolism of 50 to 60% of all currently used drugs (Evans and Relling, 1999). Four major cytochrome P450 (P450) 3A genes have been identified, CYP3A4, CYP3A5, CYP3A7, and CYP3A43. P450 3A4 is generally thought to be the predominant form expressed in the human liver (Wrighton et al., 2000). P450 3A5 can contribute as much as 50% to the hepatic 3A content in one-third of Caucasians and in one-half of African Americans (Kuehl et al., 2001). P450 3A7 has been shown to be a fetal form of 3A, and little is known about the function of P450 3A43. P450s 3A4 and 3A5 share 84% amino acid sequence homology, and thus considerable overlap exists between their substrate specificities (Wrighton and Stevens, 1992). On the other hand, there has also been evidence for preferential metabolism or inactivation of one enzyme over the other. The two enzymes have been shown to exhibit different regioselectivities toward aflatoxin (Gillam et al., 1995b), increased metabolism of midazolam, tacrolimus, and vincristine by P450 3A5 is observed compared with that by P450 3A4 (Wandel et al., 1994; Kamdem et al., 2005; Dennison et al., 2006), and differential inactivation of P450 3A4 and not 3A5 by raloxifene has also been reported (Baer et al., 2007).

Coumarin is a constituent of many plants, microorganisms, and animals and is widely used as a fragrance in various products such as cosmetics, soaps, and detergents (Soine, 1964). Coumarins are composed of an aromatic ring fused to a lactone ring and have been shown to possess a range of pharmacological and biochemical properties. Coumarin has been used in combination therapy with cimetidine in clinical trials for the treatment of malignant melanomas, metastatic renal carcinomas, and carcinomas of the prostate (Marshall et al., 1989). A number of studies have suggested that coumarin is not toxic to humans; however, species differences exist that could account for variations in both metabolism and toxicity. For example, administration of coumarin to rodents has been shown to produce liver and lung toxicity (Born et al., 1998). Coumarin and coumarin derivatives are known to be substrates for cytochrome P450s (Lake, 1999). Buters et al. (1993) used this property to develop a sensitive assay to

ABBREVIATIONS: P450, cytochrome P450; CPE, 7-coumarin propargyl ether; TFCPE, 7-(4-trifluoromethyl)coumarin propargyl ether; BFC, 7-benzyloxy-4-(trifluoromethyl)coumarin; GC, gas chromatography; MS, mass spectrometry; NMR, nuclear magnetic resonance; HPLC, high-performance liquid chromatography; LC, liquid chromatography; MS/MS, tandem mass spectrometry; APCI, atmospheric pressure chemical ionization; SRS, substrate recognition site; CPE-GS, CPE-glutathione conjugate.
measure the activities of a number of cytochrome P450s by which the O-deethylation of 7-ethoxy-(4-trifluoromethyl)coumarin results in the formation of the fluorescent 7-hydroxy-(4-trifluoromethyl)coumarin product. Thus, coumarin and coumarin derivatives are currently widely used in the drug discovery process.

Many acetylenic compounds have been shown to inactivate various cytochrome P450 enzymes (Roberts et al., 1997; Shimada et al., 1998; Strobel et al., 1999) in a mechanism-based manner that involves binding of a reactive intermediate to either the heme or the apoprotein (Ortiz de Montellano and Kunze, 1980). 2-Ethynylnaphthalene has been shown to inactivate P450s 2B1 and 2B4 through a modification of the apoprotein (Roberts et al., 1994), whereas tert-butyl acetylene and tert-butyl 1-methyl-2-propynyl ether have been shown to inactivate P450s 2E1 and 2B1, respectively, through the formation of both heme and protein adducts (Blobaum et al., 2002; Von Weymarn et al., 2004).

Naphthoﬂavones serve as substrates for a number of P450 enzymes (Cho et al., 2003), and the propargyl ether derivatives of naphthoﬂavones and adamantanes have been shown to be inhibitors of some P450 enzymes. The purpose of this project was to evaluate compounds that combine both the structural motif of P450 3A substrates and the activity of P450 enzymes. The corresponding starting material (0.027 mol of 7-hydroxycoumarin or 7-hydroxy-(4-(trifluoromethyl))coumarin purchased from Sigma-Aldrich) was dissolved in 30 ml of dry tetrahydrofuran under an atmosphere of N2. Fresh sodium hydride (3 Eq, 0.081 mol, 1.94 g) was added slowly, followed by 2 Eq of propargyl bromide (0.054 mol, 6.01 ml). The reaction mixture was left to stir for 1 week under N2 and its progress was monitored daily by thin-layer chromatography. The reaction was quenched with 50 ml of deionized water and extracted twice with 30-ml portions of methylene chloride. The organic layers were combined and washed with 10% HCl followed by water (twice). The crude material was then dried over anhydrous MgSO4, and the residual solvent was evaporated under vacuum. The crude product was puriﬁed by flash silica gel column chromatography using petroleum ether as the solvent. The pure fractions were combined, and the puriﬁed product was recrystallized from ethanol and water.

