

Direct Binding of Chloroquine to the Multidrug Resistance Protein (MRP)

POSSIBLE ROLE FOR MRP IN CHLOROQUINE DRUG TRANSPORT AND RESISTANCE IN TUMOR CELLS

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ABSTRACT. Multidrug resistance protein (MRP) transports a range of compounds that include glutathione S-conjugates, amphiphilic anionic drugs, and natural-product toxins. However, the mechanism of MRP drug binding and transport is presently unclear. We recently demonstrated the direct binding of a quinoline-based photoactive drug, N-{4-[1-hydroxy-2-(dibutylamino)ethyl] quinolin-8-yl}-4-azidosalicylamide (IAAQ), to MRP at a biologically relevant site [Vezmar et al., Biochem Biophys Res Commun 241: 104-111, 1997]. In the present report, we demonstrated that the lysosomotropic or antimalarial drug chloroquine is a substrate for MRP. Specifically, our results showed that chloroquine, similar to leukotriene C₄ (LTC₄) and 3-(3-(2-(7-chloro-2quinolinyl)ethenyl-phenyl)((3-(dimethyl amino-3-oxo propyl)thio)methyl)thio) propanoic acid (MK 571), inhibits the photoaffinity labeling of MRP by IAAQ. Furthermore, cell growth assays showed MRP-expressing multidrug-resistant cells (H69/AR and HL60/AR) to be more resistant to chloroquine than their parental cells (i.e., IC50 of 121 µM versus 28 µM chloroquine for H69/AR and H69, respectively). Moreover, MK 571, an LTD₄ receptor antagonist, reversed the resistance of H69/AR cells to chloroquine. Drug transport studies using [14C]chloroquine demonstrated that MRP-expressing cells accumulate less drug than the parental drug-sensitive cells. The reduced accumulation of [14C]chloroquine in resistant cells was ATP dependent and was due to enhanced drug efflux. Taken together, the results of this study show that MRP modulates the transport of chloroquine by direct binding. BIOCHEM PHARMACOL **56**;6:733–742, 1998. © 1998 Elsevier Science Inc.

KEY WORDS. multidrug resistance; chloroquine; multidrug resistance protein (MRP); drug transport; photoaffinity labeling

Tumor cell lines selected *in vitro* with sublethal concentrations of natural-product drugs respond by overexpressing P-gp1† (for reviews, see Refs. 1 and 2). The overexpression of P-gp1 in response to drug selection occurs frequently, and tumor cells display a pleiotropic resistance towards a diverse group of drugs. Studies with purified P-gp1 reconstituted into defined lipids have confirmed the role of P-gp1 as an ATP-dependent drug efflux pump [3, 4]. The normal function of P-gp1 remains unknown. However, P-gp1 was shown recently to function as a "flipase" of short chain lipids and to mediate the transport of normal cell metabolites and xenobiotics [5–7]. Furthermore, disruption of both P-gp1 and P-gp2 genes in mice led to an increase in drug accumulation in normal tissues [8, 9].

Cole et al. [10] described the overexpression of another membrane transporter, the MRP, in in vitro selected H69

to express MRP without P-gp1 [11]. P-gp1 and MRP are members of a large family of ABC trafficking proteins that couple ATP hydrolysis to ligand transport across a lipid membrane [12, 13]. Although gene transfer studies of P-gp1 or MRP, in previously drug-sensitive cells, were shown to confer largely similar drug resistance profiles, the mechanisms of P-gp1 and MRP-mediated MDR are thought to be different [14-17]. Thus, while P-gp1 binds directly to unmodified drugs and mediates their efflux across the cell membrane in an ATP-dependent manner [18], the mechanism of MRP-mediated MDR is not clear. Several reports have demonstrated increased efflux of unmodified drugs from MRP-overexpressing cells or cells transfected with human MRP cDNA [14, 19, 20]. In contrast, other studies have shown an ATP-dependent efflux of unmodified drugs only in the presence of GSH [21, 22]. Hence, an outstanding question concerning the role of GSH in the binding and transport of drugs by MRP remains to be resolved. In a recent report by Rappa et al. [23], MRP was shown to function as a co-transporter of GSH and natural-product toxins in embryonic stem cells. Furthermore, disruption of the MRP gene in mice led to increased sensitivity to

SCLC cells that displayed an MDR phenotype without

P-gp1. Since then, other MDR cell lines have been shown

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[†] Abbreviations: ABC, ATP binding cassette; LT, leukotriene; MDR, multidrug resistance; MRP, multidrug resistance protein; P-gp1, P-glycoprotein; and SCLC, small cell lung cancer.

