

Inhibition of CYP2S1 mediated metabolism of anticancer prodrug AQ4N by liarozole

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ABSTRACT

CYP2S1 is an orphan Cytochrome P450 whose expression is elevated in numerous epithelial derived cancers; however, little is known about CYP2S1 inhibitors. Our results indicate that liarozole acts as an effective inhibitor of CYP2S1, and decreases the turnover of the prodrug AQ4N to its active metabolites AQ4M and AQ4. Our studies also found that the active drug AQ4 is found only in the perinuclear location, whereas the prodrug AQ4N is located in the cytoplasm. In addition, we showed that AQ4N is non-genotoxic/cytotoxic whereas the bioreduced metabolite AQ4 is both genotoxic and cytotoxic. Finally, we observed that CYP2S1 metabolizes AQ4N under anoxic conditions and that increasing the CYP2S1 concentration leads to more AQ4N turnover. The important enzymatic role of CYP2S1 with regard to the metabolism of the anticancer prodrug AQ4N into AQ4 was confirmed.

INTRODUCTION

Cytochrome P450 (CYPs) constitute a large superfamily of enzymes that are haem-thiolate proteins with ubiquitous expression and distribution in diverse life forms including bacteria, fungi, plants and animals (Nelson *et al.*, 1999). CYPs play an important role in carcinogen and anti-cancer drug biotransformation (Tang *et al.*, 2015). Within cells, CYPs are localized to the endoplasmic reticulum (ER) and mitochondria (Szczesna-Skorupa *et al.*, 2008). In total, 58 human cytochrome P450s (CYPs) have been identified to date. Most CYPs have been well characterized, and biological substrates have been identified for nearly 75% of all known CYPs. However there remains 25% of CYPs called "orphan P450s" which little physiological function has clearly been identified. CYP2S1 is a gene coding for the Cytochrome P450 2S1 protein, which has been shown to metabolize AQ4N, an anticancer prodrug, under anoxic conditions (Xiao *et al.*, 2011, Nishida *et al.*, 2010). The epoxygenase activity of CYP2S1 is inhibited *in vitro* by miconazole and liarozole, an inhibitor of CYP26 and CYP2S1 (Fromel *et al.*, 2013). Several ongoing studies have

identified endogenous and exogenous substrates of CYP2S1. However, very few studies have examined the role of inhibitors on CYP2S1 mediated metabolism of substrates like AQ4N. We therefore examined the role of liarozole, a known inhibitor of CYP enzymes in CYP2S1 mediated metabolism of AQ4N. Banoxantrone (AQ4N) is an anticancer prodrug that requires both a hypoxic environment and metabolic activation to generate the ultimate toxicant, a topoisomerase II inhibitor, AQ4. Heterologous CYP2S1 expression studies have revealed that AQ4N is a substrate for CYP2S1 (Nishida *et al.*, 2010), and that CYP2S1 metabolizes non-cytotoxic AQ4N to AQ4M and cytotoxic AQ4. While there have been numerous studies conducted to identify the substrates of CYP2S1, there is very little information on its inhibitors. Liarozole is an imidazole-containing compound that inhibits the cytochrome P-450-dependent metabolism of ATRA in hamster liver microsomes (Van Wauwe *et al.*, 1992). Previous studies have shown that liarozole inhibits human epidermal retinoic acid 4-hydroxylase activities (Kang *et al.*, 1996). Liarozole acts as a blocker of retinoic acid metabolism (retinoic acid metabolism blocking agent, RAMBA) and has been shown to inhibit CYP26A1 (Nelson *et al.*, 2013). It is a known Cytochrome P450 inhibitor, which inhibits several cytochrome P450 enzymes including aromatase (CYP19), a key enzyme in the biosynthesis of estrogens and retinoic acid 4-hydroxylase (CYP26S), responsible for ATRA metabolism (PavezLorie *et al.*, 2009).