CPE. The yield after puriﬁcation was 90.5%. GC/MS showed >99% purity; m/z (%): 209 (19), 51 (100). 1H NMR (CDCl3): δ 2.60 (s, 1H), 4.75 (s, 2H), 6.28 (1H), 6.92 (d, 1H), 6.95 (s, 1H), 7.40 (d, 1H), and 7.65 (d, 1H). 13C NMR (CDCl3): δ 25.60, 56.20, 71.00, 102.13, 113.06, 113.18, 113.84, 128.84, 129.24, 129.46, 156.13, 156.59, 161.26, and 161.02. TFCPE. The yield after puriﬁcation was 25.3%. GC/MS showed >99% purity; m/z (%): 268 (1), 202 (100). 1H NMR (CDCl3): δ 2.60 (s, 1H), 4.80 (s, 2H), 6.66 (s, 1H), 6.99 (s, 1H), 7.01 (d, 1H), and 7.65 (d, 1H). 13C NMR (CDCl3): δ 25.60, 56.31, 67.98, 102.71, 107.74, 112.81, 113.87, 113.84, 122.94, 126.46, 155.64, 160.53, and 161.02.

**Materials and Methods**

Materials. 1-α-Dilaurylphosphatidylcholine, 1-α-dioleyl-sn-glycero-3-phosphocholine, phosphatidylerine, bovine serum albumin, NADPH, Tris, catalase, sodium dithionite, and Hepes were purchased from Sigma-Aldrich (St. Louis, MO). 7-Benzylxoy-4-(trifluoromethyl)coumarin (BFC) was obtained from BD Gentest (Woburn, MA). 7-Ethoxy-4-(trifluoromethyl)coumarin (EFC) was obtained from St. Louis, MO. 7-Benzylxoy-4-(trifluoromethyl)coumarin (BFC) was obtained from BD Gentest (Woburn, MA). 7-Ethoxy-4-(trifluoromethyl)coumarin was obtained from Invitrogen (Carlsbad, CA). AccuBond SPE ODS-C18 cartridges were obtained from Agilent Technologies (Wilmington, DE). All other chemicals were of the highest quality commercially available. The human liver microsomes used in the studies have been described previously (Teiber and Hollenberg, 2000).

**Purification of Enzymes.** P450s 3A4 and 3A5 were expressed in Escherichia coli DH5α cells and puriﬁed to homogeneity by the method of Gillam et al. (1993). P450 NADPH-reductase was expressed in E. coli Topp3 cells and puriﬁed according to a previously published protocol (Hamma et al., 1998). Cytochrome bs was puriﬁed from liver microsomes of rats (Lin et al., 2002).

**Synthesis of Coumarin Propargyl Ethers.** The corresponding starting

![chemical structures of CPE and TFCPE](image)
was injected onto an Aqua C18 column (150 mm OD; ODS-C18 1-ml cartridges. After application of the samples, the cartridges were washed with 2 ml of 15% trichloroacetic acid. The HPLC system consisted of 501 pumps, a 600E controller, and a 717A autosampler (Waters). The components of the reconstructed mixtures were resolved by linearly increasing the percentage of acetonitrile to 90% over 30 min. Heme was quantified by integrating the area under the peaks that absorbed at 405 nm, and mean and S.D. were calculated from four different experiments. Reduced CO spectra of the control, exposed, and inactivated samples were obtained by transferring 100 pmol of each sample into 900 μl of quench buffer containing 50 mM potassium phosphate (pH 7.4), 40% glycerol, and 0.6% Tergitol. The samples were bubbled with CO, sodium dithionite was added, and the reduced CO spectra were recorded between 400 and 500 nm on a DW2 UV/visible spectrophotometer equipped with an Otis operating system (On Line Instruments Systems, Bogart, GA).

