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Verapamil-sensitive transport of quinacrine and methylene blue via mutant PfCRT reduces the malaria parasite’s susceptibility to these tricyclic drugs

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ABSTRACT

Background. It is becoming increasingly apparent that certain mutations in the Plasmodium falciparum 'chloroquine resistance transporter' (PfCRT) alter the parasite’s susceptibility to diverse compounds. Here we investigated the interaction of PfCRT with three tricyclic compounds that have been used to treat malaria (quinacrine (QC) and methylene blue (MB)) or to study P. falciparum (acridine orange; AO).

Methods. We measured the antiplasmodial activities of QC, MB, and AO against chloroquine-resistant and chloroquine-sensitive P. falciparum, and determined whether QC and AO affect the accumulation and activity of chloroquine in these parasites. We also assessed the ability of mutant (PfCRTDd2) and wild-type (PfCRTD10) variants of the protein to transport QC, MB, and AO when expressed at the surface of Xenopus laevis oocytes.

Results. Chloroquine-resistance conferring isoforms of PfCRT reduced the sensitivity of the parasite to QC, MB, and AO. In chloroquine-resistant (but not chloroquine-sensitive) parasites, AO and QC increased the parasite’s accumulation of, and susceptibility to, chloroquine. All three compounds were shown to bind to PfCRTDd2, and the transport of QC and MB via this protein was saturable and inhibited by the chloroquine ‘resistance-reverser’ verapamil.

Conclusions. Our findings reveal that the PfCRTDd2-mediated transport of tricyclic antimalarials reduces the parasite’s susceptibility to these drugs.

Keywords. Drug resistance; Plasmodium falciparum; tricyclic drug; methylene blue; PfCRT; fluorescent uptake assay; Xenopus oocytes.
INTRODUCTION

The global campaign to control and eliminate malaria is under serious threat from the emergence and spread of *Plasmodium falciparum* parasites that are resistant to the existing antimalarial drugs [1]. The *P. falciparum* chloroquine resistance transporter (PfCRT) plays a significant role in the phenomenon of drug resistance; mutations in this protein confer resistance to the former ‘wonder drug’ chloroquine (CQ) and also affect the parasite’s susceptibility to a diverse range of molecules, including many of the drugs that have been – or are currently – deployed as antimalarials [e.g., [2] and reviewed in [3]].

PfCRT is located at the membrane of the parasite’s digestive vacuole (DV) [4, 5]. Many quinoline-related drugs, including CQ and quinine, accumulate within this acidic compartment (pH 5-5.5) via ‘weak-base trapping’ [6]. Here they exert an antimalarial effect by preventing the conversion of toxic heme monomers, arising from the parasite’s digestion of host hemoglobin, into the inert crystal hemozoin [7, 8]. Resistance to CQ and quinine is associated with a reduction in the accumulation of these drugs in the DV, and several biochemical studies produced indirect evidence for this being attributable to the efflux of CQ and quinine via mutant PfCRT [9-12]. A direct demonstration of this activity was achieved when the wild-type PfCRT (PfCRTD10) and a resistance-conferring isoform of the protein (PfCRTDd2) were expressed in *Xenopus laevis* oocytes; PfCRTDd2 was shown to possess significant CQ and quinine transport activity, whereas PfCRTD10 did not [13-15]. These findings aside, much remains to be understood about the protein’s ability to recognize and transport diverse compounds.

Here we investigated interactions of PfCRT with quinacrine (QC), methylene blue (MB), and acridine orange (AO). All three molecules contain a tricyclic scaffold (Supplementary Figure 1) and will sequester within the DV due to weak-base trapping or, in the case of the
cationic MB, via weak-base trapping and/or oxidation of the reduced forms of the drug [16-19].

AO has been employed as a fluorescent dye for the diagnosis of malaria [20] and as a probe for estimating the pH of the DV (pH_{DV}) in CQ-sensitive and CQ-resistant parasites [21-23]. The latter studies reported that CQ-resistant strains have a lower pH_{DV} than their CQ-sensitive counterparts (pH 5.21 versus pH 5.64), and that this could account for the differential accumulation of, and susceptibility to, CQ between these parasites. Alternatively, it has been suggested that the observed difference in the accumulation of AO may instead be due to its export from the DV via mutant PfCRT [24].

MB was the first synthetic antimalarial drug and QC, which was derived from MB, was the first synthetic antimalarial to be widely used. Both drugs are thought to inhibit heme detoxification [17, 18, 25], with the redox-cycling capacity of MB potentially exerting an additional parasiticidal effect [18, 26]. Although QC was set aside following the introduction of CQ, structural derivatives of QC have since been deployed (pyronaridine) or are undergoing development [27, 28] as antimalarials. Interest in MB, which until recently was a largely overlooked antimalarial, has been greatly revived by studies that have found it to be (1) a potent gametocytocidal and transmission-blocking agent [29, 30], (2) a potential partner drug that improves the clinical efficacy of amodiaquine and artesunate-amodiaquine [31, 32], and (3) a relatively well-tolerated drug [31-33].

A number of in vitro studies have observed cross-resistance between QC and CQ [34-36] and these observations, along with the finding that QC increases the accumulation of CQ in CQ-resistant but not CQ-sensitive parasites [37], implicate mutations in PfCRT as a determinant of the parasite's susceptibility to QC. The relationship between CQ and MB susceptibility in CQ-resistant P. falciparum is considerably less clear. Several studies have reported that CQ resistance (and specifically mutant PfCRT) does not impart cross-resistance to MB [38-40], whereas others have observed modest but significant positive correlations.
between the half-maximal inhibitory concentrations (IC\textsubscript{50}s) of CQ and MB in \textit{P. falciparum} [17, 41, 42].

We have sought to elucidate the mechanisms underpinning these observations by ascertaining the nature and strength of the interactions between PfCRT and QC, MB, and AO. Our findings reveal that CQ resistance-conferring isoforms of PfCRT reduce the parasite’s susceptibility to tricyclic compounds and that, at least in the cases of QC and MB, this phenomenon is due to the ability of mutant PfCRT to transport these drugs away from their main site of action.

\textbf{MATERIALS AND METHODS}

\textbf{Culture of \textit{P. falciparum}-infected Erythrocytes}

The use of human blood in this study was approved by the Australian National University’s Human Research Ethics Committee. Two CQ-sensitive strains (D10 and 3D7), three CQ-resistant strains (K1, RSA11, and Dd2), and three \textit{pfcrt} transfectant lines (C2\textsubscript{GC03}, C4\textsubscript{Dd2}, and C6\textsubscript{7G8}; [43]) were cultured and synchronized as described previously [12].

\textbf{\textit{P. falciparum} Drug Susceptibility and Drug Accumulation}

Measurements of parasite proliferation, and of [\textsuperscript{3}H]CQ and [\textsuperscript{3}H]AO in accumulation in trophozoite-infected erythrocytes, were performed according to the protocols provided in the Supplementary Materials.

\textbf{Drug Transport in \textit{Xenopus} Oocytes Expressing PfCRT}

Oocytes were harvested and prepared using the method described in detail in the Supplementary Materials. PfCRT\textsuperscript{Dd2} and PfCRT\textsuperscript{D10} were expressed at the oocyte plasma membrane using an approach outlined elsewhere [13]. The uptake into oocytes of [\textsuperscript{3}H]CQ,
well as that of unlabelled QC, MB, and AO (all of which fluoresce), was measured using the protocols described in the Supplementary Materials.

Statistics

Statistical comparisons were made using the Student’s t-test for paired or unpaired samples, or ANOVA in conjunction with Tukey’s multiple comparisons test.

