Anti-tumor Activity of Methoxymorpholinyl

Doxorubicin: Potentiation by Cytochrome P450 3A

Metabolism¹

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Running Title: P450-activation of MMDX

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Abbreviations - CHO, Chinese hamster ovary; CYP, cytochrome P450; Dex,

dexamethasone; GDEPT, gene-directed enzyme prodrug therapy; MDR, multi-drug

resistance; MMDX, methoxymorpholinyl doxorubicin; MRP, MDR-associated protein;

P-gp, P-glycoprotein.

Summary – Methoxymorpholinyl doxorubicin (MMDX) is a novel liver cytochrome P450 (CYP)-activated anti-cancer prodrug whose toxicity toward cultured tumor cells can be potentiated up to 100-fold by incubation with liver microsomes and NADPH. In the present study, MMDX was activated by a panel of human liver microsomes with potentiation ratios directly correlated to the CYP3A-dependent testosterone 6β -hydroxylase activity of each liver sample. Microsome-activated MMDX exhibited nanomolar IC₅₀ values in growth inhibition assays of human tumor cell lines representing multiple tissues of origin: lung (A549 cells), brain (U251 cells), colon (LS180 cells) and breast (MCF-7 cells). Analysis of individual cDNAexpressed CYP3A enzymes revealed that MMDX was activated by rat CYP3A1 and human CYP3A4 more efficiently than by rat CYP3A2, and that human CYPs 3A5 and 3A7 displayed little or no activity. MMDX cytotoxicity was substantially increased in CHO cells following stable expression of CYP3A4 in combination with P450 reductase. CYP3A activation of MMDX abolished the parent drug's residual cross-resistance in a doxorubicin-resistant MCF-7 cell line that over-expresses P-glycoprotein. CYP3A-activated MMDX displayed a comparatively high intrinsic stability, with a $t_{1/2}$ of ~5.5 hr at 37°C. MMDX was rapidly activated by CYP3A at low (~1-5 nM) prodrug concentrations, with 100% tumor cell kill obtained after as short as a 2 hr exposure to the activated metabolite. These findings demonstrate that MMDX can be activated by CYP3A metabolism to a potent, long-lived and cell-permeable cytotoxic metabolite, and suggest that this anthracycline prodrug may be used in combination with CYP3A4 in a P450 prodrug activation-based gene therapy for cancer treatment.

Introduction

Anthracyclines, such as daunorubucin and doxorubicin, are used to treat a wide range of malignancies, but myelosuppression, cardiotoxicity and the emergence of multidrug resistance limit their effectiveness in the clinic. Novel anthracyclines that show improved activity, lower toxicity and the ability to circumvent drug resistance are therefore sought. These efforts have led to the discovery of morpholinyl anthracyclines such as MMDX (3'-deamino-3'-[2(S)-methoxy-4-morpholinyl]doxorubicin; PNU 152243), a particularly promising candidate that is presently under clinical evaluation (Suarato et al., 1999; Sun et al., 2003). MMDX contains a methoxymorpholinyl group at the 3' position of the sugar moiety and is highly lipophilic. Unlike doxorubicin, MMDX is not cardiotoxic at optimal anti-tumor doses (Danesi et al., 1993) and shows substantially reduced cross-resistance in tumor cell lines highly resistant to doxorubicin (Bakker et al., 1997; Kuhl et al., 1993; Ripamonti et al., 1992; van der Graaf et al., 1995). Mechanistically, MMDX induces DNA strand breaks primarily through topoisomerase I cleavage, whereas doxorubicin induces DNA lesions through topoisomerase II cleavage (Duran et al., 1996).

MMDX exhibits up to ~8-fold greater toxicity than doxorubicin toward drug-sensitive tumor cells *in vitro* (Kuhl et al., 1993) but 80-fold greater potency *in vivo* (Ripamonti et al., 1992). This difference in activity reflects the fact that MMDX is a prodrug that is activated *in vivo* in a liver cytochrome P450 (CYP)-catalyzed reaction. Thus, the cytotoxicity of MMDX in cell culture can be markedly enhanced by pre-incubation with

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liver microsomes and NADPH in a metabolic process that is inhibited by the cytochrome P450 3A (CYP3A)-selective inhibitors triacetyloleandomycin and ketoconazole (Baldwin et al., 2003; Lewis et al., 1992; Quintieri et al., 2000). P450-activated MMDX induces DNA-DNA interstrand cross-links, in contrast to MMDX, which primarily induces protein-associated DNA single-strand breaks (Lau et al., 1994; Lau et al., 1989). Other studies suggest that activated MMDX differs from the parent drug in terms of its spectrum of anti-tumor activity and pattern of resistance (Geroni et al., 1994; Kuhl et al., 1993; van der Graaf et al., 1995).