### Heme Analysis.
P450 3A4 was reconstructed and inactivated as described above for the activity assays. Samples (100 pmol) of the control and inactivated enzymes were chromatographed on a C4 column (250 × 4.6 mm; Phenomenex, Torrance, CA) equilibrated with 30% acetonitrile containing 0.1% trifluoroacetic acid. The HPLC system consisted of 501 pumps, a 600E controller, and a 717A autosampler (Waters). The components of the reconstructed mixtures were resolved by linearly increasing the percentage of acetonitrile to 90% over 30 min. Heme was quantified by integrating the area under the peaks that absorbed at 405 nm, and mean and S.D. were calculated from four different experiments. Reduced CO spectra of the control, exposed, and inactivated samples were obtained by transferring 100 pmol of each sample into 900 μl of quench buffer containing 50 mM potassium phosphate (pH 7.4), 40% glycerol, and 0.6% Tergitol. The samples were bubbled with CO, sodium dithionite was added, and the reduced CO spectra were recorded between 400 and 500 nm on a DW2 UV/visible spectrophotometer equipped with an Olis operating system (On Line Instruments Systems, Bogart, GA).

### GSH Adducts.
Samples were reconstituted using 0.5 nmol of P450 3A4 as described previously and incubated in the presence of 10 mM GSH. Reaction conditions and inactivation conditions were the same as described earlier. The reactions were stopped by the addition of 60 μl of 10% trifluoroacetic acid/ml of sample and vortexed. The samples were then applied to AccuBond SPE ODS-C18 1-ml cartridges. After application of the samples, the cartridges were washed with 2 ml of water followed by elution of the GSH-CPE or GSH-TFCPE adducts with 2 ml of methanol followed by 0.3 ml of acetonitrile. The methanol and acetonitrile fractions were combined, dried under nitrogen, and suspended in 50% acetonitrile and 0.5% acetic acid. One part of the sample was injected onto an Aqua C18 column (150 mm × 4.6 mm, 5 μm; Phenomenex) equilibrated with 5% acetonitrile and 0.1% acetic acid. The sample components were resolved by eluting at 0.3 ml/min while the percentage of acetonitrile was linearly increased to 90% over the course of 70 min. The LC effluent was directed into a mass spectrometer (LCQ ion trap; Thermo Fisher Scientific, Waltham, MA) and MS spectra as well as MS/MS spectra were recorded using the data-dependent scanning routine of the Xcalibur data system (Thermo Fisher Scientific). The analyzer conditions were as follows: sheath gas, 90 arbitrary units; auxiliary gas, 30 arbitrary units; spray voltage, 4.5 kV; capillary temperature, 170°C; and capillary voltage, 30 V. One full scan between m/z 150 and 1000 was collected followed by two dependent scans between m/z 50 and 1000 at 35% relative collision energy, with an activation Q of 0.250 and an activation time of 30 ms. The second part of the sample was injected into a LC column coupled to a fluorescence detector with excitation at 350 nm and emission at 454 nm. The HPLC system consisted of an LC-20AB liquid chromatograph, autosampler SIL-20AC HT, and fluorescence detector RF-10Axl (Shimadzu). The analysis was performed using a mobile phase of acetonitrile and 0.1% acetic acid (phase B) and water and 0.1% acetic acid (phase A) and a flow rate of 0.3 ml/min. Initial conditions were 5% B, and the percentage of B was maintained for 5 min followed by a linear gradient to 30% B from 5 to 15 min to 80% B from 15 to 35 min and to 90% B from 35 to 40 min. The column was washed with 90% B for 15 min before a return to the initial conditions and equilibration for 10 min at the initial conditions before the next injection.

### Metabolism of CPE and TFCPE.
The formation of 7-hydroxycoumarin was determined to see whether P450s 3A4 and 3A5 were able to metabolize the coumarin compounds. This was done by reconstituting P450s 3A4 or 3A5 in the presence of CPE or TFCPE as described above. The reactions were allowed to proceed for 30 min after they were initiated by the addition of 1 mM NADPH. The reaction mixture was then stopped with 70 μl of 15% trichloroacetic acid. The formation of 7-hydroxycoumarin was analyzed by HPLC using a C18 Microsorb-MV column coupled to a fluorescence detector with excitation at 350 nm and emission at 453 nm. The analysis was performed by using an isocratic mobile phase of 30% methanol with 0.2% acetic acid. The flow rate was set to 1 ml/min. Quantitation was done using peak areas by comparing to a standard curve generated with 7-hydroxycoumarin.