RESULTS

Mutant Forms of PfCRT Decrease the Susceptibility of P. falciparum to QC, MB, and AO

The antiplasmodial activities of QC, MB, and AO were evaluated in vitro against erythrocytes infected with field-derived CQ-sensitive (D10 or 3D7) or CQ-resistant (RSA11, K1, or Dd2) P. falciparum strains. The D10 and 3D7 strains carry the wild-type version of PfCRT (PfCRT\textsubscript{D10}), K1 and Dd2 parasites carry resistance-conferring isoforms of the protein (PfCRT\textsubscript{K1} and PfCRT\textsubscript{Dd2}, respectively), and the PfCRT haplotype carried by RSA11 is currently unknown. CQ was included as a reference drug and the resulting IC\textsubscript{50}s are presented in Table 1. QC and MB displayed potent antimalarial activities, with IC\textsubscript{50}s against 3D7 parasites of approximately 10 and 2.5 nM, respectively. However, these values increased 1.6 to 2-fold when the drugs were tested against the Dd2 strain. AO was the least active of the test compounds, with IC\textsubscript{50}s of ~300 nM against the D10 and 3D7 strains, and these values also increased 1.9 to 2.3-fold in the CQ-resistant strains. The role of PfCRT in the observed decrease in the sensitivities of the CQ-resistant strains to QC, MB, and AO was investigated using a set of isogenic transfectant lines [43], which express the wild-type \textit{pfcr}\textit{t} allele (C2\textsuperscript{GC03}) or a mutant \textit{pfcr}\textit{t} allele from either Dd2 (C4\textsuperscript{Dd2}) or the CQ-resistant strain 7G8 (C6\textsuperscript{7G8}). Consistent with what was observed for the field-derived strains, the IC\textsubscript{50}s determined for QC, MB, AO, and CQ were higher in the CQ-resistant lines (C4\textsuperscript{Dd2} and C6\textsuperscript{7G8}) than in the CQ-sensitive C2\textsuperscript{GC03} line (Table 1). CQ-resistant
parasites can be partially re-sensitised to CQ in vitro by verapamil (VP) or chlorpheniramine (CP) ([44] and Table 1) – a phenomenon we have shown to be attributable to the ability of these drugs to inhibit CQ transport via mutant PfCRT [13, 14, 45]. In the data presented in Table 1, the presence of 1 µM VP decreased the QC and MB IC\textsubscript{50}s in all of the CQ-resistant strains and lines, and was without effect in their drug-sensitive counterparts. Taken together, these findings suggested that the decreased sensitivity of CQ-resistant parasites to QC and MB was due to the interaction of these drugs with mutant isoforms of PfCRT.

VP did not have a significant effect on the activity of AO in any of the parasite types (Table 1). We therefore explored the interaction between AO and PfCRT further by testing whether AO, like VP or CP, potentiates the activity of CQ against CQ-resistant parasites. An initial experiment revealed that the presence of 250 nM AO or 1 µM VP decreased the CQ IC\textsubscript{50} in the K1 strain by ~30% and 65%, respectively, whereas no significant change was observed in the 3D7 strain (Supplementary Table 1). Isobolograms describing the interactions of CQ with AO, and of VP with AO, against the C2\textsubscript{GCO3} and C4\textsubscript{Dd2} lines indicated these combinations are additive in C2\textsubscript{GCO3} parasites, whereas interactions suggestive of synergism were observed in the C4\textsubscript{Dd2} line (Figure 1). These results suggested that the antimalarial activity of combinations of AO with CQ or VP is dependent on the \textit{pfcrt} allele expressed and that AO may behave as a reverser of CQ resistance.

\textbf{QC and AO Increase [\textsuperscript{3}H]CQ Accumulation in CQ-resistant Parasites}

The resistance-reversing activity of VP causes an increase in the accumulation of [\textsuperscript{3}H]CQ in CQ-resistant strains, with no such change occurring in drug-sensitive parasites [44]. To investigate whether AO enhances the effects of CQ via the same mechanism, [\textsuperscript{3}H]CQ accumulation was measured in CQ-sensitive and CQ-resistant parasites in the presence or absence of AO, QC, and VP. We found that AO and QC increased the accumulation of [\textsuperscript{3}H]CQ in CQ-resistant parasites when present at 100 nM and 1 µM, respectively (Figure 2). By contrast,
1 μM AO had no effect on [3H]CQ accumulation in the drug-sensitive parasites, whereas 100 nM QC caused a modest decrease in [3H]CQ accumulation. The latter result suggests that QC reduces the amount of free heme, and therefore CQ-binding sites, within the DV of drug-sensitive parasites, whereas the observed increase in [3H]CQ accumulation in the resistant line indicates that QC possesses a degree of resistance-reversing activity. Moreover, the finding that 1 μM AO and 5 μM VP increased the accumulation of [3H]CQ in the C4Dd2 line to similar extents (Figure 2D) further supported the hypothesis that AO potentiates the activity of CQ against CQ-resistant parasites by blocking the PfCRT-mediated efflux of CQ from the DV.

QC, MB, and AO Inhibit the Transport of [3H]CQ via PfCRTDd2

Direct measurements of the abilities of QC, MB, and AO to block CQ transport via PfCRTDd2 were undertaken using the Xenopus oocyte system. In a preliminary assay in which the tricyclic compounds were each tested at 500 μM, QC and AO caused pronounced reductions in the PfCRTDd2-mediated transport of CQ, whereas MB caused a modest level of inhibition (Figure 3A). An analysis of the concentration-dependence of inhibition yielded IC50s of 13.5 ± 0.9 μM, 41.0 ± 5.1 μM, and 1.3 ± 0.2 mM for QC, AO, and MB, respectively (Figures 3B-D).

QC and MB are Substrates of PfCRTDd2

We further investigated the nature of the interactions of PfCRTDd2 with QC, MB, and AO by performing direct measurements of transport in the Xenopus oocyte system. We achieved this by re-purposing the radioisotope influx assay for the detection of fluorescent substrates (see Supplementary Materials). In all cases, there was a linear relationship between concentration and fluorescence intensity for QC, MB, and AO (Supplementary Figure 2). It was therefore possible to use the native fluorescence of these compounds to measure their uptake into oocytes expressing different forms of PfCRT.
Oocytes expressing PfCRT^{Dd2} were found to accumulate significantly more QC and MB than either non-injected oocytes or oocytes expressing PfCRT^{D10}, consistent with PfCRT^{Dd2}, but not PfCRT^{D10}, mediating the transport of these tricyclic drugs (Figures 4A and 5A; \( P < 0.001 \) at 60 minutes). The uptake of QC and MB via PfCRT^{Dd2} was linear over time for at least 2 hours (Figures 4B and 5B) and the estimated rates of PfCRT^{Dd2}-mediated transport (in picomoles hour\(^{-1}\) per oocyte) were 7.01 ± 1.34 (QC) and 4.73 ± 0.14 (MB). An analysis of the concentration-dependence of QC and MB uptake revealed, in both cases, that the component of transport attributable to PfCRT^{Dd2} was saturable (Figures 4C and 5C). A least squares fit of the Michaelis-Menten equation to the data (Figures 4D and 5D) yielded apparent Michaelis constants (\( K_m \)) of 9.8 ± 1.1 and 123 ± 11.5 µM and maximal velocities (\( V_{max} \)) of 16.3 ± 0.7 and 202 ± 5 picomoles hour\(^{-1}\) per oocyte for QC and MB, respectively. By comparison, in non-injected oocytes and those expressing PfCRT^{D10}, the rate of QC and MB influx increased in a roughly linear manner with increasing concentrations of the drug (Figures 4C and 5C). This finding, together with the observation that these two oocyte types took up QC or MB to similarly low levels, indicates that this component of accumulation is due to the simple diffusion of the uncharged drug into the oocyte.