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natural-product toxins and elevated GSH levels in MRP-expressing tissues [24].

Several studies have now shown MRP to mediate the transport of GSH-conjugated compounds and other normal cell metabolites that include the cysteinyl leukotriene (LTC₄), GSSG, and 17β-estradiol 17-(β-D-glucuronide) [21, 22, 25, 26]. These organic anions are thought to be potential physiological substrates for MRP and are consistent with findings that suggest that MRP is the same or a related protein to the multispecific organic anion transporter (MOAT) [27]. Moreover, direct binding between LTC₄ and MRP has been shown in intact cells and membrane vesicles from MRPoverexpressing cells [26, 28]. Furthermore, LTC₄ binding and transport by MRP were inhibitable with excess (3-(3-(2-(7-chloro-2-quinolinyl) ethenyl)phenyl)((3-dimethyl amino-3-oxo propyl)thio) methyl)thio) propanoic acid (MK 571), an antagonist of the LTD₄ receptor [25]. Interestingly, it has not been possible to demonstrate direct binding between MRP and other substrates of MRP (e.g. natural-product drugs [11]). Using a photoactive quinolinebased drug, $(N-\{4-[1-hydroxy-2-dibutylamino\})$ ethyl] quinolin-8-yl}-4-azidosalicylamide; IAAQ), we recently provided the first evidence for a direct and specific binding of unmodified drug to MRP in the H69/AR SCLC cells [29]. The inhibition of MRP photolabeling by IAAQ with MK 571, also a quinoline-based drug [30], led us to examine the possibility that other quinoline-based drugs may interact directly with MRP. Therefore, it was of interest to determine if the lysosomotropic or antimalarial drug, chloroquine, is a substrate for MRP. An earlier report by Cole et al. [31] had shown that MRP-overexpressing cells (H69/ AR) are less sensitive to chloroquine. However, it was not known from that study if MRP mediated chloroquine resistance. Moreover, chloroquine has been shown previously to interact with P-gp1 and to reverse P-gp1-mediated MDR [32]. In addition, chloroquine resistance in malaria is thought to be mediated by a P-gp1 homologue, Pfmdr1 [33–36]. However, the role of *Pfmdr1* in chloroguine resistance in malaria remains controversial [37–39].

In this report, we have used the photoactive quinoline-based drug IAAQ [29] to demonstrate a direct binding between chloroquine and MRP in intact cells and plasma membranes. Moreover, our results implicate MRP in the cross-resistance and transport of chloroquine in MRP-expressing tumor cells.

MATERIALS AND METHODS Materials

Iodine-125 (100.7 mCi/mL) was purchased from Amersham Biochemical Inc. The LTD₄ receptor antagonist MK 571 was provided by Dr. A. W. Ford–Hutchinson (Merck–Frosst Centre for Therapeutic Research [30]). LTC₄ was purchased from the Cayman Chemical Co. The SCLC cells (H69 and H69/AR) were gifts from Dr. Susan P. C. Cole (Cancer Research Laboratories). The HL60 and HL60/AR cells were a gift from Dr. M. Center at Kansas State

University. All other chemicals were of the highest commercial grade available.

Cell Culture and Plasma Membrane Preparation

Drug-sensitive (H69 and HL60) and -resistant (H69/AR and HL60/AR) cells were grown in RPMI 1640 medium containing 4 mM glutamine and 5–10% fetal bovine serum (HyClone). Resistant cells were cultured continuously in the presence of doxorubicin; however, cells used for drug transport studies were grown in drug-free medium for 10 days prior to the date of the experiment. Plasma membranes from H69 and H69/AR cells were prepared essentially as described by Lin *et al.* [40]. Membrane fractions were stored at -80° if not used immediately. Protein concentrations were determined by the method of Lowry *et al.* [41].

Cytotoxicity Assays

Cells were harvested and plated in 96-well plates at 0.5 to 1.0×10^4 cells/well. After allowing the cells to recover for 24 hr, cells were grown in increasing concentrations of chloroquine in the absence or presence of non-toxic concentrations of MK 571. Cells were grown for 4 days at 37° before the addition of the MTT dye [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide]. The plates were processed as previously described [42]. The effects of drugs on the viability of cells were expressed as the means \pm SD of three independent experiments in which triplicates were assayed.