### LC-APCI-MS and Metabolite Analysis.
For the metabolism of TFCPE by P450s 3A4 and 3A5, the reconstitution and reaction conditions were as described earlier. The control and the inactivated samples (900 pmol each) were extracted with 2 ml of ethyl acetate, and the organic phases were dried under nitrogen. The dried samples were resuspended in 100 μl of 50% acetonitrile for chromatographic separation. Chromatographic separation of the
metabolites was performed on a Surveyor HPLC system equipped with a quaternary pump, an autosampler, and a diode array detector. The column used was a C18 (either a Microsorb-MV 100 Å with a flow rate of 1 ml/min or a Phenomenex Jupiter 300 5 μm H9262 with a flow rate of 0.2 ml/min) reverse-phase column, equilibrated with 90% A (water and 0.1% acetic acid) and 10% B (acetonitrile and 0.1% acetic acid). Metabolites were resolved by increasing the percentage of B as follows: from initial conditions to 30% over 3 min, to 50% B over 5 min, holding at 50% B for 2 min, increasing to 70% B over 3 min and to 90% B over 5 min, and then holding at 90% B for 2 min before returning to initial conditions. A volume of 50 μl was injected for each analysis. The elution of the parent compound and the metabolites from the chromatographic column was monitored at 319 nm. Effluent from the HPLC was directed into the APCI source of a linear ion trap (LTQ; Thermo Fisher Scientific) scanning in the positive ion mode. The ionization source conditions were as follows: APCI vaporizer temperature, 450°C; corona discharge voltage, 6.0 kV; and tube lens offset voltage, 60 V. Nitrogen was used as the sheath gas (55 psi) and auxiliary gas (15 psi). Full mass spectra were recorded between m/z 100 and 2000 and data-dependent scanning was used to collect MS/MS spectra of the four most intense ions in each full scan.

Results

Inactivation Assay. In the presence of CPE or TFCPE, a time-, concentration-, and NADPH-dependent inactivation of the BFC O-debenzylation activity of P450 3A4 in the reconstituted system was observed. Pseudo first-order kinetics was observed for concentrations of CPE between 25 and 500 μM (Fig. 2) and for concentrations of TFCPE between 5 and 100 μM (Fig. 3). As calculated from the data

<table>
<thead>
<tr>
<th>P450 3A4</th>
<th>Activity Remaining</th>
<th>P450 Remaining by Reduced CO Spectra</th>
<th>Residual Heme by HPLC</th>
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<tr>
<td></td>
<td>0 min 20 min</td>
<td>0 min 20 min</td>
<td></td>
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<tr>
<td>CPE, NADPH</td>
<td>100 89 ± 2</td>
<td>100 92 ± 3</td>
<td>100</td>
</tr>
<tr>
<td>+CPE, +NADPH</td>
<td>86 ± 5</td>
<td>69 ± 7</td>
<td>70 ± 11</td>
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<tr>
<td>-TFCPE, -NADPH</td>
<td>88 ± 4</td>
<td>75 ± 4</td>
<td>100 ± 4</td>
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<tr>
<td>+TFCPE, +NADPH</td>
<td>88 ± 3</td>
<td>75 ± 6</td>
<td>100 ± 4</td>
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**TABLE 2**

Effect of CPE and TFCPE on P450 3A4 BFC O-debenzylation activity, reduced CO spectra, and residual heme as measured by HPLC

Assay conditions are described under Materials and Methods. Activity remaining was determined using 50 μM CPE or 12 μM TFCPE with an incubation time of 20 min. The amount of P450 remaining was measured using reduced CO spectra. The amount of residual heme by HPLC was calculated from the area under the heme peak as determined by absorbance at 405 nm. The area for the control sample was set to 100%. The data represent the mean and S.D. from four different experiments.

Fig. 4. Mass spectral analysis of the adduct of CPE with glutathione. P450 3A4 was reconstituted with reductase and lipid and incubated with CPE in the presence of GSH, and the samples were analyzed as described under Materials and Methods. A, total ion chromatogram showing a primary peak eluting at 22 min. B, monitoring of the column effluent fluorometrically shows elution of the fluorescent GSH adduct at 22.2 min. C, mass spectrum of the peak eluting at 11.8 with an (M+H) ion at 523.9. D, MS/MS of the 523.9 ion. E, proposed structure of the CPE-GS adduct showing the primary fragmentations.