Several known inhibitors of PfCRT^{Dd2} were tested for their effect on QC and MB uptake in oocytes expressing PfCRT^{D10} or PfCRT^{Dd2}, including VP (250 µM), CQ (750 µM), neurotensin fragment 8-13 (NT; 250 µM) and, in the case of MB uptake, saquinavir (SQV; 250 µM) – which is the most potent of the inhibitors tested [46]. None of the test compounds affected the accumulation of QC or MB in the PfCRT^{D10}-expressing oocytes (Figures 4E and 5E). By contrast, VP reduced the PfCRT^{Dd2}-mediated uptake of QC and MB by 86 ± 1.6 and 65 ± 4 percent, respectively, and SQV caused a 71 ± 3 percent decrease in the transport of MB via PfCRT^{Dd2} (Figures 4F and 5F). Compared with VP, CQ and NT were less potent inhibitors of QC transport (71 ± 3 and 48 ± 4 percent decreases, respectively) and caused only modest reductions in the PfCRT^{Dd2}-mediated uptake of MB (30 ± 5 and 33 ± 2 percent decreases,
Attempts to detect the transport of AO (75 or 150 μM) via PfCRT were performed at pH 4.5, 5.0, 5.5, and 6.0, and a second series of experiments also made measurements of [3H]AO transport over the same pH range. Although several of the assays conducted at pH 5.0 appeared to detect a small increase in AO accumulation in the PfCRTDd2-expressing oocytes, the signal was not reproduced in the majority of the assays carried out. This suggested that AO is either not a substrate of PfCRTDd2 or that the protein has a very low capacity for AO transport, such that – at least in the oocyte system – it is obscured by the rate of uptake via simple diffusion of the uncharged species (the proportion of AO in the uncharged form is relatively high, even at pH 5.0; Supplementary Table 2). The first scenario raised the possibility that PfCRTDd2 (and PfCRT7G8) decreases the parasite’s susceptibility to AO without mediating the efflux of AO from the DV. We therefore compared the accumulation of [3H]AO between the C2GC03, C4Dd2, and C67G8 lines and found there to be a modest but significant decrease in [3H]AO accumulation in the CQ-resistant lines (Figure 6). Consistent with the observed failure of VP to increase the activity of AO against CQ-resistant parasites, VP did not produce a PfCRT-dependent increase in the accumulation of [3H]AO (data not shown) and assays performed with the CQ resistance-reverser CP suggest it may, if anything, cause a small decrease in [3H]AO accumulation in the CQ-resistant lines (Supplementary Figure 3).

**DISCUSSION**

PfCRT is a significant player in the malaria parasite’s steadily expanding resistance to antimalarial drugs, yet relatively little is understood about the extent of its substrate-specificity or the pharmacology of its transport activity. In this study, assays conducted with isogenic transfectant lines, which differ only in the isoform of PfCRT they express, revealed that CQ resistance-conferring isoforms of PfCRT confer a VP-sensitive decrease in the
parasite’s susceptibility to the tricyclic drugs QC and MB. Characterization of the interaction of these drugs with a mutant isoform of PfCRT (PfCRT\textsuperscript{Dd2}) in the \textit{Xenopus} oocyte system provided a mechanistic explanation for this phenomenon; PfCRT\textsuperscript{Dd2} can efflux QC and MB out of the parasite’s DV and VP inhibits this transport activity. Our findings are also consistent with QC and MB exerting their main schizontocidal activity within the DV, where they are thought to prevent the detoxification of heme and where MB is also likely to disrupt vital redox processes.

We also found that parasites carrying a mutant isoform of PfCRT accumulated less of the pH probe AO, and were more resistant to its antiplasmodial activity, than parasites carrying the wild-type version of PfCRT – but these phenomena were not affected by VP. AO did, however, behave as a CQ resistance-reverser; it inhibited CQ transport via PfCRT\textsuperscript{Dd2} in the oocyte system (IC\textsubscript{50} of 41 µM) and chemosensitized CQ-resistant parasites to CQ. The simplest explanation for these observations is that AO binds to mutant PfCRT and is translocated at a low rate, such that it is not consistently detected above the simple diffusion of uncharged AO into the oocyte. These findings render AO unsuitable for comparing the pH\textsubscript{DV} between CQ-sensitive and CQ-resistant parasites, since the efflux of the probe via mutant PfCRT will result in non-pH-dependent changes in the accumulation of AO. Indeed, no significant difference was observed between CQ-sensitive and CQ-resistant parasites when the pH\textsubscript{DV} was measured with several other pH-sensitive dyes [24].

Our characterization of the interactions between mutant PfCRT and QC in the oocyte system revealed QC transport via PfCRT\textsuperscript{Dd2} to be a high-affinity and low-capacity process (e.g., \textit{K\textsubscript{m}}(QC) of 9.8 µM compared with \textit{K\textsubscript{m}}(CQ) of 232-245 µM [13, 15]) and QC to be a strong inhibitor of the PfCRT\textsuperscript{Dd2}-mediated transport of CQ (QC is comparable in potency to SQV; IC\textsubscript{50}s of 13.5 and 13 µM [46], respectively). These data, which indicate that QC can outcompete CQ for transport via PfCRT\textsuperscript{Dd2}, provide a mechanistic explanation for the ability of QC to increase \textsuperscript{3}H\textsubscript{CQ} accumulation in CQ-resistant parasites.
The finding that the parasite’s sensitivity to MB is reduced by the PfCRT-mediated resistance mechanism is consistent with several studies in which a significant correlation between MB and CQ susceptibilities was observed in *P. falciparum* [17, 41, 42], but is at odds with reports of there being no such relationship [38-40]. The basis for this difference is not clear, but may relate to differences in the parasite proliferation protocols between labs. For example, even when the same parasite strains were studied (CQ-sensitive ‘D6’ and CQ-resistant ‘W2’), one group observed modest but significant cross-resistance between CQ and MB [17] whereas another did not [39]. We found PfCRT<sup>Dd2</sup> to be a medium-affinity and very high-capacity transporter of MB (e.g., the $V_{\text{max}}$ values (in picomoles hour<sup>-1</sup> per oocyte) for MB and CQ are 202 and 61-67 [13, 15], respectively), which may explain, at least in part, why VP and CP inhibited MB uptake to a much lesser extent than what was observed for QC uptake. However, it is also worth noting that MB was a very poor inhibitor of CQ uptake ($IC_{50}$ of 1.3 mM) and vice versa. We recently provided evidence for there being at least two distinct, but antagonistically-interacting, substrate-binding sites within PfCRT<sup>Dd2</sup> and proposed this region of the transporter most likely resembles a cavity with multiple points for substrate interaction [14]. Hence, the relatively poor ability of MB and CQ to interfere with each other’s transport could arise from these two drugs binding to largely different sets of attachment points. Likewise, the inability of VP to decrease the AO $IC_{50}$ (and of CP to increase $[^3]H$AO accumulation) in CQ-resistant parasites may be due to there being little or no overlap between the attachment points occupied by AO and VP (or CP).

