In Vitro Activity of Antiretroviral Drugs against *Plasmodium falciparum*[⊽]

Christian Nsanzabana and Philip J. Rosenthal*

Department of Medicine, San Francisco General Hospital, University of California San Francisco, San Francisco, California

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Malaria and HIV infection are both very common in many developing countries. With the increasing availability of therapy for HIV infection, it was of interest to determine whether antiretroviral drugs exert antimalarial effects. We therefore tested the in vitro activity of 19 antiretroviral drugs against the W2 and 3D7 strains of *Plasmodium falciparum* at concentrations up to 50 µM. None of 5 tested nucleoside reverse transcriptase inhibitors demonstrated activity. Two nonnucleoside reverse transcriptase inhibitors, efavirenz (mean 50% inhibitory concentration [IC₅₀] of 22 to 30 μ M against the two strains) and etravirine (3.1 to 3.4 μ M), were active; nevirapine was not active. Also active were the fusion inhibitor enfuvirtide (6.2 to 7.9 μ M) and the entry inhibitor maraviroc (15 to 21 µM). Raltegravir was not active. However, for all active drugs mentioned above, the IC_{so} s were considerably greater than the concentrations achieved with standard dosing. The effects most likely to be clinically relevant were with HIV protease inhibitors. Of the tested compounds, activity was seen with lopinavir (2.7 to 2.9 µM), atazanavir (3.3 to 13.0 µM), saquinavir (5.0 to 12.1 µM), nelfinavir (6.5 to 12.1 µM), ritonavir (9.5 to 10.9 µM), tipranavir (15.5 to 22.3 µM), and amprenavir (28.1 to 40.8) but not darunavir. Lopinavir was active at levels well below those achieved with standard dosing of coformulated lopinavir-ritonavir. Lopinavir also demonstrated modest synergy with the antimalarial lumefantrine (mean fractional inhibitory concentration index of 0.66 for W2 and 0.53 for 3D7). Prior data showed that lopinavirritonavir also extends the pharmacokinetic exposure of lumefantrine. Thus, when used to treat HIV infection, lopinavir-ritonavir may have clinically relevant antimalarial activity and also enhance the activity of antimalarials.

Every year, 300 to 400 million people become ill and nearly 1 million die from malaria, mostly caused by Plasmodium falciparum and mostly in sub-Saharan Africa (13). HIV currently infects approximately 33 million people worldwide. The pandemic is dynamic, with 2.6 million new infections and nearly 2 million deaths per year (45). With the overlap of malaria and HIV infection in tropical regions, it is important that control measures for one disease consider impacts on the other. Importantly, the treatment of both malaria and HIV infection has changed dramatically in recent years. For malaria, new artemisinin-based combination therapies (ACTs) are now recommended for the treatment of uncomplicated falciparum malaria in nearly all countries where malaria is endemic (52). For HIV infection, antiretroviral therapy is increasingly available in the developing world, and there is a move toward initiating therapy earlier in the course of infection (28). Thus, many patients will be at risk of malaria and also treated for malaria while concomitantly receiving antiretroviral drugs.

The effects of some antiretroviral drugs on malaria have been studied previously. In particular, HIV protease inhibitors have been shown to have antimalarial activity (31, 40). Initially, saquinavir, ritonavir, and indinavir, all older protease inhibitors of relatively little importance for HIV therapy today, were shown to be active against cultured *P. falciparum* (40). Subse-

* Corresponding author. Mailing address: San Francisco General Hospital, 1001 Potrero Avenue, Building 30, Room 421, San Francisco, CA 94110. Phone: (415) 206-8845. Fax: (415) 648-8425. E-mail: prosenthal@medsfgh.ucsf.edu.

quently, a wider range of protease inhibitors was shown to be active, including lopinavir, atazanavir, and amprenavir, drugs that are more widely used than the agents studied earlier (31). Importantly, lopinavir was active at concentrations well below those that circulate in plasma when the drug is coadministered with a second protease inhibitor, ritonavir, to enhance lopinavir levels (31). Subsequent studies showed activity of multiple protease inhibitors against P. falciparum in vitro and Plasmodium chabaudi in vivo (1), activity against clinical isolates of P. falciparum and Plasmodium vivax (22), and in vitro activity of sera from patients receiving protease inhibitors against P. falciparum (36). Considering other classes of antiretroviral drugs, limited data are available. The nonnucleoside reverse transcriptase inhibitor nevirapine was not active (40). The antimalarial activity of most classes of antiretroviral drugs and of some important new antiretroviral protease inhibitors has not been reported previously (41).