In the presence of CPE or TFCPE, a time-,
in Fig. 2, the concentration required for \( K_I \) by CPE was 112 \( \mu M \), the \( k_{\text{inact}} \) was 0.05 min\(^{-1}\), and the \( t_{1/2} \) was 13.9 min (Table 1). The \( K_I \) for TFCPE was 14 \( \mu M \), the \( k_{\text{inact}} \) was 0.04 min\(^{-1}\), and the \( t_{1/2} \) was 16.5 min (Table 1). In contrast to P450 3A4, incubations of P450 3A5 with CPE or TFCPE in the presence of NADPH showed no inactivation of the BFC \(-\)-debenzylation activity (data not shown). Incubation with either CPE or TFCPE did not result in inactivation of both samples of human liver microsomes (data not shown).

The loss in the 6\( R \)-testosterone hydroxylation activity of P450 3A4 when incubated with CPE or TFCPE in a reconstituted system was also investigated and was found to be time-, concentration-, and NADPH-dependent. The inactivation was characterized by a \( K_I \) of 25 \( \mu M \), a \( k_{\text{inact}} \) of 0.014 min\(^{-1}\), and a \( t_{1/2} \) of 50 min for CPE and a \( K_I \) of 17 \( \mu M \), a \( k_{\text{inact}} \) of 0.04 min\(^{-1}\), and a \( t_{1/2} \) of 17 min for TFCPE (Table 1).

**Heme Analysis.** The effects of inactivation by CPE and TFCPE on the amount of P450 3A4 heme remaining were studied by determining the reduced CO spectra of the inactivated proteins and by determining the amount of heme remaining by comparing the area under the heme absorbance at 405 nm from chromatograms of control and the inactivated samples resolved by HPLC. The inactivation of P450 3A4 by CPE and TFCPE after a 20-min incubation resulted in approximate losses of 35 and 55% in \(-\)-debenzylation activity, respectively, compared with 0 min (Table 2). Some decrease in activity is seen in 0-min samples with the inactivator and with or without the NADPH and also in the samples with inactivators, but without NADPH. Presumably this decrease in activity is due to carryover of the inactivator from the initial reaction mixture to the assay mixture where it can act as a competitive inhibitor. These losses in the activity were accompanied by losses in the ability of the inactivated samples to form reduced CO spectra. Losses in the levels of the P450 heme as measured by the reduced CO spectra were also seen in control samples that were incubated only with inactivator. The reasons for this result are not clear. As shown in Table 2, the loss in heme was approximately 30% compared with the −NADPH control in samples incubated with CPE, suggesting that the inactivation was partly due to heme modification. For the inactivation of 3A4 by TFCPE, the loss in activity was approximately 55% with the −NADPH +TFCPE control compared with a 40% loss of heme as measured by HPLC. This suggests that the inactivation could be due partly to heme modification. In each case when the residual heme was measured by HPLC, a small peak eluting at 23.4 min corresponding to a heme adduct was detected in the inactivated samples but not in the control samples (data not shown). LC-MS analysis of the adducted hemes did not reveal the mass of these adducts.

**Analysis of GSH Adducts.** Incubations of P450 3A4 with CPE or TFCPE in the presence of GSH resulted in the formation of CPE- and TFCPE-glutathione adducts (Figs. 4 and 5). Figure 4A shows the total ion chromatogram of the CPE-GSH adduct with the primary ion eluting at 22 min that was obtained from samples incubated with CPE and GSH in the presence of NADPH. The peak detected by the fluorescence detector eluting at the same retention time is shown in Fig. 4B. The mass spectrum of the peak eluting at 22.2 min is
presented in Fig. 4C. The MS/MS spectrum of the ion with an m/z of 523.9 shown in Fig. 4D exhibited major ions at m/z 448.7 [MH-H11002/75] and 394.7 [MH-H11001/129] arising from the parent ion at 523.9 by a neutral loss of glycine and pyroglutamate, respectively, which is characteristic of glutathione-conjugated metabolites. The fragment ion at m/z 307.7 is from the protonated GSH moiety arising from the cleavage of the thioester bond. The fragments with m/z 319.0 and 291.7 are from the combined loss of glycine and glutamate and glycine, glutamate, and CO, respectively. The fragment with m/z 216.8 results from the cleavage of the C–S bond between the GSH moiety and the coumarin propargyl ether. The ion at m/z 161.8 is most likely obtained from the sum of coumarin and oxygen, whereas the ion at m/z 178.8 could be from the combined mass of coumarin, oxygen, and a methyl group. The fragmentation pattern for the major ions observed in the MS/MS spectrum of the CPE-glutathione conjugate (CPE-GS) is shown in Fig. 4E. Figure 5A shows the total ion chromatogram of the TFCPE adduct with GSH showing a primary ion eluting at 31.31 min. The peak eluting at approximately 31 min as seen by the fluorescence detector is shown in Fig. 5B. The mass spectrum of the peak eluting at 31.3 min is shown in Fig. 5C. Figure 5D shows the MS/MS spectrum of the ion with m/z 591.9. The fragment ion at m/z 573.9 is from the loss of water from the TFCPE-GSH molecule. The daughter ions with m/z values of 516.9 [MH-H11002/75], 462.8 [MH-H11001/129], and 444.8 [MH-H11001/146] correspond to losses of glycine, pyroglutamate, and pyroglutamate and a water molecule, respectively. The ion with m/z 426.9 resulted from a subsequent loss of water from the 444.8 ion. The ion with m/z 359.8 resulted from the loss of glycine, glutamate, and CO and the 341.8 ion from the further loss of water. The ion with m/z 307.6 represents the mass of GSH and was also seen in the CPE-GSH adduct profiles. The ions with m/z 268.8 and 242.5 correspond to the parent [M+H]+ molecule and to a loss of HC≡CH, respectively. The fragmentation pattern for the major ions of the TFCPE-GS adduct observed in the MS/MS spectrum is shown in Fig. 5E.