In the present report, we investigate the role of specific CYP3A enzymes in the metabolic activation of MMDX and we evaluate the cytotoxicity of CYP-activated MMDX to a range of human tumor cells. We characterize the intrinsic stability of activated MMDX and its cytotoxicity toward tumor cells resistant to doxorubicin and partially cross-resistant to MMDX. Finally, we evaluate the impact of tumor cell expression of CYP3A on MMDX toxicity. Overall, our findings indicate that MMDX has several properties that make it an ideal candidate for combination with CYP3A4 in a P450 prodrug-activation gene therapy strategy (Chen and Waxman, 2002; Jounaidi, 2002).

Materials and Methods

<u>Chemicals</u> - NADPH (tetrasodium salt, cat# N6505) was purchased from Sigma-Aldrich (St. Louis, MO). MMDX hydrochloride (PNU-152243) was a gift from Pharmacia & Upjohn (Milan, Italy). MMDX was prepared as a 10 mM (6.8 mg/ml) stock solution in

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ddH₂O and then diluted to 10 μ M and stored in aliquots at -20°C, which retained activity for at least 3 months. NADPH was prepared fresh for each experiment as a 120 mM stock solution in ddH₂O and kept on ice until use. Aqueous solutions of MMDX and NADPH were filter-sterilized immediately after preparation by passage through a 0.22 μ M syringe filter (Pall Corp. East Hills, NY; cat# 4454). Except for the initial weighingout of chemicals, all solutions used for cell culture studies were prepared under sterile conditions. Crystal violet solution for cell staining was prepared by dissolving 1.25 g crystal violet (Sigma cat# C3886) in 50 ml of 37% formaldehyde and 450 ml methanol and then stored at room temperature.

Liver Microsomes - Liver microsomes prepared from dexamethasone-induced, phenobarbital-induced and uninduced male Sprague-Dawley rats were purchased from In Vitro Technologies, Inc. (Baltimore, MD) and used in the experiments shown in Fig. 1 and Fig. 5. Dexamethasone-induced male Fischer 344 rat liver microsomes prepared in this laboratory by differential centrifugation were used in all other experiments. Microsomes prepared from individual human liver donors and microsomes from baculovirus-infected insect cells expressing individual P450 cDNAs (SupersomesTM) were purchased from BD-Gentest Corp. (Woburn, MA). Donor information were provided by the supplier for 3 of the liver donors, HLM samples 42, 89 and 112, which were from female donors ages 48, 71 and 2 yr-old, and had CYP3A4 protein contents of 306, 177 and 331 pmol P450/mg protein, respectively. Donor 112 had been treated with phenobarbital and several other medications during hospitalization. The SupersomesTM

combination with human P450 reductase (P450R) and cytochrome b_5 (BD-Gentest cat# P202); rat CYP3A1 with rat P450R and cytochrome b_5 (cat# P501); rat CYP3A2 with rat P450R and cytochrome b_5 (cat# P502); human CYP3A5 with human P450R (cat# P235); human CYP3A7 with human P450R and cytochrome b_5 (cat# P237); and control Supersomes, which are devoid of P450, P450R and cytochrome b_5 (cat# P201). All microsomes were stored in aliquots at -80°C.

Cell Culture - The rat gliosarcoma cell line 9L is a brain tumor-derived adherent cell line, and was cultured in Dulbecco's Modified Eagle Medium (DMEM) (Gibco-BRL Life Technologies, Rockville, MD) supplemented with 10% fetal bovine serum (FBS), 100 U/ml penicillin and 100 µg/ml streptomycin (Gibco). A549, a human lung carcinoma cell line, and U251, a human glioma cell line, were cultured in Roswell Park Memorial Institute (RPMI) medium 1640 containing 5% FBS and penicillin/streptomycin antibiotic mixture (1:100, Gibco). LS180, a human colon adenocarcinoma cell line, was grown as a monolayer in 10% Minimal Essential Medium (MEM) containing penicillin/streptomycin. The human breast carcinoma cell line MCF7, and its doxorubicin-resistant subline, MCF7-ADR, were cultured in RPMI 1640 medium containing 5% FBS plus penicillin/streptomycin. CHO cell lines expressing human P450R alone, or P450R in combination with CYP3A4 (~8 pmol P450/mg, and CYP3A4-dependent testosterone 6βhydroxylase activity of 572 pmol/min/mg) were kindly provided by Dr. T. Friedberg (Univ. of Dudee, Dundee, Scotland) (Ding et al., 1997). CHO-hP450R cells were cultured in MEM plus 10% FBS, 1:100 diluted penicillin/streptomycin mixture and hypoxanthine/thymidine supplement (Gibco). CHO-3A4-hP450R cells were cultured in

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 α -MEM medium containing 10% dialyzed fetal bovine serum (Gibco) and penicillin/streptomycin (Ding et al., 1997).