Another important consideration concerning antimalarial and antiretroviral therapy is drug interactions leading to changes in pharmacokinetics or antimalarial efficacy. Considering pharmacokinetics, the HIV protease inhibitor ritonavir is widely used to enhance the circulating concentrations of other protease inhibitors, but it may also affect exposure to many other drugs that are metabolized by cytochrome P_{450} 3A4. Indeed, in normal volunteers, exposure to the antimalarial lumefantrine, a component of the ACT artemether-lumefantrine, was markedly enhanced by coadministration of lopinavirritonavir (12), and exposure to a number of other antimalarial drugs may be affected by ritonavir or other agents (41). Considering antimalarial efficacy, HIV protease inhibitors have

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Class ^a	Drug	IC_{50} (μ M) for <i>P</i> . <i>falciparum</i> strain:		Serum concn (µM) ^b					
		W2	3D7	Standard dosing			Ritonavir coadministration		
				C _{min}	$C_{\rm max}$	Reference	$\overline{C_{\min}}$	C_{\max}	Reference
NRTIs	Abacavir	>50	>50	0.1	10.5	26			
	Emtricitabine	>50	>50	0.7	8.3	50			
	Lamivudine	>50	>50	0.3	8.3	7			
	Stavudine	>50	>50	0.1	1.7	17			
	Zidovudine	>50	>50	0.4	4.1	47			
NNRTIs	Efavirenz	21.9	29.7	0.4	33.9	21			
	Etravirine	3.4	3.1	0.4	0.9	4			
	Nevirapine	>50	>50	16.7	26.2	8			
PIs	Amprenavir	28.1	40.8	0.7	9.5	2	2.9	14.3	2
	Atazanavir	3.3	13	0.2	3.6	27	0.7	4.5	49
	Darunavir	>50	>50	0.7	8.7	48	4.1	12.8	4
	Lopinavir	2.7	2.9	5.7	10.4	10	9.5	19.4	20
	Nelfinavir	6.5	12.1	1.3	8.3	43	0.9	6.4	18
	Ritonavir	9.5	10.9	3.6	14.8	38			
	Saquinavir	5	12.1	0.1	1.4	30	0.5	3.7	6
	Tipranavir	15.5	22.3	0.8	22.5	25	32.4	131.3	24
Entry/fusion inhibitor	Enfuvirtide	7.9	6.2	0.1	1.3	44			
	Maraviroc	15.2	21.1	0	0.8	19			
Integrase inhibitor	Raltegravir	>50	>50	0.27	2.71	4			

TABLE 1. Activity of antiretroviral drugs against cultured P. falciparum

^a Abbreviations: NRTI, nucleoside reverse transcriptase inhibitors; NNRTIs, nonnucleoside reverse transcriptase inhibitors; PIs, protease inhibitors.

^b Serum concentrations achieved with standard dosing, including those for protease inhibitors boosted by coadministration with ritonavir, are shown, with the sources of this information cited. Values represent the means of two assays, each performed in duplicate.

been shown to potentiate the activity of chloroquine (14, 15, 23) and to either augment (29) or antagonize (16) the activity of artemisinins. To help clarify the impacts of antiretroviral therapy on malaria, we systematically evaluated the activity of representatives of all currently used classes of antiretroviral drugs against two strains of *P. falciparum*. We also studied the effects of combinations of standard antimalarial drugs with lopinavir, the antiretroviral drug that appears to have the most important antimalarial activity.