Drug Transport

Drug-sensitive and -resistant cells (1 \times 10⁶ cells/sample) were washed three times with PBS (pH 7.2) containing 1.0 μM [14C]chloroquine (52 mCi/mmol) and 5 mM D-glucose or 2-deoxy-glucose and 100 nM sodium azide. Drug accumulation in cells was stopped at different time points (0-60)min) with the addition of 1 mL of ice-cold PBS, and then cells were washed three times with the same solution. The cell pellets were lysed in 100 µL of 1 M NaOH followed by neutralization with equal volumes of 1 M HCl, and the accumulated radiolabel was determined by fluorography using 1219 Racbeta model counters (LKB, Wallace). For drug efflux, cells were loaded with 10 µM [14C]chloroquine in the presence of 100 nM sodium azide and 10 mM 2-deoxy-glucose for 30 min at 37°. Then cells were washed with PBS containing 5 mM D-glucose, and the incubation was continued for 60 min at 37°. Samples were collected at different times, up to 60 min. Cells were washed in ice-cold PBS, and the cell pellets were lysed and processed as indicated above.

Photoaffinity Labeling

For photoaffinity labeling of cells, H69 or H69/AR cells were washed with PBS and preincubated for 20 min at 37°

in the presence of excess vinblastine, chloroquine, LTC₄, or MK 571 before the addition of 0.1 to 0.25 μ M IAAQ [29]. Cells were incubated at room temperature in the dark for 30 min and then were transferred to ice for 10 min. Following the latter incubation on ice, cells were irradiated for 10 min on ice with a UV source at 254 nm (Stratagene UV crosslinker, Stratagene) and lysed in 20 μ L of 50 mM Tris (pH 7.4) containing 1% NP-40, 5 mM MgCl₂, and protease inhibitors (3 μ g/mL of leupeptin and 2 mM phenylmethylsulfonyl fluoride). IAAQ-photolabeled proteins were isolated by a brief centrifugation and processed for SDS-PAGE. For photoaffinity labeling of plasma membranes, 15- to 30- μ g aliquots of plasma membranes prepared from H69 and H69/AR cells were photoaffinity labeled as described above.

SDS-PAGE and Western Blotting

Samples containing IAAQ-labeled proteins were mixed with 1:5 (v/v) of buffer I [10 mM Tris–HCl, pH 8.0, containing 2% SDS, 50 mM dithiothreitol (DTT), 1 mM EDTA] and an equal volume of buffer II [2x buffer I and 9 M urea]. The solubilized proteins were then resolved by SDS–PAGE using the Fairbanks gel system with some modifications [43]. Gel slabs containing the resolved proteins were fixed in 50% methanol, dried, and exposed to XAR Kodak film at -70° . For Western blotting, proteins were transferred to nitrocellulose membrane and probed with the MRP-specific monoclonal antibody QCRL-1 (0.5 μ g/mL) essentially as outlined by Towbin *et al.* [44]. Immunoreactive proteins were visualized by chemiluminescence using the ECL method (Amersham Inc.).

RESULTS

MRP has been shown to confer resistance to a broad spectrum of dissimilar drugs [11]. Our recent demonstration that the photoactive drug IAAQ and MK 571, both quinoline-based drugs (Fig. 1), bind directly to MRP led us to examine the possibility that other quinoline-based drugs also may be substrates for MRP [29]. Of several possible quinoline-based drugs, we were interested to know if the antimalarial drug chloroquine (Fig. 1) is a substrate for MRP. To determine if chloroquine interacts directly with MRP, IAAQ was used to photoaffinity label MRP in the absence and in the presence of excess chloroquine, MK 571, LTC₄, and vinblastine (Fig. 2). The results in Fig. 2A (lanes 1 and 2) show the photoaffinity labeling of MRP with IAAQ in H69/AR but not in H69 cells. Further, a molar excess of chloroquine, MK 571, LTC₄, and, to a lesser extent, vinblastine inhibited MRP photoaffinity labeling in intact H69/AR cells (Fig. 2A). Similar photoaffinity labeling results of MRP were also obtained using plasma membranes from H69 and H69/AR cells (Fig. 2B). In intact cells, LTC₄ at 75-fold molar excess completely inhibited the photoaffinity labeling of MRP with IAAQ (lane 9, Fig. 2A). Chloroquine and MK 571 were equal in

FIG. 1. Organic structures of quinoline-based drugs: MK 571, chloroquine, and IAAQ (N-{4-[1-hydroxy-2-(dibutylamino)ethyl] quinolin-8-yl}-4-azidosalicylamide).

their ability to inhibit IAAQ photolabeling of MRP (Fig. 2A). Vinblastine at 300- to 1000-fold excess of IAAQ also inhibited MRP photoaffinity labeling but to a much lesser extent than LTC₄, MK 571, or chloroquine (lanes 3 and 4 of Fig. 2, A and B). Together, these results show that chloroquine, LTC₄, and MK 571 interact directly with MRP.