Microsomes Prodrug Co-culture Assay - Tumor cell lines were seeded in 96 well micro-plates at a density of 1000 cells/well (9L, A549, U251, CHO-hP450R and CHO-3A4-hP450R cells), or 3000 cells/well (MCF7, MCF7-ADR and LS180 cells), and allowed to attach for 24 hr. MMDX diluted in fresh culture medium was then added to each well at concentrations specified in each experiment. Each well also contained 0.3 mM NADPH (final concentration) and microsomal protein (rat or human liver microsomes; 2 µg/well) or insect cell microsomes containing cDNA-expressed CYPs (0.2 pmol P450/well), as indicated. The final volume of each well was 200 µl. Cells were routinely cultured for 4 d after the addition of MMDX, NADPH and microsomes. The culture medium was then removed and the wells gently washed with PBS. Crystal violet staining solution (100 μ l; see above) was added to each well and the plates were shaken gently for 20 min on an orbital shaker. Excess stain was then removed and the plates were washed by immersion in water and air-dried. The cell-adhering crystal violet stain was suspended in 100 μ l of 70% ethanol over 20 min with gentle shaking. A₅₉₅ values were determined using an SLT SPECTRA Shell Reader (SLT Lab Instruments, Austria). Background A₅₉₅ values, based on an average of triplicate cell-free wells on each plate, were subtracted from the A_{595} value of each experimental sample and determined in triplicate. Relative cell survival, expressed as a percentage of the corresponding drugfree control, was calculated as follows:

Cell Survival (%) =
$$\frac{(A_{595 \text{ drug treated}} - A_{595 \text{ background}})}{(A_{595 \text{ drug-free control}} - A_{595 \text{ background}})} \times 100$$

 IC_{50} values (drug concentration that decreases cell survival to 50% of control levels) were determined from a semi-logarithmic graph of the data points.

To evaluate the time course for MMDX activation, 9L cells were seeded in 96 well tissue culture microtiter-plates at a density of 1000 cells/well and allowed to attach for 24 hr. MMDX at 0.5, 1 or 5 nM and dexamethasone-induced rat liver microsomes (2 µg/well) plus NADPH were then added to each well. Cells were cultured for different time periods, after which the culture medium containing drugs, microsomes and NADPH was removed and replaced with fresh DMEM plus 10% FBS. Cells were then cultured in the absence of the microsomal activation system for the remainder of the experiment, i.e., for a total of 4 d after beginning MMDX treatment, at which time the cells were stained with crystal violet. Relative cell survival as a function of time of incubation with MMDX was calculated as described above, with the drug-free control taken as the zero time point.

<u>Half-life of CPA3A-activated MMDX</u> - MMDX at 0.5 and 1 nM was pre-incubated in DMEM (without FBS) and dexamethasone-induced rat liver microsomes (10 μ g/ml) plus NADPH for 30 min at 37°C. The reaction was terminated by spinning down microsomal proteins at 12,000 g for 5 min. The supernatant was filtered through a 0.22 μ m syringe filter and the filtrates were divided into aliquots. One aliquot was immediately removed and a portion diluted 2-fold into fresh culture medium. The diluted and undiluted filtrates

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were then added to 9L cells as a zero time point. The remaining aliquots were incubated at 37°C for times up to 12 hr, at which point the samples and 2-fold dilutions prepared from each sample were added to 9L cells, which were cultured for 4 days. Cell survival was determined by crystal violet staining as described above. Data were graphed as log of percent cell kill (i.e., 1- cell survival percentage) vs. time of incubation at 37°C, and $t_{1/2}$ values were calculated from the log-linear decay portion of each curve.

<u>Statistical Analysis</u> - IC₅₀ values and 95% confidence interval were calculated from semi-logarithmic dose-response curves using a variable slope sigmoid equation (Prism 4, GraphPad Software Inc., San Diego, CA). Values differed significantly at P < 0.05 when their 95% confidence intervals did not overlap. The relationship between IC₅₀ values and testosterone 6β-hydroxylase activities was analyzed by Pearson correlation.

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Results

Cytotoxicity of MMDX activated by rat and human liver microsomes - The

cytotoxicity of MMDX towards 9L gliosarcoma cells can be increased by incubation with dexamethasone-induced rat liver microsomes in an NADPH-dependent manner (Baldwin et al., 2003). In agreement with this finding, the cytotoxicity of MMDX toward 9L cells in cells cultured was increased in a 4-day growth inhibition assay carried out in the presence of NADPH and either uninduced, phenobarbital-induced or dexamethasoneinduced rat liver microsomes and NADPH. The potentiation ratio (ratio of IC_{50} of MMDX in the absence of microsomes to IC_{50} of MMDX incubated with microsomes) was 87 for dexamethasone-induced liver microsomes, 22 for phenobarbital-induced microsomes and 8 for uninduced liver microsomes. Strong increases in MMDX cytotoxicity were also obtained using a panel of microsomes prepared from individual human liver donors (Fig. 1B). These microsomes varied in the cytochrome P450 protein content and activity profiles, reflecting individual differences due to factors such as age, sex and drug exposure of the liver donors. Potentiation ratios varied from ~10 (HLM-066) to ~60 (HLM-112), and the IC₅₀ values ranged from 0.66 to 3.8 nM (Table 1). Several of the human liver microsome (HLM) samples were similar in activity to the highly active dexamethasone-induced rat liver microsomes (c.f., Fig. 1A). Moreover, the capacity of each HLM to activate MMDX correlated directly with its CYP3A activity, assayed by testosterone 6β -hydroxylation, a CYP3A-selective liver microsomal activity $(r = 0.945 \text{ for IC}_{50} \text{ (MMDX)} \text{ vs. testosterone } 6\beta\text{-hydroxylase activity for } n = 5 \text{ HLM}$ samples; p<0.05).