MATERIALS AND METHODS

Parasites. The *P. falciparum* strains studied, which were obtained from the Malaria Research and Reference Reagent Resource Center, were 3D7, which is sensitive to all standard antimalarial drugs, and W2, which is resistant to many agents.

Drugs. All antimalarial drugs tested were obtained from Sigma Chemical Co., and all the antiretroviral drugs used in this study were obtained through the NIH AIDS Research and Reference Reagent Program.

In vitro drug susceptibility assay. P. falciparum (W2 and 3D7 strains) was cultured in human erythrocytes at 2% hematocrit in RPMI medium supplemented with 0.5% Albumax, 2 mM L-glutamine, 100 mM hypoxanthine, and 5 μ g/ml gentamicin. The parasites were synchronized by serial treatments with 5% D-sorbitol. Ninety-six-microwell culture plates were prepared with serial dilutions of HIV drugs diluted in dimethyl sulfoxide (DMSO), with concentrations ranging from 50 nM to 50 μ M and negative controls containing the same concentrations of DMSO. Parasites were then cultured for 48 h beginning at the ring stage. After 48 h, cultures were fixed with 4% formaldehyde overnight and then stained with 4 nM YOYO-1 dye (Molecular Probes) in 0.1% Triton X-100 in phosphate-buffered saline. Antimalarial activity was assessed by fluorescence-activated cell sorter analysis as previously described (39). Fifty percent inhibitory concentrations (IC_{s0}s) were calculated by nonlinear regression with the Prism 5.0 program (GraphPad Software).

Drug combination assay. Drug interaction studies were performed using the checkerboard technique as previously described (51). Briefly, parasites were

cultured in wells containing different combinations of antimalarial drugs (0.05 to 10,000 nM) and lopinavir (5 to 10,000 nM). Combinations based on serial dilutions of each drug, each set up in duplicate, were evaluated. Concentrations that yielded 45 to 55% of control parasitemia were plotted on isobolograms and used to calculate fractional inhibitory concentrations (FICs) as the ratio of the IC_{50} of the drug in combination and the IC_{50} of the drug alone (11). The FIC index for two drugs was the FIC of drug A plus the FIC of drug B.

RESULTS

In vitro antimalarial activity. We surveyed in vitro activity against the P. falciparum W2 and 3D7 strains of representatives of all available classes of drugs to treat HIV infection. Nineteen drugs were assessed at concentrations up to 50 µM, including nucleoside reverse transcriptase inhibitors, nonnucleoside reverse transcriptase inhibitors, protease inhibitors, entry/fusion inhibitors, and an integrase inhibitor (Table 1). None of the nucleoside reverse transcriptase inhibitors displayed antimalarial activity. From the nonnucleoside reverse transcriptase inhibitor class, efavirenz showed midmicromolar and etravirine low micromolar antimalarial activity. However, for both active nonnucleoside reverse transcriptase inhibitors, the IC₅₀s were well above the plasma concentrations achieved with standard dosing. HIV protease inhibitors have previously been shown to have antimalarial activity; we broadened previous observations by including a wide range of protease inhibitors, including newer agents not previously studied. Of these agents, lopinavir was most active, and atazanavir, nelfinavir, ritonavir, and saquinavir also had fairly good activity, with IC_{50} s of <10 µM against at least one of the plasmodial strains tested. However, only for lopinavir was the IC₅₀ for both

 TABLE 2. Activities of antimalarial drugs against the strains of

 P. falciparum used in the isobologram analyses

Antimalarial	IC_{50} (nM) for <i>P. falciparum</i> strain:			
	W2	3D7		
Lumefantrine	31.86	42.27		
Quinine	89.50	21.53		
Chloroquine	29.04	9.30		
Monodesethylamodiaquine	9.51	2.70		
Mefloquine	11.22	7.32		
Dihydroartemisinin	1.31	1.23		
Piperaquine	9.98	12.08		
Pyronaridine	0.99	1.24		

strains well below achievable plasma concentrations. In the case of lopinavir, the drug circulates at ~10 to 20 μ M with standard dosing of lopinavir/ritonavir, well above the calculated IC₅₀s of 2.6 to 2.9 μ M for the two tested strains. Tipranavir showed modest activity, with IC₅₀s of 15 to 22 μ M for the two strains. However, those values are below the plasma concentrations achieved when tripranavir is boosted by ritonavir