The high-capacity transport of MB via mutant PfCRT, and the resulting reduction in the parasite’s susceptibility to the drug, could have future clinical relevance given the potential of MB to serve as a potent transmission-blocking agent and its inclusion in two candidate combination therapies (artesunate-amodiaquine-MB and amodiaquine-MB). It is conceivable that parasites carrying mutant isoforms of PfCRT will be selected by MB-containing treatments, forming a population from which high-level resistance could evolve. Whether this
transpires will depend on a number of parameters (none of which are currently known), such
as the approximate concentration of MB within the DV and the extent to which it can be
lowered by mutant PfCRT, how this fluctuates during the treatment course, and for what
period the level of MB within the DV falls beneath the minimum parasiticidal concentration.
Moreover, these factors are likely to be affected by the co-administration of MB with
amodiaquine, since the latter is suspected to be a substrate of mutant PfCRT [12, 47]. It is also
possible that the gametocytocidal activity of MB could be affected by the PfCRT-mediated
resistance mechanism; although mature gametocytes no longer digest hemoglobin, they
retain both PfCRT as well as vesicles that could be the remnant of, or replacement for, the DV
[48, 49].

The data presented here have (1) broadened the known substrate range of mutant
PfCRT to include tricyclic compounds and (2) established that the PfCRT-mediated efflux of
QC and MB from the DV decreases the parasite’s sensitivity to these molecules. When taken
together with previous studies [13-15], these results reveal a shared mechanism of resistance
for CQ, quinine, quinidine, QC, and MB. Our work therefore adds further support to the
proposal that PfCRT behaves as a ‘multidrug resistance carrier’ [3, 50] and highlights the
potential for there being an interaction between mutant PfCRT and existing and upcoming
antimalarials with quinoline- and acridine-related structures (e.g., ferroquine and
pyronaridine).
**FIGURE LEGENDS**

**Figure 1.** Isobolograms describing the interactions of verapamil (VP) with chloroquine (CQ), acridine orange (AO) with CQ, and VP with AO, against erythrocytes infected with C2GCO3 or C4Dd2 parasites. The data represent the mean ± SEM from 3 independent experiments (performed on different days), within which measurements were averaged from 3 replicates. Where not shown, error bars fall within the symbols. In general, a fractional inhibitory concentration (FIC) of 1.0 suggests that the effects of the two compounds are additive, a FIC ≤ 0.5 suggests synergism, and a FIC ≥ 4.0 indicates antagonism. The mean FICs for the C2GCO3 line were as follows: VP and CQ combination, 1.04; AO and CQ combination, 1.00; VP and AO combination, 0.99. None of these FICs appeared different from 1.0, consistent with all three drug combinations being additive in the CQ-sensitive parasite line. The mean FICs for the C4Dd2 line were as follows: VP and CQ combination, 0.54; AO and CQ combination, 0.94; VP and AO combination, 0.84. In all three cases, the mean FICs were below 1.0, but were also above the accepted threshold for categorizing an interaction as synergistic. This finding suggests that the drug combinations were slightly synergistic in the CQ-resistant parasite line.

**Figure 2.** Effects of quinacrine (QC), acridine orange (AO), and verapamil (VP) on the accumulation of radiolabelled chloroquine ([3H]CQ) by erythrocytes infected with mature trophozoite-stage parasites. Panels A, B, and C show data for CQ-sensitive parasites (C2GCO3, D10, and 3D7, respectively). Panels D, E, and F show data for CQ-resistant parasites (C4Dd2, RSA11, and K1, respectively). CQ accumulation is expressed in terms of the fold difference in the [3H]CQ ‘distribution ratio’, which corresponds to the concentration of [3H]CQ within the infected cells relative to the concentration in the extracellular medium. In panels A and D, the data represent the mean ± SEM from 6 independent experiments (performed on different days), within which measurements were averaged from 3 replicates. Panels B, C, E, and F
show the mean + range/2 of 2 independent experiments (performed on different days), within which measurements were averaged from 3 replicates. In panel E, the SEM for the 100 nM AO treatment was very small and the error bar is therefore not visible. The asterisks denote a significant difference in $[^3]$H CQ accumulation between the control treatment and that measured in the presence of QC, AO, or VP: *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$ (ANOVA).

**Figure 3.** Quinacrine (QC), acridine orange (AO), and methylene blue (MB) inhibit the transport of radiolabelled chloroquine ($[^3]$H CQ) via the *Plasmodium falciparum* chloroquine resistance transporter (PfCRT) in *Xenopus laevis* oocytes. A, $[^3]$H CQ uptake into oocytes expressing the Dd2 isoform of PfCRT (PfCRT$^{Dd2}$; black bars) or the D10 isoform of PfCRT (PfCRT$^{D10}$; white bars) in the presence of 500 μM unlabelled CQ, QC, AO, or MB. B-D, Concentration-dependent inhibition of the uptake of $[^3]$H CQ into oocytes expressing PfCRT$^{Dd2}$ (black circles) or PfCRT$^{D10}$ (white circles) by QC, AO, or MB. In all panels, $[^3]$H CQ uptake is expressed relative to that measured in the PfCRT$^{Dd2}$-expressing oocytes under control conditions. The data represent the mean ± SEM from 3 to 5 independent experiments (performed on different days and using oocytes from different frogs), within which measurements were made from 10 oocytes per treatment. Where not shown, error bars fall within the symbols. Note that non-injected oocytes and oocytes expressing PfCRT$^{D10}$ take up $[^3]$H CQ to similar (low) levels via simple diffusion of the neutral species; this represents the “background” level of CQ accumulation in oocytes. In panel A, the asterisks denote a significant difference in $[^3]$H CQ uptake between the control PfCRT$^{Dd2}$ treatment and that measured in the presence of CQ, QC, AO, or MB: ***, $P < 0.001$ (ANOVA). Abbreviation: IC$_{50}$, half-maximal inhibitory concentration.

**Figure 4.** Quinacrine (QC) transport properties of the *Plasmodium falciparum* chloroquine
resistance transporter (PfCRT) in *Xenopus laevis* oocytes. A, Oocytes expressing the Dd2 isoform of PfCRT (PfCRT\textsuperscript{Dd2}; black circles) showed a marked increase in QC accumulation relative to non-injected oocytes (black triangles) and to those expressing the D10 isoform of PfCRT (PfCRT\textsuperscript{D10}; white circles). Rates of QC uptake (picomoles hour\textsuperscript{-1} per oocyte; \(n = 3 \pm \) SEM, estimated from uptake at 60 minutes) were as follows: non-injected, 1.74 ± 0.37; PfCRT\textsuperscript{D10}, 1.50 ± 0.20; PfCRT\textsuperscript{Dd2}, 8.62 ± 1.58. B, The PfCRT\textsuperscript{Dd2}-mediated uptake of QC (obtained by subtracting the average of the uptake measured in non-injected and PfCRT\textsuperscript{D10}-expressing oocytes from that measured in PfCRT\textsuperscript{Dd2}-expressing oocytes) was approximately linear with time for at least 2 hours. The rate of QC transport via PfCRT\textsuperscript{Dd2}, estimated from uptake at 60 minutes, was 7.01 ± 1.34 picomoles hour\textsuperscript{-1} per oocyte (\(n = 3 \pm \) SEM). C, The effect of QC concentration on QC uptake by non-injected oocytes (black triangles) and oocytes expressing PfCRT\textsuperscript{D10} (white circles) or PfCRT\textsuperscript{Dd2} (black circles). The accumulation of QC by the control oocytes increased linearly with increasing concentrations of QC, although a second component – attributable perhaps to the binding of QC to the oocyte surface – may have been present below 2.5 μM. D, The concentration-dependence of the PfCRT\textsuperscript{Dd2}-mediated transport of QC (in picomoles hour\textsuperscript{-1} per oocyte). E, Uptake of QC into oocytes expressing PfCRT\textsuperscript{D10} (white bars) or PfCRT\textsuperscript{Dd2} (black bars) in the presence of 750 μM chloroquine (CQ), 250 μM verapamil (VP), or 250 μM neurotensin (NT). F, Inhibition of the PfCRT\textsuperscript{Dd2}-mediated transport of QC by VP, CQ, or NT. The extracellular concentration of QC was 40 μM (panels A and B), 2.5-40 μM (panels C and D), or 10 μM (panels E and F). In all panels except D, QC uptake is expressed relative to that measured in the relevant PfCRT\textsuperscript{Dd2} control. The data represent the mean ± SEM from 3 to 11 independent experiments (performed on different days and using oocytes from different frogs), within which measurements were made from 12 oocytes per treatment. Where not shown, error bars fall within the symbols. In panels E and F, the asterisks denote a significant difference in QC uptake between the control PfCRT\textsuperscript{Dd2} treatment and that measured in the presence of VP, CQ, or NT: ***, \(P < 0.001\) (ANOVA).
Abbreviations: $K_m$, apparent Michaelis constant; $V_{max}$, maximum velocity.