**Metabolism of CPE and TFCPE.** The formation of 7-hydroxycoumarin as a result of O-deethylation of CPE or TFCPE by 3A4 or

<table>
<thead>
<tr>
<th>P450</th>
<th>Formation of 7-Hydroxycoumarin pmol/pmol P450/min</th>
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<tbody>
<tr>
<td>3A4</td>
<td>CPE 48.1 ± 5</td>
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<tr>
<td></td>
<td>TFCPE 97.2 ± 1</td>
</tr>
<tr>
<td>3A5</td>
<td>CPE 85.5 ± 7</td>
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<tr>
<td></td>
<td>TFCPE 78.8 ± 11</td>
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**TABLE 3**

Formation of 7-hydroxycoumarin by P450s 3A4 and 3A5

The P450s were reconstituted in the presence of CPE or TFCPE as indicated under Materials and Methods. The formation of 7-hydroxycoumarin was measured by HPLC analysis with fluorescence detection. The data points represent the mean and average from three different experiments.
3A5 was measured by reverse-phase HPLC with fluorescence detection. As shown in Table 3, both P450s 3A4 and 3A5 were able to metabolize both of the coumarin propargyl ether compounds to give 7-hydroxycoumarin. The amount of 7-hydroxycoumarin formed by 3A4 in presence of CPE was approximately half of what was seen with TFCPE. With P450 3A5, the amount of 7-hydroxycoumarin formed was approximately the same with both propargyl compounds. Overall, the rate of formation of 7-hydroxycoumarin by P450 3A5 suggests that both coumarin compounds bind to 3A5 as well as to 3A4 and are readily metabolized by 3A5 even though they do not inactivate it.

**LC-APCI-MS and MS/MS Analysis of the Metabolites of TFCPE by P450 3A4, TFCPE and its metabolites were monitored at 319 nm. Figure 6A shows the UV trace of the effluent from the HPLC that was directed into the atmospheric pressure chemical ionization. The total ion chromatogram revealed two distinct peaks: the parent peak eluting at 22.78 min and a metabolite of TFCPE eluting at approximately 20.3 min. This metabolite peak was seen only in the samples incubated with P450 3A4 and not in the samples incubated with P450 3A5. The inset shows the MS spectrum of the parent peak eluting at 22.78 min which revealed an intense [M + H]+ ion with m/z 269.0 expected for TFCPE. MS/MS analysis of the parent compound generated fragment ions at m/z 240.9 from loss of CO and m/z 213 due to cleavage of the propargyl ether. The fragmentation pattern of the parent peak is shown in Fig. 6B. The single metabolite peak eluting at 20.34 min was only observed in 3A4 samples incubated with NADPH and not in control samples (Fig. 7A). The mass spectrum of this peak showed ions with m/z values at 271.6 and 231.0 as shown in Fig. 7A. The fragmentation of the ion with m/z 271.6 yielded daughter ions at m/z 253.3 from the loss of water, at m/z 242.99 from the loss of CO, and at m/z 214.33 from the loss of C3H4 (Fig. 7B). We are unable to explain the origin of the ions with the m/z of 197.10 and 161.25. The fragmentation of the ion at m/z 231.0 yielded a daughter ion with a m/z 186.95, which is derived from the ring opening of the compound with a subsequent loss of CO2 (Fig. 7E). At present we are unable to explain the origin of the ion with m/z 166.91. The fragmentation patterns for these metabolites are shown in Fig. 7, C and E. No similar peak or any other metabolite peak was observed under these conditions for CPE. The incubation of P450 3A5 with TFCPE did not result in any metabolite as seen by LC-APCI-MS, suggesting that the product formed by P450 3A4 is unique to this enzyme.