To determine if MRP confers resistance to chloroguine, we measured the effect of increasing concentrations of chloroquine on the growth of H69/AR and HL60/AR MDR cells versus the parental H69 and HL60 cells (Fig. 3A). The results show H69/AR cells to be less sensitive to chloroquine than the parental H69 cells (IC_{50} of 121 μM vs 28 μM chloroquine, respectively). Similar results were also observed with another MRP-expressing cell line [HL60/ AR; Fig. 3A (b)]. However, HL60/AR cells were more sensitive to chloroquine than the H69/AR cells (IC₅₀ 12 μ M vs 121 μ M, respectively). Differences in the sensitivity of HL60/AR and H69/AR to chloroquine and other cytotoxic drugs (data not shown) are likely due to differences in MRP expression between the two cell lines (Fig. 3C). To further confirm the role of MRP in the cross-resistance of H69/AR cells to chloroquine, cells were grown in increasing concentrations of chloroquine plus non-toxic concentrations of MK 571. The results in Fig. 3B show that MK 571 (at 10 or 30 μM) increased the sensitivity of H69/AR cells to chloroquine (from IC₅₀ of 121 µM in the absence of 736 M. Vezmar and E. Georges

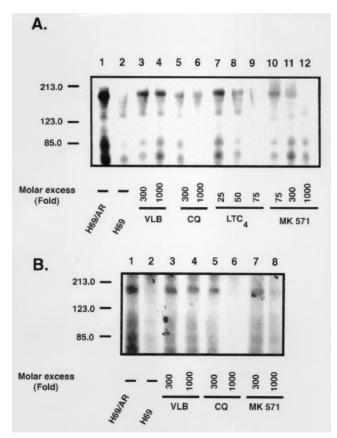


FIG. 2. Effects of chloroquine, vinblastine, MK 571, and LTC₄ on photoaffinity labeling of MRP in intact cells (A) and in plasma membranes (B). H69 or H69/AR cells were photoaffinity labeled with 0.25 μ M IAAQ in the absence (lanes 2A and 1A, respectively) or presence of 300- and 1000-fold molar excess of vinblastine (VLB), chloroquine (CQ), or 25- to 75-fold of LTC₄ or 75- to 1000-fold MK 571 (lanes 3A–12A, respectively). Similarly, plasma membranes from H69 and H69/AR cells were photoaffinity labeled with 0.25 μ M IAAQ in the absence (lanes 2B and 1B, respectively) and the presence of 300- to 1000-fold molar excess of vinblastine (VLB), chloroquine (CQ), or MK 571 (lanes 3B–8B, respectively). The numbers on the left of each panel are M_r standards in kilodaltons.

MK 571 to IC_{50} of 89 or 73 μ M in the presence of 10 or 30 μ M of MK 571, respectively). Similar concentrations of MK 571 did not affect the sensitivity of H69 cells to chloroquine [Fig. 3B (a)].

The above results show a correlation between MRP expression and reduced sensitivity to chloroquine in H69/AR and HL60/AR cells. However, it was not clear if the observed effects were due to differences in chloroquine transport or metabolism. Panels A and C of Fig. 4 show the accumulation of [14C]chloroquine in drug-resistant (H69/AR and HL60/AR) and -sensitive (H69 and HL60) cells. Both H69/AR and HL60/AR cells accumulated less [14C]chloroquine than the H69 and HL60 parental cells (Fig. 4, A and C). Moreover, the measured decrease in [14C]chloroquine accumulation in resistant cells was reversed completely when cells were preincubated for 30 min with sodium azide and 2-deoxy-glucose (Fig. 4, B and D,

respectively). Interestingly, and in view of the differences in chloroquine cross-resistance and MRP expression, H69/AR and HL60/AR showed similar levels of chloroquine accumulation (Fig. 4). However, it should be mentioned that differences in the subcellular distribution of MRP have been described previously between these two cell lines [45, 46].