Figure 5. Methylene blue (MB) transport properties of the *Plasmodium falciparum* chloroquine resistance transporter (PfCRT) in *Xenopus laevis* oocytes. A, Oocytes expressing the Dd2 isoform of PfCRT (PfCRT\textsuperscript{Dd2}; black circles) showed a marked increase in MB accumulation relative to non-injected oocytes (black triangles) and to those expressing the D10 isoform of PfCRT (PfCRT\textsuperscript{D10}; white circles). Rates of MB uptake (picomoles hour\textsuperscript{-1} per oocyte; $n = 3 \pm$ SEM, estimated from uptake at 60 minutes) were as follows: non-injected, 1.61 ± 0.07; PfCRT\textsuperscript{D10}, 1.59 ± 0.08; PfCRT\textsuperscript{Dd2}, 6.33 ± 0.22. B, The PfCRT\textsuperscript{Dd2}-mediated uptake of MB (obtained by subtracting the average of the uptake measured in non-injected and PfCRT\textsuperscript{D10}-expressing oocytes from that measured in PfCRT\textsuperscript{Dd2}-expressing oocytes) was approximately linear with time for at least 2 hours. The rate of MB transport via PfCRT\textsuperscript{Dd2}, estimated from uptake at 60 minutes, was 4.73 ± 0.14 picomoles hour\textsuperscript{-1} per oocyte ($n = 3 \pm$ SEM). C, The effect of MB concentration on MB uptake by non-injected oocytes (black triangles) and oocytes expressing PfCRT\textsuperscript{D10} (white circles) or PfCRT\textsuperscript{Dd2} (black circles). The accumulation of MB by the control oocytes increased linearly with increasing concentrations of MB between 150 μM and 2 mM MB. A second component was observed below 150 μM; this probably reflects the binding of MB to the oocyte surface. D, The concentration-dependence of the PfCRT\textsuperscript{Dd2}-mediated transport of MB (in picomoles hour\textsuperscript{-1} per oocyte). E, Uptake of MB into oocytes expressing PfCRT\textsuperscript{D10} (white bars) or PfCRT\textsuperscript{Dd2} (black bars) in the presence of 250 μM saquinavir (SQV), 750 μM chloroquine (CQ), 250 μM verapamil (VP), or 250 μM neurotensin (NT). F, Inhibition of the PfCRT\textsuperscript{Dd2}-mediated transport of MB by SQV, VP, CQ, or NT. The extracellular concentration of MB was 75 μM (panels A, B, E and F) or 0.025-2 mM (panels C and D). In all panels except D, MB uptake is expressed relative to that measured in the relevant PfCRT\textsuperscript{Dd2} control. The data represent the mean ± SEM from 3 to 5 independent experiments (performed on different days and using oocytes from different frogs), within
which measurements were made from 12 oocytes per treatment. Where not shown, error bars fall within the symbols. In panels E and F, the asterisks denote a significant difference in MB uptake between the control PfCRT<sup>Dd2</sup> treatment and that measured in the presence of SQV, VP, CQ, or NT: **, \( P < 0.01 \); ***, \( P < 0.001 \) (ANOVA). Abbreviations: \( K_m \), apparent Michaelis constant; \( V_{\text{max}} \), maximum velocity.

**Figure 6.** Accumulation of radiolabelled chloroquine ([³H]CQ) or radiolabelled acridine orange ([³H]AO) by erythrocytes infected with mature trophozoite-stage parasites. The white, black, and gray bars show data for the C2<sup>GCO3</sup>, C4<sup>Dd2</sup>, and C6<sup>7G8</sup> parasite lines, respectively. The accumulation of [³H]drug was expressed as a ‘distribution ratio’ (i.e., the concentration of [³H]drug within the infected cells relative to the concentration in the extracellular medium) and is shown relative to the ratio measured in the C2<sup>GCO3</sup> parasites. Under control conditions, the distribution ratios for CQ and AO in the C2<sup>GCO3</sup> line were 773 ± 95 and 299 ± 31, respectively. The data represent the mean ± SEM of 4 independent experiments (performed on different days), within which measurements were averaged from 2 replicates. The asterisks denote a significant difference in [³H]drug accumulation between the C2<sup>GCO3</sup> line and that measured in the C4<sup>Dd2</sup> or C6<sup>7G8</sup> lines: *, \( P < 0.05 \); **, \( P < 0.01 \); ***, \( P < 0.001 \) (ANOVA, with “experiment” nominated as a “blocking factor”).

**Acknowledgements**

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**References**


26. Ehrhardt, K., E. Davioud-Charvet, H. Ke, A.B. Vaidya, M. Lanzer, M. Deponte. The antimalarial activities of methylene blue and the 1,4-naphthoquinone 3-[4-


A

B

C

D

E

F

Uptake of MB (% PICRT<sup>Δgs</sup> at 1 hr)

Time (min)

PICRT<sup>Δgs</sup>-mediated uptake of MB (% PICRT<sup>Δgs</sup> at 1 hr)

Time (min)

Uptake of MB (% PICRT<sup>Δgs</sup> at 1 hr)

[MB] (mM)

Uptake of MB (% PICRT<sup>Δgs</sup> at 1 hr)

[MB] (mM)

K<sub>m</sub> = 123 ± 11.5 μM
V<sub>max</sub> = 202 ± 5 pmol / oocyte / hr

Uptake of MB (% PICRT<sup>Δgs</sup> control)

Stages

Control + 250 μM SQV + 250 μM VP + 750 μM CQ + 250 μM NT

Uptake of MB (% PICRT<sup>Δgs</sup> control)

Stages

Control + 250 μM SQV + 250 μM VP + 750 μM CQ + 250 μM NT
Table 1. In Vitro Antiplasmodial Activities of Chloroquine (CQ), Quinacrine (QC), Acridine Orange (AO), and Methylene Blue (MB) against CQ-Sensitive and CQ-Resistant *P. falciparum* Parasites

<table>
<thead>
<tr>
<th>Strain/lineb</th>
<th>Chloroquine</th>
<th>Quinacrine</th>
<th>Acridine orange</th>
<th>Methylene blue</th>
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<tr>
<td></td>
<td>Control + 1 μM VP</td>
<td>Control + 1 μM VP</td>
<td>Control + 1 μM VP</td>
<td>Control + 1 μM VP</td>
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<td>D10</td>
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<td>27 ± 2.5</td>
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<td>-</td>
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<tr>
<td></td>
<td>21 ± 0.5</td>
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<td></td>
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<td>309 ± 34</td>
<td>294 ± 24</td>
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<td>2.5 ± 0.08</td>
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<tr>
<td>RSA11</td>
<td>354 ± 39c</td>
<td>43 ± 5.5f</td>
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<td>-</td>
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<tr>
<td></td>
<td>586 ± 39g</td>
<td>563 ± 84</td>
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<tr>
<td>K1</td>
<td>255 ± 26c</td>
<td>63 ± 8.4f</td>
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<td>-</td>
</tr>
<tr>
<td>Dd2</td>
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<td>26 ± 2.4f</td>
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<td>549 ± 30</td>
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<td>491 ± 19</td>
<td>501 ± 20</td>
<td>4.7 ± 0.05c</td>
<td>2.5 ± 0.03f</td>
</tr>
</tbody>
</table>

Abbreviations: IC<sub>50</sub>, half-maximal inhibitory concentrations; VP, verapamil.