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Activation of MMDX by cDNA-expressed CYP3A enzymes - We next evaluated a panel of cDNA-expressed CYP3A enzymes (SupersomesTM) to determine the potential of these enzymes to activate MMDX (Fig. 2). CYP3A4, a human P450 enzyme, exhibited the highest activity ($IC_{50} = 1.4 \text{ nM}$) when tested in a 4-day 9L growth inhibition assay. The rat enzymes CYP3A1 and CYP3A2 also exhibited significant activity ($IC_{50} = 2.5 \text{ nM}$ and 5.4 nM, respectively). Two other human P450 3A enzymes, CYP3A5 and CYP3A7, exhibited little or no MMDX activation when compared to the CYP-deficient insect microsome control. Thus, CYP3A4, the major CYP3A enzyme in human liver, is the most active catalyst of MMDX activation.

Experiments were carried out using CHO cells that stably express CYP3A4 and P450R (Ding et al., 1997) to determine if MMDX cytotoxicity can also be increased when CYP3A4 is expressed intracellularly. A potentiation ratio of 22 was obtained for MMDX incubated with CHO-3A4-hP450R cells compared to the CYP3A4-deficient control cell line CHO-hP450R (Fig. 3). This MMDX potentiation ratio was 2.8-fold higher than when CHO-hP450R cells were incubated with MMDX in the presence of dexamethasone-induced microsomes (potentiation ratio = 8; Fig. 3 and Table 2). Based on the measured IC₅₀ of 0.8 nM MMDX for the CHO-3A4-hP450R cells (Table 2), a cellular CYP3A4 content of 8 pmol P450/mg protein (Ding et al., 1997) and an estimated content of ~1-2 μ g/well (0.008-0.016 pmol CYP3A4 protein/well), in comparison to an IC₅₀ of 1.4 nM for MMDX activation by 0.2 pmol of extracellular CYP3A4 (Fig. 2), we

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estimate that the cells are at least 12-25-fold more sensitive to MMDX when the prodrug is activated intracellularly.

Sensitivity of human tumor cell lines to activated MMDX - We next investigated whether the strong potentiation of MMDX cytotoxicity seen with dexamethasoneinduced microsomes and rat 9L cells is also observed with human tumor cells. MMDX activated by dexamethasone-induced microsomes was highly toxic toward the human tumor cell lines A549 (lung), U251 (brain), MCF-7 (breast) and LS180 (colon) (Fig. 4), with potentiation ratios ranging from 11 to 43 (Table 2). A 6-fold difference in sensitivity to unactivated MMDX was seen when the most sensitive tumor cell line (U251) was compared to the least sensitive cell line (LS180). Cell line-based differences in the IC₅₀ values of activated MMDX were more narrow, ranging only ~ 2.3 fold, and did not correlate with the sensitivity of each cell line to MMDX itself (Table 2). This finding is consistent with earlier findings that MMDX and activated MMDX kill tumor cells by distinct mechanisms (Lau et al., 1994; Lau et al., 1989).

Evaluation of cross-resistance of MCF7-ADR cells to MMDX and to CYP3A-

activated MMDX - MCF7-ADR is an MCF-7 cell line derivative that overexpresses the drug efflux pump P-glycoprotein and is 500-fold more resistant to doxorubicin (Table 3) (Chen and Waxman, 1995; Fairchild et al., 1987). MCF7-ADR cells are cross-resistant to many other, structurally distinct drugs, which are also substrates for P-glycoprotein (P-gp) and are efficiently pumped out of the cells. Previous reports indicate that MMDX is active against cultured tumor cell lines that overexpress P-glycoprotein and are highly

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resistant to doxorubicin, however, some cross-resistance to MMDX is apparent (Bakker et al., 1997; Kuhl et al., 1993; van der Graaf et al., 1995). In contrast, *in vivo* studies indicate that MMDX retains substantial activity against doxorubicin-resistant tumor cells (Ripamonti et al., 1992). To clarify this issue, we evaluated the cytotoxicity of MMDX and of activated MMDX toward MCF-7 cells and toward the P-glycoproteinoverexpressing subline MCF7-ADR. As shown in Fig. 5 and Table 3, MCF7-ADR cells were ~5.2-fold more resistant to MMDX when compared to MCF-7 cells. The cytotoxicity of MMDX toward MCF7-ADR cells was substantially improved in the presence of dexamethas one-induced liver microsomes (potentiation ratio = 15), such that activated MMDX was equally active against MCF-7 and MCF7-ADR cells (resistance factor = 1). In the case of doxorubicin, however, incubation with microsomes led to a ~7-fold *decrease* in activity toward MCF-7 cells (potentiation ratio = 0.14), indicating that doxorubicin is inactivated by microsomal metabolism. The cytotoxicity of doxorubicin toward MCF7-ADR cells was also decreased by microsomal metabolism, albeit to a smaller extent. We conclude that CYP3A activation of MMDX, but not doxorubicin, fully reverses the resistance phenotype of MCF7-ADR cells.