To determine if the reduced chloroquine accumulation in resistant cells was due to enhanced efflux, cells were loaded with [14C]chloroquine (see Materials and Methods), and drug efflux was measured in the absence and in the presence of sodium azide and 2-deoxy-glucose. The results in Fig. 5A show a more rapid efflux of [14C]chloroquine from HL60/AR than from HL60 cells. By contrast, there was no enhanced efflux of [14C]chloroquine from H69/AR cells compared with the parental H69 cells (Fig. 5B). However, depletion of ATP levels with sodium azide and 2-deoxy-glucose inhibited the efflux of chloroquine from both sensitive (HL60 or H69) and resistant (HL60/AR or H69/AR) cells (Fig. 5, A and B). To determine if the reduced accumulation, without enhanced efflux, of [14C]chloroquine in H69/AR cells is due to an enhanced metabolism of the drug, cells were incubated with [14C]chloroquine, and the fate of the [14C]radiolabel was monitored by HPLC, as previously described [47]. The mobility of the extracted radiolabeled material from H69/AR or H69 cells was compared with the unaltered [14C]chloroquine by HPLC using a C₁₈ reverse-phase column [47]. Analysis of the HPLC results showed the elution time of the extracted radiolabel to be identical to that of unmodified [14C]chloroquine (data not shown). Thus, differences in [14C]chloroquine transport between H69/AR and HL60/AR are not due to the metabolism or organic modification of the drug.

DISCUSSION

P-gp1 and MRP are members of a large family of evolutionarily conserved ABC trafficking proteins [12, 13]. Although a certain overlap in P-gp1 and MRP substrate specificity exists, the two proteins are thought to differ in their mechanisms of drug transport [11]. For example, unlike P-gp1, MRP binds to and mediates the transport of GSHconjugated drugs [28, 48, 49]. In addition, a GSH-dependent vincristine transport was shown in membranes from cells transfected with MRP [21]. However, it was not clear if GSH was required for drug binding and transport by MRP. Our previous demonstration that a quinoline-based drug binds directly to MRP in intact cells and in membrane fractions suggests that GSH is not required for drug binding [29]. In the present study, we showed that chloroquine, another quinoline-based drug, also binds directly to MRP. This is based on the ability of chloroquine to inhibit the photoaffinity labeling of MRP by IAAQ in purified membranes. In addition, we also found that the binding of chloroquine to MRP did not require direct modification of the drug (data not shown). Similar inhibition of MRP