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Liarozole also displays antitumor activity against androgen-dependent and independent rat prostate carcinomas (Smets *et al.*, 1995). A hypoxic environment is exhibited by solid tumors due to poor vascularization resulting from the rapid growth of tumors. Tirapazamine (Brown *et al.*, 1999), mitomycin (Tomasz 1995), NLCQ-1 (Papadopoulou and Bloomer 2003), KS119 (Seow *et al.*, 2005) and AQ4N (Patterson *et al.*, 2002, Lalani *et al.*, 2007) are a few anticancer agents that are selectively activated in an hypoxic environment (Nishida *et al.*, 2010). AQ4 itself is not suitable as an anticancer drug because it is highly toxic to normal cells (Nishida *et al.*, 2010) whereas AQ4N has negligible topoisomerase inhibitory activity (Patterson *et al.*, 1994, Smith *et al.*, 1997). When AQ4N reaches hypoxic tumor cells the drug is selectively converted into its active cytotoxic form AQ4. The active form of the drug AQ4 has been shown to be a potent topoisomerase II inhibitor and DNA intercalator (Patterson *et al.*, 2000). This work focuses on the role of the liarozole inhibitor on CYP2S1 and its subsequent effect on metabolism of AQ4N.

MATERIALS AND METHODS

Cell Lines and Reagents

A549 and BEAS2B were obtained from ATCC. A549s were grown in DMEM (Invitrogen, Cat. No. 31053-036) with 10% FBS media (Invitrogen, Cat. No. 16000-044) and BEAS2B were grown in LHC9 media (Invitrogen, Cat. No. 12680-013). The cell lines were kept in 37° C incubator with 5% CO₂ conditions. The media was changed every other day until the cells were 80 – 90% confluent, and then the cells were split using Trypsin 0.25% EDTA (Invitrogen, Cat. No. 25200-056) for A549s and TrpLE for Beas2Bs (Invitrogen, Cat. No. 12604-021). Phosphate buffer Saline (PBS) was used to wash the cells (Invitrogen, Cat. No. 10010-023).

Cell Genotoxicity Assays

Genotoxicity (DNA damage) in BEAS2B was evaluated using Trevigen Comet Assay™kit (Trevigen Inc., Gaithersburg, MD, Catalog No. # 4250-050-K) according to the manufacturers protocol. BEAS2B was plated in six well plates at a density of ~1.2 X 10⁶ cells/ml for treatment with AQ4N, AQ4 or Mitoxantrone. Three different concentrations of drug 1, 10 and 100 µM, were used. BEAS2B adherent cells were trypsinized and cells were resuspended in ice cold PBS to a concentration of 1 X 10⁵ cells/ml. The cells were combined with low melting agarose (at 42°C) at a ratio of 1:10 (v/v) and immediately pipetted (50 µl) onto CometSlide™. The slides were incubated in the dark at 4 °C for 10 min and them immersed in 4°C Lysis Solution for 60 min. The slides were then transferred to a beaker with Alkaline Unwinding Solution for 20 min at room temperature or 1 hr at 4°C, in the dark for denaturation. Approximately 850 ml of prechilled Alkaline Electrophoresis Solution was added to Comet Assay® ES unit. Then, the slides were subjected to electrophoresis at 1 V/cm, 300 mA for 40 min in the dark. After electrophoresis, the slides were removed and gently drained of excess

electrophoresis solution, and washed by immersing twice in neutralization buffer for 5 min each and in 70% ethanol for 5 min followed by air-drying at room temperature for 10-15 min. 100 µl of diluted SYBR® Gold was placed onto each circle of dried agarose and stained for 30 min at room temperature in the dark. The slides were allowed to dry completely at 37°C and analyzed using View slides by epifluorescence microscopy using Nikon e600 microscope using fluorescein filter with excitation/emission at 496 nm/522 nm. The relative DNA damage (Genotoxicity) was analyzed using olive tail moment, a product of tail length and the fraction of total DNA in the tail. Prism statistical software (Graphpad Prism 5.0) was used to graph the results obtained.