Intrinsic activity and stability of CYP3A-activated MMDX - In the growth inhibition assays described above, tumor cells were cultured with liver microsomes and MMDX for a total of 96 hr, at which time drug-induced tumor cell kill was evaluated. Experiments were therefore carried out to determine whether strong cytotoxic responses could be achieved after shorter times of incubation and microsomal metabolism. Fig. 6 shows that a 2 hr incubation of MMDX (5 nM) with dexamethasone-induced rat liver microsomes

generated sufficient active metabolite to kill ≥95% of the 9L tumor cells by the end of the 4-day assay period. Thus, the MMDX activation reaction is rapid and does not require full 96 hr period to generate activated, cytotoxic metabolites. Even at a 5-fold lower MMDX concentration, ~85% tumor cell kill was achieved within a 6 hr incubation period. Interestingly, under these latter conditions, the formation of cytotoxic microsomal metabolites was not detectable during the initial 1 hr period. This apparent lag period suggests that the concentration of activated MMDX may need to reach a minimum threshold level, perhaps reflecting saturation of cellular binding sites not linked to the cytotoxic response.

Next, we investigated the intrinsic stability of activated MMDX. Initial studies showed that a protein-free filtrate, prepared after preincubation of MMDX with dexamethasone-induced microsomes and NADPH for 30 min, killed >90% of 9L tumor cells in a 4 d growth inhibition assay, but that a 12 hr pre-incubated MMDX sample was totally inactive. This loss of activity after 12 hr incubation with microsomes could reflect the intrinsic (chemical) instability of MMDX, or alternatively, may be the result of further microsomal metabolism of activated MMDX to yield inactive, secondary metabolites. To distinguish these possibilities, we evaluated the intrinsic stability of activated MMDX incubated at 37°C in the absence of liver microsomes. Protein-free filtrates were prepared from samples of MMDX preincubated with dexamethasone-induced liver microsomes and NADPH for 30 min. The protein-free filtrates, containing activated MMDX, were then incubated at 37°C for times ranging from 2 hr to 12 hr, at which point aliquots were added to 9L cells, which were then cultured for 4 d in a growth inhibition

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assay. Fig. 7 shows that the growth inhibitory activity of CYP3A-activated MMDX was retained for an extended period of time. The length of this 'plateau' period of stability was directly related to the extent of MMDX activation during the 30 min pre-incubation period, as indicated by the cytotoxic activity at time zero (y-intercept value). At the end of each plateau period, the cytotoxic activity of each activated MMDX sample declined in a log-linear fashion, giving a series of parallel lines whose slopes indicate first-order kinetics for decomposition of activated MMDX with a $t_{1/2}$ (apparent) of 5.5 \pm 1.6 hr.

Discussion

CYP3A-selective inhibitors, such as troleandomycin, markedly decrease the therapeutic efficacy of MMDX in mice bearing M5076 liver metastases (Quintieri et al., 2000) and substantially inhibit liver microsome-catalyzed activation of MMDX to tumor cell cytotoxic species *in vitro* (Baldwin et al., 2003; Lewis et al., 1992). These findings were extended by the present study, which revealed a strong correlation between CYP3A-catalyzed testosterone 6β -hydroxylase activity and the potentiation of MMDX cytotoxicity in a panel of human liver microsomes, and demonstrated that the human enzyme CYP3A4 and its rodent counterparts CYP3A1 and CYP3A2 are active catalysts of the prodrug activation reaction. These findings are consistent with the observation that the metabolism of MMDX by human liver microsomes can be competitively inhibited by the CYP3A4 substrates paclitaxel, tamoxifen and cyclosporin (Beulz-Riche et al., 2002).

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CYP3A metabolism resulted in a striking potentiation of MMDX cytotoxicity at nanomolar prodrug concentrations, both when the CYP enzyme was present extracellularly and when it was expressed endogenously in CHO cells stably transfected with CYP3A4. Comparison of the increase in MMDX cytotoxicity conferred by intracellular vs. extracellular CYP3A4 expression suggested than intracellular MMDX activation is at least 12-25-fold more potent than the extracellular activation system, a phenomenon that has been seen with other anti-cancer prodrugs (Lawrence et al., 1998). Extratumoral (i.e., hepatic) activation of MMDX is nevertheless likely to be the major site of pro-drug activation in vivo, given the very low level of CYP3A4 expression and activity found in human tumors (Iscan et al., 2001; Kivisto et al., 1996; Knupfer et al., 1999; Martinez et al., 2002; Murray et al., 1999). These findings point to the potential advantages of using gene therapy vectors to increase the content of CYP3A4 within tumor cells and thereby augment the toxicity, and the tumor cell selectively, of MMDX. A P450 prodrug activation-based gene therapy designed to achieve this goal has recently been developed with CYP2B6 for the anti-cancer prodrug cyclophosphamide and has shown promise in preclinical and in initial phase I/II clinical trials (Chen and Waxman, 2002; Jounaidi, 2002).