**Discussion**

The importance of human P450 3A enzymes arises from their abundant expression and their role in the first-pass metabolism of many clinically important drugs. Until recently, P450 3A4 was thought to be the major human drug-metabolizing enzyme. However, it has now been shown that P450 3A5 may represent more than 50% of the total P450 3A content in some individuals (Kuehl et al., 2001). The contribution of P450 3A5 to the metabolic clearance of P450 3A substrates in the liver has not been well characterized. Both P450s 3A4 and 3A5 show similar Km and Vmax values for midazolam 1’-hydroxylation (Gibbs et al., 1999). Both enzymes metabolize nifedipine (Gillam et al., 1995), lidocaine (Bargetzi et al., 1989), and dextromethorphan (Gorski et al., 1994). However, a few studies have shown marked differences in the inhibition kinetics between P450s 3A4 and 3A5. For example, significant differences have been seen for the mechanism-based inactivation of P450s 3A4 and 3A5 by mifepristone where P450 3A5 did not exhibit irreversible inhibition (Khan et al., 2002). Recently, verapamil was shown to be a selective inhibitor of P450 3A4 (Wang et al., 2005). In the search for additional specific probes that are capable of differentiating between these two major human 3A enzymes, two coumarin propargyl ether compounds were synthesized.

In this investigation, incubation of P450 3A4 with two structurally similar propargyl ethers, CPE and TFCPE, resulted in the loss of BFC O-debenzylation activity in a time- and concentration-dependent manner. The inactivation of P450 3A4 by CPE and TFCPE was characterized by KI values of 112 and 14 μM, kinact values of 0.05 and 0.04 min⁻¹, and half-times of 13.9 and 16.5 min, respectively (Table 1). Because mechanism-based inactivation can be substrate-dependent and studies have shown that P450 3A4 can accommodate more than one of the same or different substrate molecules in its active site (Korzekwa et al., 1998), we used testosterone as an additional substrate to measure the effect of the two propargyl ethers on the hydroxylation activity of P450 3A4. CPE and TFCPE both caused a loss of the 6β-testosterone hydroxylation activity of P450 3A4 in a time-, concentration-, and NADPH-dependent manner and demonstrated pseudo-first-order kinetics. The inactivations were characterized by KI of 25 μM, kkinact of 0.014 min⁻¹, and a t1/2 of 50 min for CPE and KI of 17 μM, kkinact of 0.04 min⁻¹, and a t1/2 of 17 min for TFCPE. This type of substrate-dependent effect on P450 3A4 inhibition has been shown before (Kenworthy et al., 1999). Wang et al. (2000) have also shown that 3A4 inhibition as measured by different probes such as testosterone, nifedipine, and midazolam is substrate-dependent. Thus, differences in the kinetic parameters seen with CPE could be substrate-dependent. Interestingly, neither CPE nor TFCPE showed any effect on the BFC activity of P450 3A5. Because P450 3A5 does not efficiently catalyze the hydroxylation of testosterone (White et al., 1990) or progesterone (Aoyama et al., 1989), we did not investigate the effects of CPE or TFCPE on steroid metabolism by 3A5. Although neither CPE nor TFCPE compounds resulted in the loss of BFC activity, they both were O-dealkylated to form 7-hydroxycoumarin by 3A4 and 3A5, as detected by HPLC with fluorescence detection (Table 3).