(33 to 165 μ M). Among newer agents, the two entry/fusion inhibitors had modest activity, although their IC₅₀s were well above typical plasma concentrations. The integrase inhibitor raltegravir did not show antimalarial activity.

Interaction between lopinavir and antimalarials. Consistent with previous studies, our broad survey identified lopinavir as the HIV drug most likely to offer clinically relevant activity against P. falciparum. To add insight into studies of the antimalarial activity of lopinavir, we assessed interactions between lopinavir and antimalarial agents. We used standard methods to test activities of combinations of lopinavir and chloroquine, monodesethylamodiaquine (the active metabolite of amodiaquine), quinine, mefloquine, lumefantrine, piperaquine, pyronaridine, and dihydroartemisinin, based on the IC₅₀s determined for each drug against W2 and 3D7 strain parasites (Table 2). One combination, lopinavir and lumefantrine, showed a trend toward synergism, with the mean FIC index \pm standard deviation being 0.53 ± 0.23 for the 3D7 strain and 0.66 ± 0.32 for W2 (Fig. 1). All other combinations studied showed additive effects, in some cases with modest trends toward synergism or antagonism.

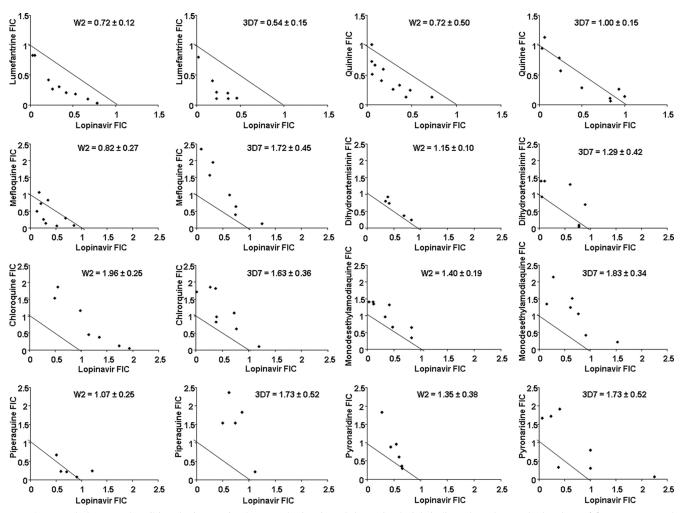


FIG. 1. Isobolograms describing the interaction between lopinavir and the antimalarials indicated on the graphs for the *P. falciparum* W2 and 3D7 strains. Mean fractional inhibitory concentration (FIC) index \pm standard deviation is shown for each combination and each strain.

DISCUSSION

We assessed the antimalarial activity of different HIV drugs. No activity was seen at concentrations up to 50 µM for all 5 nucleoside reverse transcriptase inhibitors tested, the nonnucleoside reverse transcriptase inhibitor nevirapine, the protease inhibitor darunavir, or the integrase inhibitor raltegravir. However, since nucleoside reverse transcriptase inhibitors function as prodrugs (35), we cannot rule out the possibility that the drugs exert antimalarial activity after intracellular phosphorylation. Among nonnucleoside reverse transcriptase inhibitors, efavirenz and etravirine showed antimalarial activity, although their IC_{50} s were well above the levels of the drugs that circulate with standard therapy. The fusion inhibitor enfuvirtide and entry inhibitor maraviroc were also active although, as with the active nonnucleoside reverse transcriptase inhibitors, concentrations with meaningful antimalarial activity are probably not achieved with standard dosing. Consistent with older reports (22, 31, 36, 40), the most meaningful antimalarial activity was seen with HIV protease inhibitors. All tested protease inhibitors except darunavir showed antimalarial activity. Also consistent with older reports, lopinavir was the most potent of these agents. When provided in standard dosing with ritonavir, lopinavir circulates in the plasma at 10 to 19 μ M, well above the IC₅₀s of 2 to 3 μ M determined in this study and 1 to 5 μ M seen in other studies using a variety of methods (1, 31, 32, 33). Serum from patients treated with lopinavir-ritonavir also exerted in vitro antimalarial activity (36). Thus, when administered in modern regimens to treat HIV infection, lopinavir is likely to exert clinically relevant antimalarial activity.