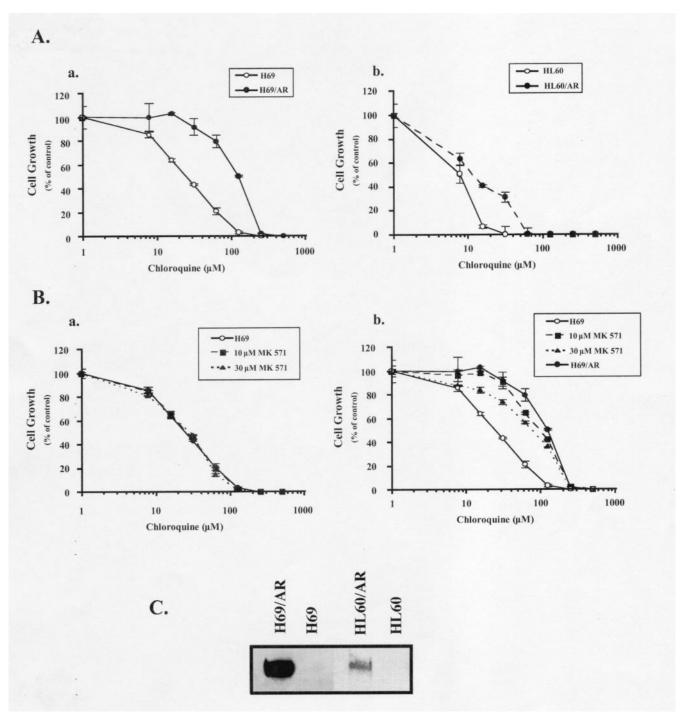


FIG. 3. Chloroquine cross-resistance and reversal with MK 571 in tumor cell lines. Drug-sensitive (H69 or HL60) and -resistant (H69/AR or HL60/AR) cells were plated at 0.5 to 1.0×10^4 and incubated in increasing concentrations of chloroquine (1–500 μ M) (Fig. 3A, a and b, respectively). The effects of 10 and 30 μ M MK 571 on the chloroquine cross-resistance of H69 and H69/AR are shown in Fig. 3B, a and b, respectively. The IC₅₀ values were determined from the graph. Each value is the mean \pm SD of three experiments in which triplicates were assayed. Figure 3C shows a Western blot of total cell lysates from MRP-overexpressing cells (H69/AR and HL60/AR) and their parental cells (H69 and HL60) probed with QCRL-1 monoclonal antibody.

photolabeling with IAAQ in intact cells and plasma membranes was also observed in the presence of LTC_4 , MK 571, and, to a lesser extent, vinblastine (Fig. 2). Thus, although the above results are consistent with a direct binding between MRP and quinoline-based drugs (IAAQ, MK 571, and chloroquine), a catalytic role of GSH in MRP drug

binding cannot be ruled out. Nevertheless, based on these results, we speculate about the ability of MRP to recognize compounds that contain a quinoline moiety. It is interesting in this respect that earlier studies [50] with P-gp1 have suggested that most drugs that bind to P-gp1 contain a planar benzene ring and a cationic charge (i.e. a quinoline

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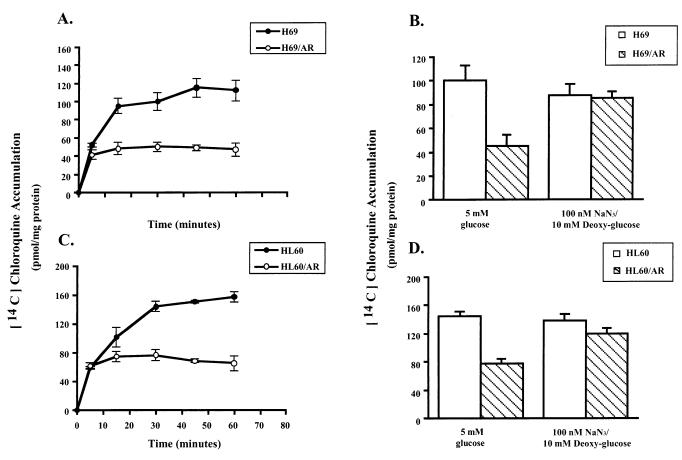


FIG. 4. [14 C]Chloroquine accumulation in MRP-expressing cells and their parental cells. Drug sensitive (H69 or HL60) and -resistant (H69/AR and HL60/AR) cells were incubated in the presence of 1 μ M [14 C]chloroquine at 37°. Samples were removed, and the associated radioactivity was determined at 5–60 min by fluorography (Fig. 4, A and C). Similarly, cells were incubated in the presence of 1 μ M [14 C]chloroquine in the absence and in the presence of 100 nM sodium azide and 10 mM 2-deoxy-glucose (Fig. 4, B and D). Each value is the mean \pm SD of three experiments in which triplicates were assayed.

moiety). Moreover, both P-gp1 and MRP bind to chloroquine, MK 571, and IAAQ.* Thus, it is likely that P-gp1 and MRP recognize certain structural features in quinoline-based drugs.

The finding that MRP-expressing cells (H69/AR and HL60/AR) are less sensitive to chloroquine than their parental (H69 and HL60) cells is consistent with our photoaffinity labeling results in Fig. 2. Furthermore, incubation of resistant cells with increasing concentrations of chloroquine plus MK 571 reversed the sensitivity of H69/AR, but not H69 cells, to chloroquine. Although MK 571 was shown previously to reverse MRP-mediated drug resistance [51], its inability to reverse completely the cross-resistance of H69/AR cells to chloroquine may be due to its poor drug reversal capacity. Alternatively, it is possible that more than one mechanism mediates the resistance of H69/AR cells to chloroquine.

The mechanism by which MRP mediates resistance to structurally dissimilar drugs is not well understood. It has been suggested that, similar to P-gp1, MRP functions as an energy-dependent drug efflux pump [16, 20, 21, 28, 45, 52].