Cell Cytotoxicity Assays

The cell cytotoxicity assays were performed using AlamarBlue® cell viability reagent (Life Technologies, Cat. No.: DAL1100) according to the manufacturer's protocol. BEAS-2B cells were plated at 2,000 to 4,000 cells per well, in triplicates, of at least four per treatment or genotype, in a 96 well plate. The cell numbers for plating were calculated using the hemocytometer after trypsinizing BEAS2B cells in a T-75 flask grown to 90% confluence. The cells were allowed to grow overnight at 5% CO₂ at 37°C.

The following day media with three different concentrations (1, 10 or 100 µM) of the drug (AQ4N, AQ4 or Mitoxantrone) was added to three different plates each representing one of the drug treatments. Drug preparations were done in DMSO as described elsewhere in experimental procedures. For negative controls, DMSO and untreated media were used. For the positive control, fully reduced alamarBlue® prepared by autoclaving media with 10% alamarBlue® was used. Finally, media without cells was used as a blank to establish the background. The 96 well plates were now placed in a hypoxic chamber for 24 hrs at 1% O₂, 5% CO₂ / 37°C, and control samples were placed in a 5% CO₂ incubator at 37°C. After 24 hr, alamarBlue® was added to a final concentration of 10%, after which the cells were incubated for four more hours at previously described conditions.

The fluorescence measurements of the plates were recorded with the following parameters: excitation wavelength at 530-560nm and emission wavelength at 590 nm. Percentage reduction of alamarBlue® was calculated and these values were used to compare cells. All data points were normalized to blank media wells (alamarBlue® background fluorescence levels). The experiments were conducted at least three times under identical conditions.

Fluorescence measurements were analyzed using Prism statistical software (Graphpad Prism 5.0), analyzing linear cytotoxicity levels at different concentrations of the drug.

Isolation of CYP2S1 Microsomal Fractions

CYP2S1 overexpressing microsomes were isolated according to methods as described by Fromel *et al.*, (2013), with some modifications to the protocol. BEAS2B stable cell lines

overexpressing CYP2S1 (~5 fold) were grown until approximately 90% confluent in 75-ml flasks and harvested by trypsinization. Following centrifugation at 5000 rpm for 10 min, the cell pellet was collected. Microsomal fractions were prepared by repeat centrifugation. The cell pellet was homogenized in 3 volumes of KCl (50mM, pH 7.4) containing EDTA (1mM) and MgCl₂ (3mM) and centrifuged at 13000 rpm for 10 min at room temperature. Following centrifugation at 41000g for 30 min at 4°C, microsomes were resuspended in KCl containing 0.25M sucrose. Microsomal preparations (0.25mg) were then pre-incubated with AQ4N/AQ4 for 5 min.

The reaction was initiated by the addition of an NADPH regenerating system at a final concentration of 1.2mM (Stock 100mM). Reactions were carried out in a shaking water bath for 30 min and terminated by the addition of methanol (1 ml). Samples were vortexed for 2 min and the upper organic phase removed into a clean HPLC vial following centrifugation at 3500rpm for 15 min at 4°C.

Drug Incubations

400 µg of microsomal protein was preincubated for 5 min at 37°C under anaerobic conditions in 0.1M sodium/potassium phosphate buffer (pH 7.6) containing 5 mM NADPH. Metabolic reactions were initiated with the addition of AQ4N to a final concentration of 100 µM. The total reaction volume was 250 µl. After 60 min, reactions were terminated by the addition of 50% v/v acetonitrile (250µl) and deproteinized by centrifugation (12000g) for 5 min. Control reactions were performed in the absence of microsomes or NADPH, and under aerobic conditions.