*The IC<sub>50</sub>s are shown as the mean ± SEM from 3 to 11 independent experiments (performed on different days), within which measurements were averaged from 2 or 3 replicates.

*Field-derived CQ-sensitive strains: D10 (Papua New Guinea); 3D7 (Africa). Field-derived CQ-resistant strains: RSA11 (South Africa); K1 (Thailand); Dd2 (Indochina/Laos). Isogenic pfcr<sup>7</sup>t transfectant lines: C2<sup>GCO3</sup> (CQ-sensitive); C4<sup>Dd2</sup> (CQ-resistant); C6<sup>7G8</sup> (CQ-resistant).

*P* values determined from Student *t*-test comparisons with the relevant CQ-sensitive strain or line (and within the same drug treatment) were less than c0.001, d0.01, or e0.05.

*P* values determined from Student *t*-test comparisons with the control treatment from the same CQ-sensitive strain or line (and within the same drug treatment) were less than f0.001, g0.01, or h0.05.
Supplementary Materials for

Verapamil-sensitive transport of quinacrine and methylene blue via mutant PfCRT reduces the malaria parasite’s susceptibility to these tricyclic drugs

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Supplementary Materials and Methods

Supplementary Tables 1 and 2

Supplementary Figures 1, 2, and 3

References
Supplementary Materials and Methods

Measurements of *P. falciparum* Drug Susceptibility

Parasite proliferation was measured in 96-well plates using either the \[^3\text{H}\]hypoxanthine incorporation method or the lactate dehydrogenase (pLDH) assay. The \[^3\text{H}\]hypoxanthine incorporation assay was performed over 72 hours using a protocol described elsewhere [1], and the starting hematocrit and parasitemia were both approximately 1%. The Malstat\textsuperscript{TM} reagent (Flow Inc.) was used to measure pLDH activity in accordance with the method of Makler and colleagues [2], and these assays were performed over 48 hours with a starting hematocrit and parasitemia of approximately 1% and 2%, respectively. The level of parasite proliferation in the presence of each drug concentration was expressed as a percentage of the proliferation measured in the absence of drug. The IC\textsubscript{50} (i.e., the concentration of drug at which the inhibition of parasite growth is half-maximal) was determined in SigmaPlot by a least-squares fit of the equation \( y = a/[1+([\text{drug}]/\text{IC}_{50})c] \) to the data, where \( y \) is the percent parasite proliferation, \( a \) is the maximum change in the percent parasite proliferation, and \( c \) is a fitted constant. The IC\textsubscript{50}s obtained using the \[^3\text{H}\]hypoxanthine incorporation method were indistinguishable from those generated with the pLDH assay.

Antimalarial drug combinations were assessed using fixed-ratio isobolograms [3] as described previously [4]. Fractional inhibitory concentrations (FICs) were calculated for each drug within a combination as follows: FIC (A) = the IC\textsubscript{50} for drug A in a combination divided by the IC\textsubscript{50} for drug A alone; FIC (B) = the IC\textsubscript{50} for drug B in a combination divided by the IC\textsubscript{50} for drug B alone. The isobolograms were constructed by plotting FIC (A) against FIC (B) for each combination. It is generally accepted that a drug combination is synergistic if the FIC is \( \leq 0.5 \), antagonistic if the FIC is \( \geq 4.0 \) [5], and additive if the FIC is not significantly different from 1.
In all cases, at least three independent experiments were performed (on different days), and within each experiment measurements were averaged from three replicates.

**P. falciparum Drug Accumulation Assays**

The accumulation of $[^3\text{H}]\text{CQ}$ and $[^3\text{H}]\text{AO}$ was measured in mature trophozoite-infected erythrocytes (~36 hours post-invasion). In the experiments giving rise to the data presented in Figure 2, hematocrit and parasitaemia were approximately 1% and 5%, respectively, and the initial extracellular concentration of $[^3\text{H}]\text{CQ}$ was 2 nM (7 Ci/mmol; Moravek Biochemicals). Prior to the addition of $[^3\text{H}]\text{CQ}$, the cell suspension was incubated for 15 minutes at 37°C in the presence or absence of the test compound (unlabelled VP, AO, or QC). The 1-hour incubation was terminated by the addition of 100 μL of dibutyl phthalate to the cell suspension and immediate centrifugation ($17000 \times g$, 1 minute) to sediment the cells. The supernatant solution was aspirated and the microcentrifuge tube tip containing the pelleted cells was cut off and placed in a scintillation vial (PerkinElmer). The cells were lysed with 100 μL of Solvable (Packard; 30 minutes at room temperature) and the sample decolourised by the addition of 100 μL of hydrogen peroxide (Sigma; 30 minutes at room temperature). Scintillation fluid (2 mL of Ultima Gold; PerkinElmer) was added to each vial and the samples were shaken overnight. The radioactivity was measured with a Packard Tri-Carb 4640 liquid scintillation analyser. The data presented in panels A and D of Figure 2 were derived from three independent experiments (performed on different days), within which measurements were averaged from three replicates. Panels B, C, E, and F of Figure 2 show the mean of two independent experiments (performed on different days), within which measurements were averaged from three replicates.

The experiments giving rise to the data presented in Figure 6 and Supplementary Figure 3 were performed according to a protocol described in detail elsewhere [6] with minor modifications. Briefly, the hematocrit was 2% and the parasitaemia ranged between 3.2 and
8.4%, and the initial extracellular concentrations of $[\text{H}]CQ$ (20 Ci/mmol) and $[\text{H}]AO$ (25 Ci/mmol) (both from American Radiolabeled Chemicals, ARC) were 20 nM and 50 nM, respectively. The 1-hour incubation at 37°C was terminated by transferring an aliquot of the suspension (200 µL, in duplicate) to a microcentrifuge tube containing 300 µL of dibutyl phthalate (Sigma-Aldrich), which was centrifuged immediately (17,000 $\times g$, 2 minutes) to sediment the cells. The supernatant solution was removed by aspiration and residual radioactivity on the sides of the tube was removed by rinsing four times with water, after which the majority of the dibutyl phthalate was aspirated. The cell pellet was dissolved in 1 M NaOH (100 µL, 2 hours at 55°C) and the sample decolourised by the addition of 12.5% w/v bleach (25 µL, 2 hours at room temperature) prior to dilution with water (1 mL). The sample was then transferred to a scintillation vial, mixed with scintillation fluid (1.5 mL of Irga-Safe Plus; PerkinElmer), and the radioactivity measured with a PerkinElmer Tri-Carb 2810 liquid scintillation analyser. The data presented in Figure 6 were derived from four independent experiments (performed on different days), within which measurements were averaged from two replicates.