The precise nature of the active, cytotoxic metabolite of MMDX is not presently known. P450-activated MMDX is highly cytotoxic and induces DNA-DNA crosslinks (Lau et al., 1994), in contrast to MMDX itself, which, like doxorubicin, induces DNA strand breaks (Ross et al., 1979). Six to 8 distinct metabolites are formed when MMDX is incubated with liver microsomes, depending on the species and on the MMDX concentration

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(Beulz-Riche et al., 2001). Among these, the only characterized metabolite detected both in human patients and in animals is MMDX-ol (13-dihydro-3'-deamino-3'-[2(S)methoxy-4-morpholinyl]doxorubicin, or PNU-155051). However, this metabolite is formed by an aldo-keto reductase without the involvement of cytochrome P450 and is less cytotoxic than MMDX (Breda et al., 2000). Several other characterized and uncharacterized MMDX metabolites are formed by human liver microsomes in a reaction that can be inhibited by CYP3A4 substrates (Beulz-Riche et al., 2001; Beulz-Riche et al., 2002). However, the potential role of these metabolites in MMDX cytotoxicity remains to be determined.

Analysis of four human tumor cell lines revealed a 6-fold range in sensitivity to MMDX, with IC₅₀ values ranging from 12 nM MMDX for U251 brain cancer cells to 74 nM for LS180 colon cancer cells. Notably, both the IC₅₀ values and the cell line to cell line differences in drug sensitivity were decreased for CYP3A-activated MMDX. The more uniform response of the tumor cells to P450-activated MMDX may reflect differences in the mechanism of cell killing, with activated MMDX able to induce a direct, and rapid, alkylation of tumor cell DNA (Lau et al., 1992). P450-activated MMDX may not display improved tumor selectivity compared to MMDX, however, given its comparable cytotoxicity against tumor cells and hematopoietic progenitors (Ghielmini et al., 1998; Kuhl et al., 1993). Consistent with this observation, the CYP3A inhibitor troleandomycin markedly inhibits MMDX induced myelosuppression *in vivo* (Quintieri et al., 2000).

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e.g., using a tumor-targetted gene therapy vector delivering CYP3A4, may therefore be useful in increasing the therapeutic index of MMDX.

In addition to the differences in mechanism of action noted above, activated MMDX differs from MMDX in its pattern of drug resistance. The doxorubicin-resistant MCF-7 subline MCF7-ADR displays 500-fold resistance to doxorubicin associated with overexpression of the drug efflux pump P-gp (Fairchild et al., 1987). Other factors, including a decrease in topoisomerase activity may also contribute to the drug-resistance of this cell line (Sinha et al., 1988). MCF7-ADR cells displayed a much lower degree of resistance (~5-fold) to MMDX than to doxorubicin (Fig. 5), which may reflect the higher lipophilicity of MMDX, a lower affinity for P-gp, or perhaps the formation of DNA strand breaks through topoisomerase I, rather than topoisomerase II (Duran et al., 1996). The cyclosporin derivative and P-gp inhibitor SDZ-PSC833 partially reverses resistance to MMDX in four models of multi-drug resistance involving P-gp overexpression (Michieli et al., 1996). Moreover, intracellular levels of MMDX are reduced 2-fold in a doxorubicin-resistant, P-gp-over-expressing cell line compared to those in the drugsensitive parent cell line (Bakker et al., 1997). These results suggest that P-gp may prevent MMDX from reaching its nuclear target, particularly when the drug concentration is low. The elevated glutathione peroxidase activity in MCF7-ADR cells (Chen and Waxman, 1995) could also contribute to the residual MMDX cross-resistance, as suggested by the correlation between increased glutathione peroxidase activity and decreased MMDX activity in a panel of human melanoma cell lines (Alvino et al., 1997). Alterations in topoisomerase I activity in MCF7-ADR cells could also contribute to the

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cross-resistance to MMDX. Interestingly, although cross-resistance to MMDX is also seen in a doxorubicin-resistant cell line that over-expresses MDR-associated protein (MRP) (van der Graaf et al., 1995), MMDX is not a substrate of the drug transporter MRP (Bakker et al., 1997), suggesting that other mechanisms are responsible for the cross-resistance seen in P-gp-negative MDR cell lines. However, in contrast to the crossresistance of MCF7-ADR cells to MMDX, no cross-resistance to P450-activated MMDX was seen in the present study. This finding is consistent with the report that a cyanomorpholino derivative of doxorubicin that has alkylating properties similar to the active metabolites of MMDX is not cross-resistant with doxorubicin in MDR cells (Scudder et al., 1988). Together, these findings suggest that P450-activated MMDX is not a substrate for the drug-efflux pump P-gp and may be effective against tumors that exhibit a multi-drug resistance phenotype.