These results suggest that metabolism of both of the propargyl ethers by P450 3A5 occurs but may not produce the reactive intermediate necessary to bring about a loss in function of this enzyme. Several reports have shown differences in the inactivation or metabolism of various substrates by P450 3A4 and 3A5. Similar findings have been identified when P450 3A5 has been shown to oxidize vincristine and tacrolimus more efficiently than 3A4 (Dennison et al., 2006) and differential inactivation of 3A4 and 3A5 by raloxifene has been observed (Baer et al., 2007). Both compounds also inactivated rat P450 2B1 in the reconstituted system in a time-, concentration-, and NADPH-dependent manner (data not shown). Incubation of these compounds at concentrations up to 100 μM in the presence of NADPH showed no inactivation of human P450 2C9 and 2E1. When incubated with human liver microsomes, neither CPE nor TFCPE showed any effect on the BFC and TFCPE assay (data not shown). This could be because other P450s present in the microsomes may play predominant roles in the metabolism of these compounds so that relatively little metabolism occurs by 3A4 or 3A5 and therefore the amount of inactivation is negligible. We have seen

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**Fig. 7.** LC-APCI-MS/MS analysis of the metabolites of TFCPE formed by P450 3A4. The incubation conditions and analysis procedures are as described under Materials and Methods. A, mass spectrum of the metabolite eluting at 20.34 min. B and D, MS/MS analyses of the peaks with m/z values of 271 and 231 peaks, respectively. C and E, proposed fragmentation patterns for the peaks with m/z values of 271 and 231, respectively.
similar results with some other mechanism-based inactivators we have studied. Spectral analysis showed some loss of heme upon inactivation, suggesting that the inactivation of P450 3A4 could be due to both heme and apoprotein modifications (Table 2). A small peak corresponding to the heme adduct was detected by HPLC when the sample was monitored at 405 nm. The diode array spectrum of the peak was similar to that of the native heme spectrum (data not shown). However, LC-MS analysis of the TFCPE-modified P450 3A4 did not reveal a heme adduct, suggesting that the modified heme may not have been stable under the LC-MS conditions used. We have not yet been able to identify the adducts formed by CPE or TFCPE with P450 primarily because modified P450 3A4 has a tendency to aggregate, and it is difficult to analyze the inactivated protein using whole protein mass spectrometry. For this reason, we chose to determine the structures of the CPE- or TFCPE-reactive intermediates using GSH. GSH adducts of CPE and TFCPE were observed by LC-MS after incubation with P450 3A4, indicating that reactive electrophilic intermediates were generated during the metabolism and inactivation. Figures 4 and 5 show the GSH adducts of the propargyl ether compounds and their fragmentation patterns. As shown in these figures, the typical ions that are characteristic of GSH adducts were identified. However, fragmentation of the coumarin compounds themselves was not observed. As shown in Figs. 4 and 5, the masses and MS/MS ion fragmentation patterns appear to be consistent with addition of GSH, although it is not clear whether adduct formation occurs at the internal or the external carbon of the ethynyl moiety. Similar retention times were observed using a fluorescence detector that confirmed the presence of a fluorescent GSH adduct. The metabolism of TFCPE by P450 3A4 in a reconstituted system was investigated using LC-APCI-UV analysis. After a 60-min incubation with P450 3A4, one NADPH-dependent peak was observed using UV detection at 319 nm with a retention time of 20.34 min. The retention time of the parent compound TFCPE was 22.78 min. The metabolite peak was observed only in the samples that were incubated with P450 3A4 in presence of TFCPE. No similar peak or any other metabolite peak was observed under these conditions. The proposed structures of the observed metabolite and the parent are depicted in Figs. 6 and 7.

Using homology modeling, Lewis et al. (2006) have shown that metabolism of coumarin by several P450s can be rationalized in terms of likely interactions between coumarin and the active sites of the enzymes. They have shown that a combination of hydrogen bonding and π-π stacking with key amino acids within the heme environment may be responsible for positioning the coumarin substrate for metabolism. In comparing the residues that are in alignment between the SRS regions based on the P450 2C5 model, it has been suggested that phenylalanine at position 205 (in SRS2) may be associated with the coumarin aromatic ring. Consequently, π-π stacking may occur between coumarin and the phenylalanine 205 residue. In addition, they suggested that at least one hydrogen bond forming amino acid should interact with the coumarin molecule to facilitate the proper orientation in the active site and determine the site of metabolism. Although relatively closely related with respect to sequence, P450s 3A4 and 3A5 differ in 78 of a total of 503 amino acids. Of these variations 17 lie within the putative SRS domains and include the one at position 205. Thus, this active site difference could affect the binding of some substrates by these two enzymes and thus account for the different catalytic outcomes that were observed when P450s 3A4 and 3A5 metabolize the same propargyl compounds.

In conclusion, the synthesis of CPE and TFCPE, which combines the structure of a known substrate of the 3A4 P450s with a functional group that can be metabolized by these enzymes to a reactive inter-

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