AQ4N Assays

Chromatographic analysis of AQ4N using High Performance Liquid Chromatography (HPLC) was done as described by Loadman *et al.*, (2001). A Hichrom RPB (25-cm × 4.6-mm i.d.) column (Hichrom Ltd., Reading, UK) was used for the separation. A Waters 996 Photodiode Array Detector ($\lambda_1 = 240$ nm, $\lambda_2 = 612$ nm) with Millenium Software (Waters Ltd., Watford, UK) was used for spectral analysis of the peaks of interest. The flow rate was set at 1.2 ml/min with Waters model 510 HPLC pump and samples introduced via Waters 717 plus Autosampler. The mobile phase used was 78% ammonium formate buffer (0.05 M)/22% acetonitrile with final pH adjusted to 3.6 with formic acid.

Liarozole Inhibition Studies

5-[(3-Chlorophenyl)-1H-imidazol-1-ylmethyl]-1H-benzimidazole hydrochloride (liarozole) (Tocris biosciences, Cat. No. 2705) was used for inhibition studies. A stock concentration of 50 mM was prepared by diluting in DMSO. 20 µM CYP2S1 protein was pre-incubated with 50 µM Liarozole for 1 hr at room temperature.

Nitrogen gas was bubbled through the sample. The samples were incubated for 5 min at 37 °C under anaerobic conditions in 0.1M sodium/potassium phosphate buffer (pH 7.6)

containing 5 mM NADPH and P450 reductase (Life Technologies, Cat. No. P2309). Metabolic reactions were initiated with the addition of AQ4N to a final concentration of 100 µM. The total reaction volume was 250 µl. After 60 min, reactions were terminated by the addition of 50% v/v acetonitrile (250µl) and deproteinized by centrifugation (12000g) for 5 min. Control reactions were performed in the absence of liarozole. The control and test samples were run on HPLC as described earlier. The AQ4N, AQ4M and AQ4 peak areas obtained from HPLC were graphed using MS-Excel and Prism statistical software (Graphpad Prism 5.0) to compare the relative turnover of AQ4N in the presence and absence of liarozole.

RESULTS and DISCUSSION

AQ4, the active drug, is found in the perinuclear location whereas the prodrug AQ4N is located in the cytoplasm.

To understand the bioreduction of AQ4N metabolism, we performed fluorescence imaging studies of BEAS-2B and A549 cell lines. AQ4N undergoes two-electron reduction to get metabolized into the cytotoxic AQ4 (Fig.1A), which acts as a topoisomerase II inhibitor. Liarozole (Fig. 1B), a known CYP inhibitor, was used to inhibit CYP2S1 mediated metabolism of AQ4N to AQ4.

In order to access the location of the prodrug AQ4N and the active form of the drug AQ4, fluorescence images of BEAS2-B and A549 (Fig. 2) treated with AQ4N and AQ4 were taken using epifluorescence microscopy. Our results indicate that the prodrug AQ4N (Fig.2: A to F) is located in the cytoplasm in both BEAS2B and A549 cell lines and did not penetrate into the nucleus, even after 60 minutes of incubation.

The active drug AQ4; however, is found in the perinuclear location (Fig.2: G to L). Our results are consistent with the function of the active drug AQ4 that acts by intercalating DNA and inhibiting topoisomerase II.

The prodrug AQ4N is non-genotoxic/cytotoxic whereas the bioreduced metabolite AQ4 (active form) is both genotoxic and cytotoxic

The differences in genotoxicity and cytotoxicity of AQ4N and AQ4 have been confirmed using alkaline comet assay to detect DNA damage and alamar Blue assays for the cytotoxicity studies. BEAS2Bs treated with the control AQ4N, an inactive prodrug, did neither show any genotoxicity nor cytotoxicity whereas the active form of the drug AQ4 was both genotoxic as well as cytotoxic to the cells. There was also a dose dependent increase in the genotoxicity with the increasing levels of AQ4; however, AQ4N did not show any toxicity even with the increasing dosage (Fig. 3A). The results obtained from the genotoxicity studies are in agreement with the results obtained from cytotoxic studies. Our cytotoxicity studies of AQ4N and AQ4 indicate that the active form of the drug AQ4 is cytotoxic and the toxicity is dose-dependent, whereas the prodrug AQ4N is non-cytotoxic even with increasing concentrations of the drug (Fig. 3B).