In all cases, the drug accumulation ratio (i.e., the concentration of $[\text{H}]$drug within the infected cells relative to the concentration in the extracellular medium) was calculated using a previous estimate of the water volume of a trophozoite-infected erythrocyte (75 fl.) [7].

Harvest, Preparation and Microinjection of X. laevis Oocytes

Ethical approval of the work performed with the X. laevis frogs was obtained from the Australian National University Animal Experimentation Ethics Committee (Animal Ethics Protocol Number A2013/13) in accordance with the Australian Code of Practice for the Care and Use of Animals for Scientific Purposes. Sections of ovary were surgically removed using a procedure adapted from the Laboratory Xenopus handbook [8]. Adult female X. laevis frogs (purchased from NASCO) were anaesthetised by submersion in a solution of ethyl 3-
aminobenzoate methanesulfonate salt (1.6 g/L of tap water; Sigma-Aldrich) and 1 mM NaHCO$_3$ (Sigma-Aldrich). Anaesthesia was complete after approximately 20 minutes, or when the frog could be turned onto its back without eliciting a response. The frog was then placed onto a container of ice covered with two sheets of moist paper towel. A moist sheet of paper towel was also placed over the frog’s head and another over its legs, leaving the abdomen exposed. Ice was piled onto the frog’s head and legs to prevent bleeding. After the abdomen was swabbed with a cotton bud sprayed with 80% ethanol, a sterile scalpel was used to make a 1 cm incision directly above the position of an ovary – first through the skin and then though the muscle layer. Sections of the ovary were removed with sterile, surgical-grade tweezers and scissors and placed in a pre-weighed petri dish containing ‘calcium-free oocyte ringer’ (OR$_2^-$) buffer (82.5 mM NaCl, 2.5 mM KCl, 1 mM MgCl$_2$, 1 mM Na$_2$HPO$_4$, 5 mM HEPES; pH 7.8). The incision was closed with one stitch of absorbable suture (B. Braun) in the muscle layer, and two stiches of non-absorbable suture (B. Braun) in the skin layer. The wound was swabbed with a cotton bud sprayed with 80% ethanol, after which the frog was returned to an empty tank. The frog then recovered under a sheet of moist paper towel, and was submersed in tap water once fully awake.

The petri dish was re-weighed and the mass of the ovary calculated. The ovary sections were cut into pieces containing around 200 oocytes. If the total mass of the ovary tissue was 17 g or less, the sections were transferred to a 100 mL Erlenmeyer flask, whereas a 250 mL Erlenmeyer flask was used when the total mass was greater than 17 g. The ovary pieces were washed with OR$_2^-$ buffer (five times, or until the buffer was no longer cloudy) before being incubated for 16 hours at 16°C on an orbital shaker in OR$_2^-$ buffer supplemented with 0.5 M Na$_2$HPO$_4$ (Sigma-Aldrich), bovine serum albumin (250 mg/mL, Sigma-Aldrich), and collagenase D (Roche). The volume of the buffer was 25 mL and 50 mL for the 100 mL and 250 mL flasks, respectively. The amount of collagenase D added to the solution was described by the equation $y = 0.44x + 11.5$, where $x$ was the mass of ovary tissue in g and $y$ was the mass
of collagenase in mg. This ratio of collagenase D to ovary mass typically produced good yields of healthy, single, and de-folliculated oocytes when the ovary mass ranged from 11 to 24 g. The collagenase-treated oocytes were washed ten times with OR\(^{-2}\) buffer and five times with ‘calcium-containing oocyte ringer’ (OR\(^{2+}\)) buffer (OR\(^{-2}\) buffer supplemented with 1 mM CaCl\(_2\) and 50 μg/mL gentamycin). An ovary mass of 14-20 g usually yielded 2000-5000 healthy, de-folliculated oocytes of a suitable size and age (developmental stages V and VI) for microinjection with cRNA (20 ng per oocyte) encoding PfCRT. The injections were performed using a Nanoliter 2000 Injector and Micro4 Controller (World Precision Instruments) with fine-tipped glass capillaries that were shaped on a Flaming/Brown micropipette puller (Sutter Instrument Co.). Oocytes were stored at 16-18°C in OR\(^{2+}\) buffer (which was replaced daily) and the solute uptake experiments were performed 3-6 days post-injection.

**Measurements of Drug Transport in Xenopus Oocytes**

The direction of drug transport in the PfCRT oocyte expression system is from the acidic extracellular medium (pH 5.0 to 6.0) into the oocyte cytosol (pH 7.2) [9], which corresponds to the efflux of drug from the acidic digestive vacuole (pH 5 to 5.5) [10-12] into the parasite cytosol (pH 7.3) [13]. A key advantage of this system is that it allows PfCRT to be studied directly and in isolation, without confounding effects such as the binding of drugs to heme or interactions of the compound with other targets or transporters.

Unless specified otherwise, the drug influx assays were performed over 1-2 hours at 27.5°C and in ND96 buffer (96 mM NaCl, 2 mM KCl, 1 mM MgCl\(_2\), 1.8 mM CaCl\(_2\), 10 mM MES, and 10 mM Tris-base). The uptake of \(^{[3]}\text{H}\)CQ (0.25 μM; 20 Ci/mmol; ARC) was measured at pH 6.0 and in the presence of 15 μM unlabelled CQ. A subset of experiments attempted to measure the transport of \(^{[3]}\text{H}\)AO (0.2 μM; 25 Ci/mmol; ARC) into oocytes in the presence of unlabelled AO (1, 2.5, 5, or 10 μM) at pH 4.5, 5.0, 5.5, and 6.0. For each treatment, 10 oocytes were transferred to a 5 mL polystyrene round bottom tube (Falcon) and washed twice with
3.5 mL of ND96 buffer, with the residual buffer removed by pipette. Influx commenced with the addition of 100 µL of ND96 buffer supplemented with [3H]CQ (or [3H]AO), unlabelled CQ (or AO) and, where specified, an unlabelled inhibitor (e.g., QC, AO, MB, or VP). The assay was terminated by removing the reaction buffer with a pipette and washing the oocytes twice with 3.5 mL of ice-cold ND96 buffer. Each oocyte was transferred to a separate scintillation vial and incubated overnight at room temperature in 200 µL of 10% SDS (Sigma-Aldrich). The lysed oocyte was then combined with 1.5 mL of Irga-Safe Plus and the radioactivity measured with a PerkinElmer Tri-Carb 2810 liquid scintillation analyser.

The IC₅₀ values presented in Figures 3B-D were determined in SigmaPlot by a least-squares fit of the equation $y = y_{\text{min}} + \left( \frac{y_{\text{max}} - y_{\text{min}}}{1 + \left( [\text{inhibitor}] / IC_{50} \right) c} \right)$ to the data, where $y$ is PfCRT_Dd2-mediated CQ transport, $y_{\text{min}}$ and $y_{\text{max}}$ are the minimum and maximum values of $y$, and $c$ is a fitted constant. PfCRT_Dd2-mediated CQ transport was calculated by subtracting the uptake measured in oocytes expressing PfCRT_D10 from that in oocytes expressing PfCRT_Dd2. Note that non-injected oocytes and oocytes expressing PfCRT_D10 take up [3H]CQ to similar (low) levels via simple diffusion of the neutral species; this represents the 'background' level of CQ accumulation in oocytes (refer to [9] for full data and a detailed discussion).