Phase I and phase II clinical trials and pharmacokinetic studies of MMDX showed a terminal half-life of 40 hr after a rapid distribution phase (Bakker et al., 1998; Vasey et al., 1995). The area under the plasma concentration-time curve calculated up to 24 hr after dosing at 1.25 mg/m² MMDX was ~20 ng/h/ml, and the C_{max} ranged from 2 to 6 ng/ml (3 to 9 nM) (Bakker et al., 1998; Vasey et al., 1995). These latter concentrations are greater than or similar to the IC₅₀ values presently reported for MMDX toxicity to human tumor cells following incubation with liver microsomes and NADPH (Table 2). Investigation of the time course of MMDX activation *in vitro* revealed that CYP3A-activated MMDX metabolites accumulate within 2 hr at 5 nM prodrug to levels that are sufficient to induce maximum cytotoxicity, suggesting that strong MMDX anti-tumor

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activity may be readily achieved *in vivo*. Although the plasma MMDX concentration during the terminal elimination phase is less than 1 ng/ml (1.5 nM) (Bakker et al., 1998; Vasey et al., 1995), active metabolites generated by liver CYP3A enzymes during this time period could still be highly toxic, given the very slow elimination of MMDX, which at 40 hr is 7-times longer than the 6 hr presently required for efficient activation of 1 nM MMDX by liver microsomes in vitro (Fig. 6). Furthermore, we observed that activated MMDX generated during a 30 min liver microsomal activation reaction maintained its initial, maximal level of cytotoxicity for as long as 3-6 hr when incubated in the presence of cells, after which activity declined with first-order kinetics and a half-life of 5.5 hr. Thus, CYP3A-activated MMDX can be formed rapidly and decays relatively slowly, suggesting that sufficient levels of active drug may accumulate in vivo to sustain antitumor activity. However, the highly toxic nature of liver CYP3A-activated MMDX is likely related to the dose-limiting hematotoxicity (Fokkema et al., 2000; Sessa et al., 1999). This problem may be minimized in the case of tumors containing high CYP3A4 expression and activity, whether expressed endogenously or achieved using gene therapy vectors that augment intratumoral CYP3A4 expression.

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Figure Legends

Fig. 1 - Activation of MMDX by rat and human liver microsomes. Panel A – Growth inhibition assay of 9L cells treated with MMDX alone ('no microsomes'), or with MMDX in the presence of drug-induced rat liver microsomes and NADPH. Liver microsomes were prepared from dexamethasone-induced rats (Dex), phenobarbitalinduced rats (PB), or uninduced rats. Each cell culture well contained 200 μ l cell culture medium, 300 μ M NADPH, 2 μ g rat liver microsomal protein and MMDX at the indicated concentrations. Data were obtained in a 4 d growth inhibition assay, as described under Materials and Methods. Each data point is the mean \pm S.D. values based on n=3 independent samples. Panel B – 9L growth inhibition assay carried out as in panel A, except that human liver microsomes from individual donors (0.5 μ g microsomal protein/well) were used in place of rat liver microsomes. IC₅₀ values based on panel B are shown in Table 1. Data shown are mean values \pm S.D. (n=3).

<u>Fig. 2</u> – <u>Activation of MMDX by cDNA-expressed CYP3A enzymes</u> – 9L growth inhibition assays were carried out as a function of MMDX concentration, as described in Fig. 1, using insect cell microsomes containing individual cDNA-expressed human or rat CYP3A proteins, as indicated, with co-expressed P450R and cytochrome b5 ('Supersomes'; 0.2 pmol P450 protein/well) in place of liver microsomes. Insect cell microsomes not containing P450 served as a negative control. Data shown are mean values \pm S.D. (n=3).

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<u>Fig. 3</u> – <u>Impact of intracellular CYP3A4 expression on MMDX cytotoxicity</u> - MMDX toxicity to CHO cells expressing human P450R alone (CHO-hP450R cells), or P450R in combination with CYP3A4 (CHO-3A4-hP450R cells) was determined in a 4 d growth inhibition assay. The cytotoxicity of MMDX to CHO-hP450R cells incubated in the presence of dexamethasone-induced rat liver microsomes (0.5 µg protein/well) and NADPH is shown for comparison. Data shown are mean values \pm S.D. (n=3). IC₅₀ values and potentiation ratios based on these data are shown in Table 2.

<u>Fig. 4</u> – <u>Sensitivity of human tumor cell lines from multiple tissues of origin to</u>

<u>microsome-activated MMDX</u> – Data shown are based on a 4 d growth inhibition assay similar to Fig. 1A in the absence of NADPH and dexamethasone-induced rat liver microsomes (2 μ g protein/well) (open symbols) or in the presence of NADPH and liver microsomes (closed symbols). Data shown are mean values ± S.D. (n=3). IC₅₀ values and potentiation ratios based on these data are shown in Table 2.

<u>Fig. 5</u> - <u>Sensitivity of MCF7 and MCF7-ADR cells to MMDX (panel A) or doxorubicin</u> (panel B) in the absence (open symbols) or presence (closed symbols) of microsomal <u>activation system</u> – Four day growth inhibition assay was carried out in the presence or absence of dexamethasone-induced rat liver microsomes, as described in Fig. 1A. Data shown are mean values \pm S.D. (n=3). IC₅₀ values based on these data are shown in Table 3.