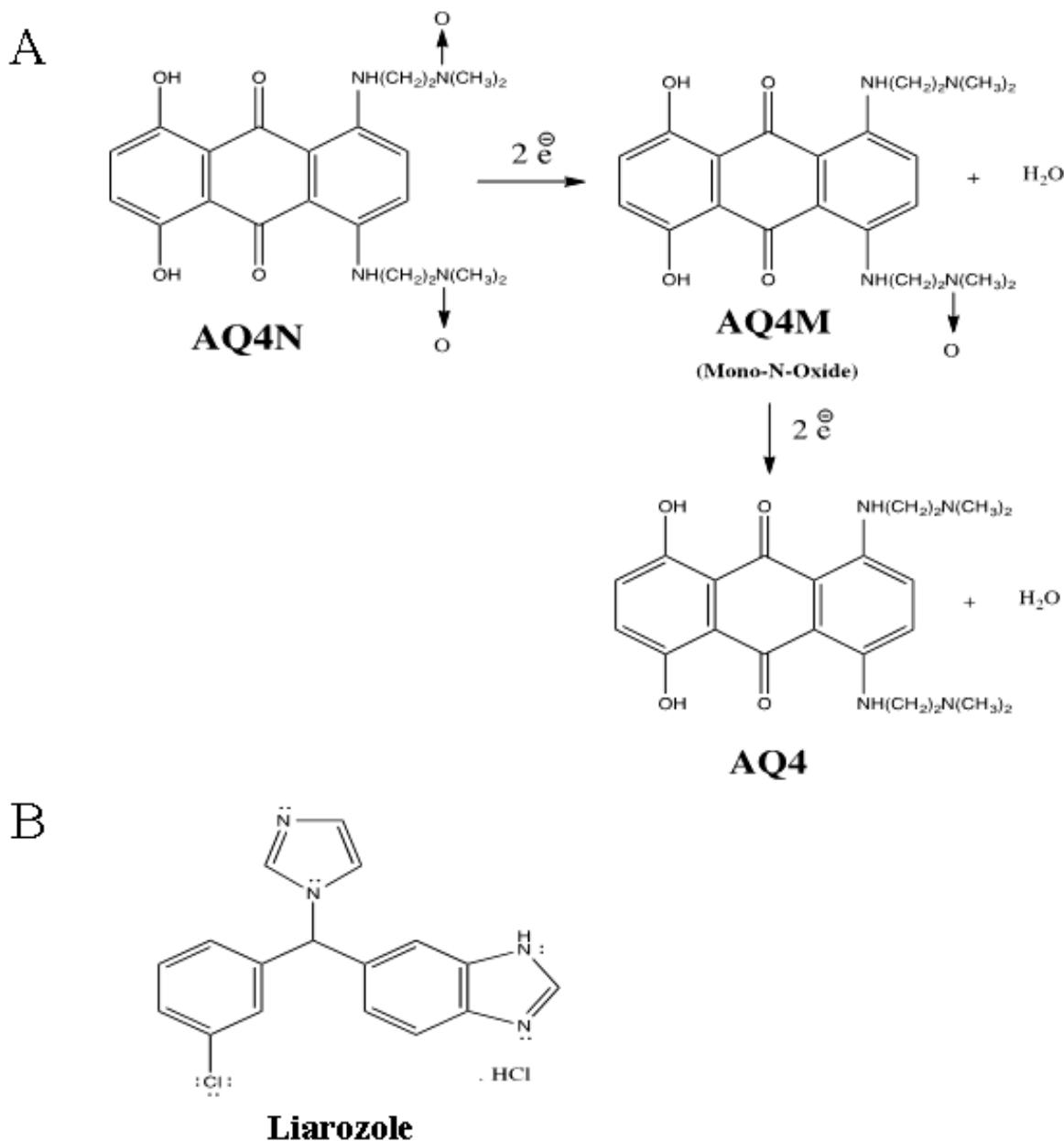


Fig. 1: AQ4N, AQ4M, AQ4 and liarozole(A). Bioreduction of AQ4N into its metabolites AQ4M and AQ4. AQ4N undergoes two-electron reduction to get metabolized into the cytotoxic AQ4. (B). Structure of liarozole.