Consistent with this function, our drug transport studies with [14C]chloroquine showed an energy-dependent decrease in the accumulation of drug in H69/AR and HL60/AR compared with H69 and HL60 cells. The decrease in [14C]chloroquine accumulation in H69/AR versus H69 cells was not due to differences in the intracellular pH since the addition of 10 mM NH₄Cl to both cell lines did not alter their drug accumulation (data not shown). In addition, the intracellular pH of both cell lines was determined previously to be identical, with a pH of 7.55 [53]. Furthermore, we show that in HL60/AR cells [14C]chloroquine efflux is energy dependent. In this study, chloroquine efflux from HL60 and HL60/AR cells was 30 and 60%, respectively. The latter results are consistent with those from an earlier study [54] showing a 15 and 40% efflux of ³Hldaunorubicin from HL60 and HL60/AR cells, respectively. The absence of enhanced chloroquine efflux from H69/AR cells relative to the parental H69 cells is not clear. However, a similar decrease in drug accumulation without detectable drug efflux was also observed in other MRPexpressing cells (e.g. MCF7/VP; [55]). Moreover, previous drug transport studies with H69/AR cells showed both drug

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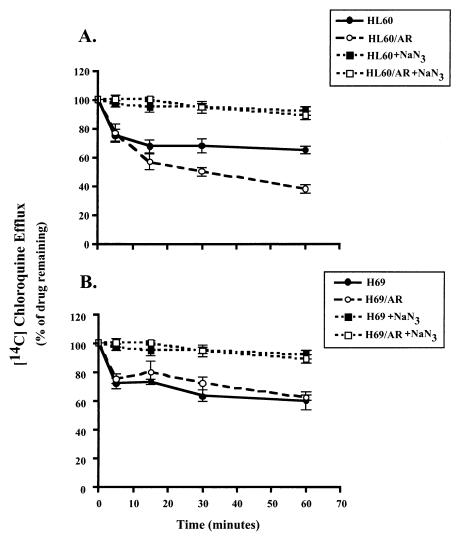


FIG. 5. [14C]Chloroquine efflux from MRP-overexpressing cells and their parental cells. Drug-sensitive (H69 or HL60) and -resistant (H69/AR or HL60/AR) cells were loaded with 10 µM [14C]chloroquine in the absence or presence of 10 mM sodium azide for 30 min after which drug extrusion was potentiated with the administration of 5 mM D-glucose to one half of the samples. For the remaining half of the samples, [14C]chloroquine levels were determined in the presence of sodium azide. Samples were collected, and the amount of drug remaining in the cells was measured by fluorography. Each value is the mean \pm SD of two experiments in which triplicates were assayed.

efflux and accumulation to be unchanged relative to H69 drug-sensitive cells [56]. However, cells transfected with MRP cDNA isolated from H69/AR cells showed a large decrease in vincristine accumulation but only a slight difference in drug efflux [16]. Although the latter findings remain unclear, the localization of MRP to the cell membrane and endosomal vesicles was suggested to be responsible for differences in drug accumulation versus efflux [16]. Taken together, although differences in chloroquine transport in H69/AR and HL60/AR are not clear, these differences may be due to differences in the subcellular distribution of MRP in these two cell lines [45, 46]. Alternatively, other cellular changes in HL60/AR and H69/AR [57-59], in addition to MRP, may be responsible for the differences in chloroquine transport between these two cell lines. Work is underway to confirm these results using cells transfected with human MRP cDNA.

Chloroquine has been extremely effective in the control and prophylaxis of malaria. Unfortunately, resistance to chloroquine and other quinoline-related drugs has rendered these drugs ineffective in many regions where malaria is endemic [60, 61]. Although the mechanism of chloroquine

resistance in the parasite is not well understood, it has been known for some time that resistant malaria accumulates less chloroquine than drug-sensitive malaria [62]. Furthermore, the reduced accumulation of chloroquine is energy dependent [63]. Given the latter characteristics of chloroquineresistant malaria, which are similar to P-gp1-mediated MDR, it was suggested earlier that a P-glycoprotein homologue (Pfmdr1) could mediate chloroquine resistance in malaria [34–36]. However, the role of Pfmdr1 in chloroquine resistance in Plasmodium falciparum remains controversial [37, 64-66]. Given the overlap in MRP and P-gp1 substrate specificity, we speculate about the possible role of an MRP homologue in chloroquine resistance in falciparum malaria. MRP homologues have been isolated from Leishmania tarentolae [67, 68], yeast [69], and Caenorhabditis elegans [70]. Moreover, these MRP homologues were shown to confer resistance to metal oxyanions and antimony [67, 68, 70].

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