The influx of unlabelled QC (10 or 40 µM) and MB (75 µM) was detected using the intrinsic fluorescence of these drugs and the uptake assays were performed at pH 6.0 and pH 5.0, respectively. The composition of the pH 5.0 ND96 buffer was 96 mM NaCl, 2 mM KCl, 1 mM MgCl₂, 1.8 mM CaCl₂, and 20 mM homo-PIPES. A subset of experiments attempted to measure the transport of unlabelled AO (75 or 150 µM) at pH 4.5, 5.0, 5.5, and 6.0. The fluorescence-based experiments were performed as described for the radioisotope uptake assays with the following modifications. At the completion of the incubation, the washed oocytes were transferred to separate wells of a flat 96-well plate (NUNC or PerkinElmer) that had been preloaded with 200 µL of 10% SDS. Each plate included a standard curve that consisted of 10 different concentrations of the fluorescent substrate under study, with each
sample being prepared in a 10% SDS solution containing a lysed oocyte (see Supplementary Figure 2). The plates were incubated overnight at room temperature and mixed on an orbital shaker the following day for approximately 5 minutes. The fluorescence intensity was measured using either a BMG Labtech FLUOstar Optima or a Tecan Infinite M1000 PRO plate-reader. The excitation and emission wavelengths were as follows: QC, 440 and 510 nm; AO, 485 and 520 nm; MB, 668 and 682 nm.

The kinetic parameters for QC and MB transport via PfCRT\textsuperscript{Dd2} (Figures 4D and 5D, respectively) were determined in SigmaPlot by a least-squares fit of the Michaelis-Menten equation \( v = \frac{V_{\text{max}} [\text{substrate}]}{K_m + [\text{substrate}]} \) to the data.

In all cases, at least three independent experiments were performed (on different days and using oocytes from different frogs), and within each experiment measurements were made from 10-12 oocytes per treatment.
Supplementary Table 1. In Vitro Antiplasmodial Activity of Chloroquine (CQ) against 3D7 and K1 *P. falciparum* Parasites in the Absence or Presence of Acridine Orange (AO) or Verapamil (VP).

<table>
<thead>
<tr>
<th>Strain&lt;sup&gt;b&lt;/sup&gt;</th>
<th>IC&lt;sub&gt;50&lt;/sub&gt; (nM)&lt;sup&gt;a&lt;/sup&gt;</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>CQ</td>
</tr>
<tr>
<td>3D7</td>
<td>17 ± 4</td>
</tr>
<tr>
<td>K1</td>
<td>131 ± 11</td>
</tr>
</tbody>
</table>

Abbreviations: IC<sub>50</sub>, half-maximal inhibitory concentration.

<sup>a</sup>The IC<sub>50</sub>s are shown as the mean ± SEM from 3 independent experiments (performed on different days), within which measurements were averaged from 3 replicates.

<sup>b</sup>3D7 is a field-derived CQ-sensitive strain from Africa and K1 is a field-derived CQ-resistant strain from Thailand.

<sup>c</sup>A student *t*-test comparison with the CQ IC<sub>50</sub> determined for the K1 strain in the absence of AO or VP yielded a *P* value less than 0.05.
Supplementary Table 2. Percentages of Chloroquine, Quinacrine, and Acridine Orange in their Neutral, Monoprotonated (H\(^+\)), or Diprotonated (H\(_2\)\(^{2+}\)) Forms in Solutions of Different pH.

<table>
<thead>
<tr>
<th>pH</th>
<th>CQ</th>
<th>QC</th>
<th>AO(^a)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>% CQ</td>
<td>% CQH(^+)</td>
<td>% CQH(_2)(^{2+})</td>
</tr>
<tr>
<td>5.0</td>
<td>0.00000005</td>
<td>0.0799997</td>
<td>99.9199998</td>
</tr>
<tr>
<td>5.5</td>
<td>0.00000050</td>
<td>0.2525445</td>
<td>99.7474505</td>
</tr>
<tr>
<td>6.0</td>
<td>0.0000497</td>
<td>0.7942781</td>
<td>99.2056722</td>
</tr>
<tr>
<td>6.4</td>
<td>0.0003100</td>
<td>1.9714568</td>
<td>98.0282332</td>
</tr>
<tr>
<td>6.9</td>
<td>0.0029731</td>
<td>5.9792417</td>
<td>94.0177852</td>
</tr>
<tr>
<td>7.4</td>
<td>0.0263210</td>
<td>16.7393496</td>
<td>83.2343294</td>
</tr>
<tr>
<td>8.4</td>
<td>1.0392882</td>
<td>66.0955322</td>
<td>32.8651795</td>
</tr>
</tbody>
</table>

The percentages were calculated using the Henderson-Hasselbach equation (pH = pK\(_a\) + log ([A\(^-\)]/[HA])) using the pK\(_a\) values shown in Supplementary Figure 1.

Abbreviations: CQ, chloroquine; CQH\(^+\), monoprotonated chloroquine; CQH\(_2\)\(^{2+}\), diprotonated chloroquine; QC, quinacrine; QCH\(^+\), monoprotonated quinacrine; QCH\(_2\)\(^{2+}\), diprotonated quinacrine; AO, acridine orange; AOH\(^+\), monoprotonated acridine orange.

\(^a\)Note that the concentration of the neutral form of AO at pH 5.0 is 32-fold greater than that of neutral QC at pH 6.0.
Supplementary Figure 1. The structures of the compounds used in this study. Abbreviation: pKₐ, the negative logarithm to the base 10 of the acid dissociation constant (Kₐ).
Supplementary Figure 2. The fluorescence intensities of quinacrine (QC), acridine orange (AO), and methylene blue (MB) are linear with concentration in a 10% SDS solution containing a lysed oocyte. The drug influx assays entail dissolving each oocyte in 200 µL of a 10% SDS solution to release the accumulated drug. Given that the resulting oocyte debris could reduce the level of excitation energy that reaches the compound and/or absorb the emitted fluorescence, it was necessary to prepare each concentration standard in 200 µL of a 10% SDS solution that contained a lysed oocyte. A, The fluorescence intensity of QC, which was measured at an excitation wavelength of 440 nm and an emission wavelength of 510 nm, was linear with concentration between 0.001 and 1 µM. B, The fluorescence intensity of AO, which was measured at an excitation wavelength of 485 nm and an emission wavelength of 520 nm, was linear with concentration between 0.0625 and 2 µM. C, The fluorescence intensity of MB, which was measured at an excitation wavelength of 668 nm and an emission wavelength of 682 nm, was linear with concentration between 0.2 and 100 nM. The data represent the mean ± SEM from 3 to 11 independent experiments, performed on different days and using oocytes from different frogs. The error bars for each of the data points fell within the symbols.
Supplementary Figure 3. The effect of chlorpheniramine (CP) on the accumulation of radiolabelled chloroquine ([3H]CQ) or radiolabelled acridine orange ([3H]AO) by erythrocytes infected with mature trophozoite-stage parasites. The white, black, and gray bars show data for the C2GCO3, C4Dd2, and C67G8 parasite lines, respectively, under control conditions or in the presence of 2.5 µM CP. The accumulation of [3H]drug was expressed as a ‘distribution ratio’ (i.e., the concentration of [3H]drug within the infected cells relative to the concentration in the extracellular medium) and is shown relative to the ratio measured in the C2GCO3 parasites in the absence of CP. The data represent the mean + range/2 of 2 independent experiments (performed on different days), within which measurements were averaged from 2 replicates.
References