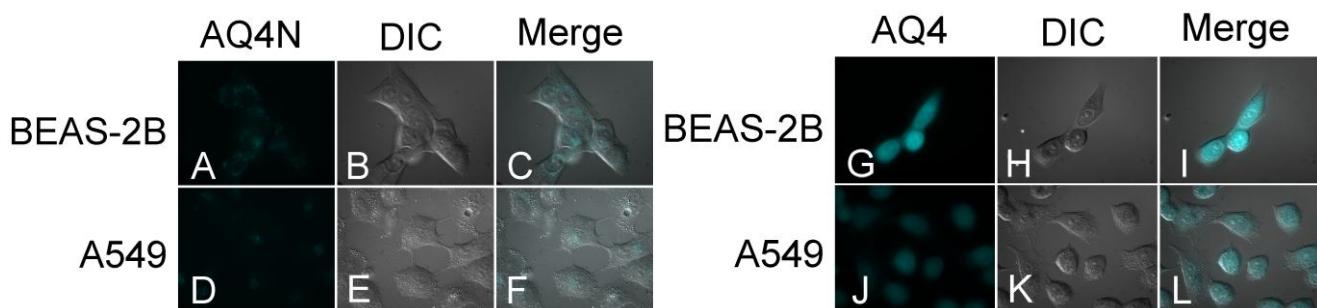


Fig. 2: Fluorescence images of BEAS-2B and A549 treated with AQ4N and AQ4. Cells were grown in glass bottom 35-mm dishes and treated with 50 µM of AQ4N or AQ4, incubated for 60 min. and were scanned with appropriate excitation (wavelength, 633 nm) and emission (650 nm LP) filters using Zeiss epifluorescence microscope. Differential Interference Contrast (DIC) shows unstained transparent samples. It can be seen from Fig. A-F that the prodrug AQ4N is located in the cytoplasm in both BEAS2B and A549 cell lines and did not penetrate into the nucleus, even after 60 minutes of incubation. The active drug AQ4; however, is located in perinuclear location (Fig. G-L).