<u>Fig. 6</u> – <u>MMDX growth inhibition as a function of time of incubation with</u> <u>dexamethasone-induced liver microsomes</u> – 9L cells were incubated with MMDX (1 or 5

nM), 2 μ g dexamethasone-induced rat liver microsomes/well and NADPH (0.3 mM) for the indicated periods of time, at which point the cells were washed and placed in drugand microsome-free culture medium for a 4 day culture period as described under Materials and Methods. Data shown are the mean values \pm S.D. (n=3).

Fig. 7 – Intrinsic stability of CYP3A-activated MMDX – MMDX (0.5 or 1 nM) was

preincubated for 30 min with dexamethasone-induced rat liver microsomes (10 µg protein/ml culture medium) and NADPH (0.3 mM). The samples were then filtered and the protein-free filtrate (diluted 2-fold into fresh culture medium where indicated; "1/2") was added to 9L cells to assay for cytotoxic activity in a 4 d growth inhibition assay. Cell killing was calculated as described under Materials and Methods. A half-life value of 5.5 ± 1.6 hr was calculated for activated MMDX from the portions of each curve showing a time-dependent, log-linear decline in cell killing activity (mean \pm S.D., n = 3 curves). No decline in cytotoxicity was observed for the 1 nM MMDX sample over the 12 hr incubation period, indicating that the amount of activated MMDX initially present in this sample was at least in 4-fold excess over that required to kill all of the 9L cells, i.e., no decline in activity was apparent after incubation at 37°C for 12 hr (~2 half-lives) prior to addition to the cells.

<u>Table 1</u> – IC₅₀ values and potentiation ratios for MMDX incubated with 9L cells and a panel of human liver microsomes.

| HLM # (Liver donor number) | IC ₅₀ (nM)* | Potentiation ratio [†] | Testosterone 6β-hydroxylase activity [‡] |
|----------------------------------|---------------------------|------------------------------------|---|
| no microsomes | 38.75 | 1 | _ |
| 66 | 3.84 | 10 | 5800 |
| 70 | 1.86 | 21 | 8960 |
| 89 | 1.46 | 27 | 11730 |
| 42 | 0.74 | 52 | 13150 |
| 112 | 0.66 | 59 | 17520 |

* Data are based on triplicate samples analyzed as shown in Fig. 1B.

 \dagger IC_{50} of MMDX incubated with control microsomes/IC_{50} of MMDX incubated with HLM.

‡ Testosterone 6β-hydroxylase data, provided by BD-Gentest, exhibited a significant Pearson correlation with the potentiation ratio values shown in column 3 (r = 0.9452; p<0.05).

<u>Table 2</u> - IC_{50} values and potentiation ratios of MMDX or incubated MMDX with dexamethasone-induced rat liver microsomes plus NADPH on human tumor cell lines and CHO cell lines either express human P450 reductase or CYP3A4 supplemented with human P450 reductase.

| | IC | | | |
|----------------|------|---------------------------------------|---------------------------------|--|
| Cell Line | MMDX | MMDX + dexamethasone microsomes | Potentiation ratio ^b | |
| CHO-hP450R | 17.6 | 2.2 | 8 | |
| CHO-3A4-hP450R | 0.8 | Not determined | _ | |
| U251 | 11.8 | 1.1 | 11 | |
| A549 | 24.2 | 0.99 | 24 | |
| MCF7 | 36.6 | 2.3 | 16 | |
| LS180 | 73.6 | 1.7 | 43 | |

^a Each value is based on triplicate samples analyzed as shown in Fig. 3 and Fig. 4.

^b IC₅₀ of MMDX / IC₅₀ of (MMDX + dexame has one-induced liver microsomes)

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<u>Table 3</u> - IC₅₀ values, potentiation ratios and resistance factors for MMDX and

doxorubicin toward MCF-7 and MCF7-ADR cells incubated in the presence or

absence of dexamethasone-induced rat liver microsomes and NADPH.

| | IC ₅₀ valu | | |
|---------------------------------|-----------------------|-----------------------------------|-----|
| | (Potentiati | Resistance Factor ^c | |
| | MCF7 | MCF7-ADR | |
| Doxorubicin | 12.6 | 6,600 | 524 |
| Doxorubicin + Dex microsomes | 86.6 (0.4) | 10,300 (0.64) | 119 |
| MMDX | 31 | 161 | 5.2 |
| MMDX + Dex microsomes | 10.2 (3.0) | 10.7 (15.0) | 1.0 |

^a Data based on triplicate samples as shown in Fig. 5 and data not shown.

^b IC_{50} of drug/ IC_{50} of drug + dexame thas one-induced liver microsomes.

 $^{\rm c}$ IC_{50} for MCF7-ADR / IC_{50} for MCF-7

Library: MMDX_Hong.enl

Style: Mol Pharm