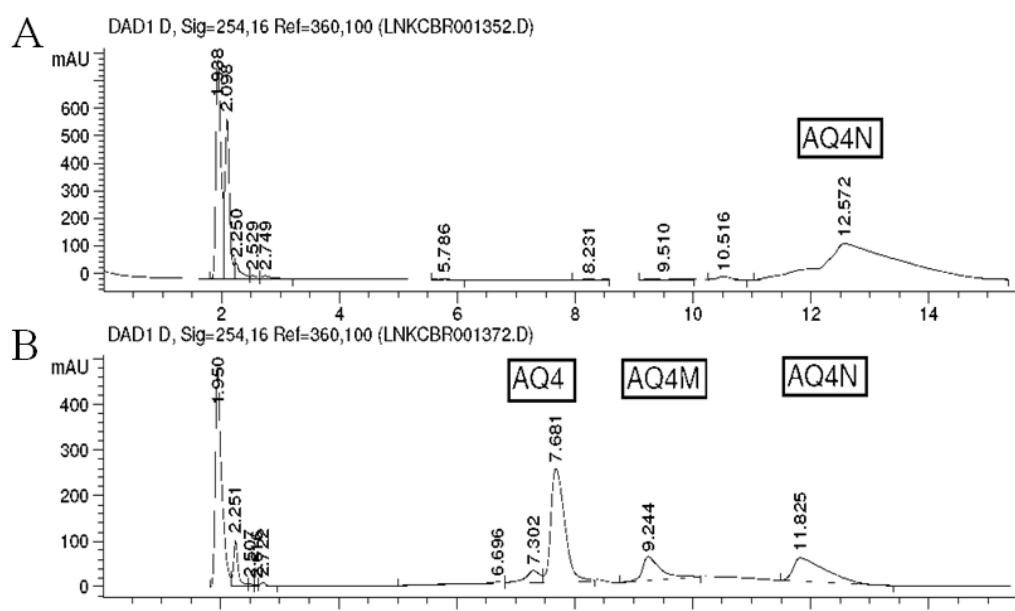
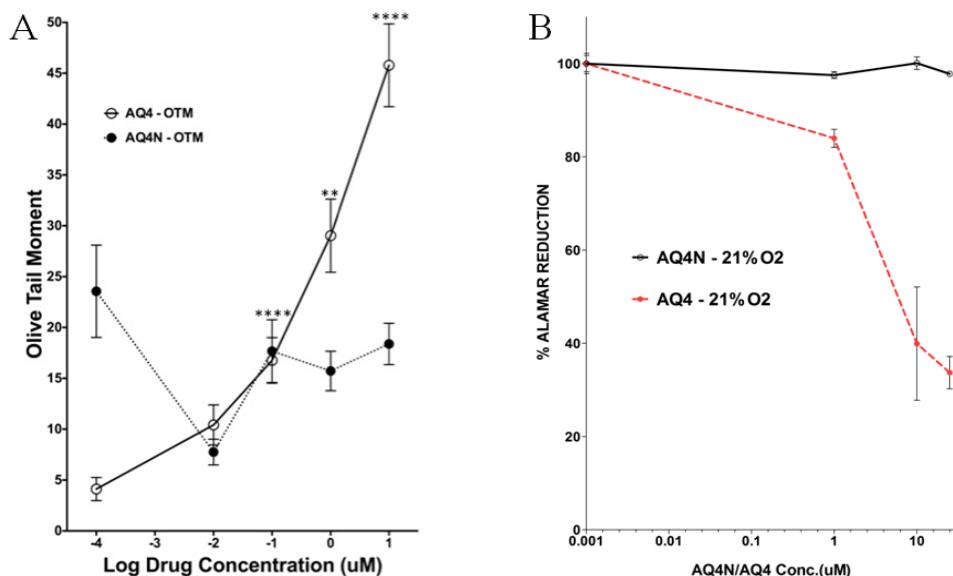


Fig. 4: HPLC analysis of bioreduction of AQ4N by CYP2S1 under anoxic conditions. (A). Representative HPLC chromatogram of control reaction with AQ4N treated with PBS instead of CYP2S1 showing AQ4N not metabolized into AQ4M or AQ4. (B) Representative HPLC chromatogram of AQ4N metabolism using CYP2S1, into AQ4M and AQ4 under hypoxic conditions. X-axis represents retention time in min, and the Y-axis is absorbance in mAU.

CYP2S1 metabolizes AQ4N under anoxic conditions and increasing CYP2S1 concentration leads to more AQ4N turnover.

Anticancer prodrugs can only be activated by bioreduction into the cytotoxic compound under hypoxic environment. Hypoxia enables the reductive activation of anticancer prodrugs that cannot be similarly activated in normoxic tissues due to competing oxidative pathways (Nishida *et al.*, 2008). Our results from HPLC analysis to determine whether CYP2S1 is able to turnover AQ4N to AQ4M and AQ4 indicate that CYP2S1

metabolizes AQ4N under anoxic conditions to AQ4M and AQ4. When compared with the control samples, where PBS was used instead of the CYP2S1, there was no conversion of AQ4N to AQ4M or AQ4—there was only a single peak (retention time: 12.572 min) corresponding to the AQ4N (Fig. 4A). However, in the reactions that involved CYP2S1, three distinct peaks of AQ4N, AQ4M and AQ4 were observed indicating the turnover of prodrug AQ4N to the active cytotoxic product AQ4 through CYP2S1 under anoxic conditions (Fig. 4B). Next, we tested whether increasing concentration of CYP2S1 protein will lead to dose

dependent increase in the rate of metabolism of AQ4N. Our results indicate that the increasing concentration of CYP2S1 leads to increased turnover of AQ4N (Fig. 5).

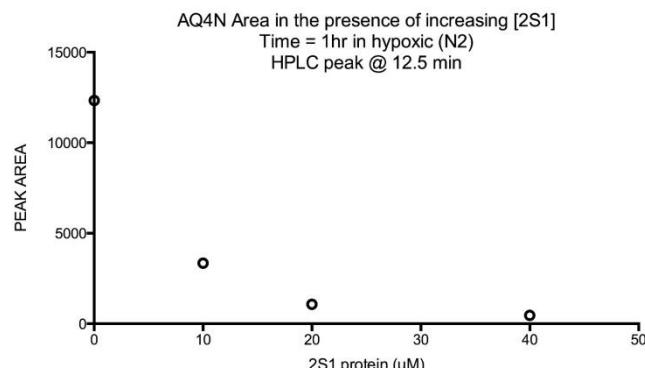


Fig. 5: HPLC analysis of bioreduction of AQ4N with increasing concentration of CYP2S1 under anoxic conditions. Increasing concentration of CYP2S1 lead to the increased turnover of AQ4N into AQ4M and AQ4 and relative reduction in the peak area of AQ4N.

Liarozole inhibits CYP2S1 and reduces the turnover of AQ4N to AQ4M and AQ4.

To understand more about the physiological function of CYP2S1 inhibition by liarozole, we treated microsomes from BEAS2B with liarozole followed by incubation with P450 reductase and NADPH under anoxic conditions. In our studies with CYP2S1 overexpressing microsomes, liarozole effectively inhibited the CYP2S1-mediated metabolism of AQ4N (Fig. 6). As can be seen in Fig. 6, In the presence of liarozole, CYP2S1 mediated metabolism of AQ4N to AQ4 is reduced, as is evident from the decreased turnover of AQ4N to its metabolites.

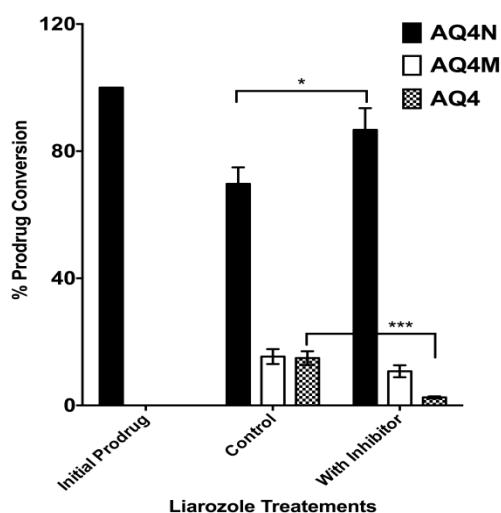


Fig. 6: CYP2S1 is inhibited by liarozole. Microsomes from BEAS2B were treated with liarozole followed by incubation with P450 reductase and NADPH under anoxic conditions. 10μM AQ4N was added and the reactions were incubated at room temperature for an hour followed by deproteinization. The supernatants were analyzed by HPLC for the conversion to AQ4M and AQ4. Liarozole was not added to the control samples. Liarozole inhibits CYP2S1 and prevents the binding of AQ4N to the enzyme, thus resulting in decreased turnover of AQ4N into AQ4M and AQ4 when compared to untreated control samples. Statistical significance was determined using a Student's *t* test. Data are represented as mean ± S.D., ***, *p* ≤ 0.001, *, *p* ≤ 0.05.

CONCLUSIONS

Cytochrome P450 metabolic studies play an important role in understanding the disposition and metabolism of various exogenous and endogenous metabolites. Identification of inhibitors for orphan CYPs is a useful tool in deducing their significance in the turnover of various substrates and also for effective enzyme studies. Our results indicate that liarozole is an effective inhibitor of CYP2S1, which has been tested against the substrate AQ4N. This study reveals part of the enzymatic role of CYP2S1 and ultimately its physiological role and significance in cancers.

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