The antimalarial activities of different HIV protease inhibitors vary greatly. Indeed, among newer agents not previously studied, tipranavir had only modest activity, although due to high circulating concentrations of this drug, the activity may be clinically important. Darunavir was inactive at 50 μ M. Thus, lopinavir remains the HIV protease inhibitor with the most promising antimalarial activity. This is a fortuitous result, as lopinavir-ritonavir is increasingly available in countries where malaria is endemic, and it is available in a heat-stable formulation (9).

The antimalarial mechanism of action of HIV protease inhibitors is uncertain. Their action differs from that of the generic aspartic protease inhibitor pepstatin, as unlike pepstatin, they did not display synergy with cysteine protease inhibitors or enhanced activity against a cysteine protease knockout parasite (32). These results suggest that, although lopinavir and ritonavir inhibited the food vacuole hemoglobinase plasmepsin II (32), the HIV protease inhibitors may target other plasmodial proteases. P. falciparum contains 10 aspartic protease genes, which encode 4 food vacuole hemoglobinases (plasmepsins I to IV [3]), a protease that processes proteins for extracellular export (plasmepsin V [5, 37]), and other putative proteases with unknown functions. HIV protease inhibitors probably target plasmepsins, but this has not been confirmed, and it is unclear which plasmepsins are targeted or if all active protease inhibitors target the same enzymes.

We also evaluated the antimalarial activities of combinations of lopinavir and standard antimalarial agents. Interestingly, modest synergy was seen between lopinavir and lumefantrine. Synergy was not seen between lopinavir and any other tested antimalarial. Most combinations appeared to have additive effects. There was a trend toward antagonism between lopinavir and the aminoquinolines chloroquine, amodiaquine (studied as its active metabolite monodesethylamodiaquine), and piperaquine; for piperaquine and mefloquine, a trend toward antagonism was seen only for the chloroquine-resistant strain W2 and not the sensitive strain 3D7. Of interest, prior studies showed potentiation of chloroquine activity by HIV protease inhibitors (14, 15, 23), although the results with lopinavir were modest. It is unclear whether differences in results between our groups are due to the different protease inhibitors studied, differences in methodology, or other factors. Other groups have reported synergy (29) or antagonism (16) between HIV protease inhibitors and artemisinins, although only older protease inhibitors and not lopinavir were studied; we found additive effects between lopinavir and dihydroartemisinin.

Our data and the results of other recent studies suggest that there are three means by which the use of lopinavir-ritonavir may have an impact upon the incidence of malaria. First, the antimalarial activity of lopinavir, with drug levels boosted by ritonavir, may kill erythrocytic parasites before infections progress to clinical illness. Second, after therapy for a prior infection, inhibition of cytochrome P450 3A4 by ritonavir may extend exposure to antimalarial drugs, as has been demonstrated for lumefantrine (12), thereby prolonging the period during which a drug circulates at concentrations adequate to prevent new infections. Third, through synergistic effects, cocirculating antimalarials and antiretrovirals may prevent new infections more effectively than the antimalarials alone. For the last two described mechanisms, the evidence best supports an effect of lopinavir-ritonavir on the posttreatment prophylactic activity of lumefantrine. Therefore, it is of interest to evaluate the effect of lopinavir-ritonavir on those at high risk of malaria and on those treated for malaria in high-transmission areas with artemether-lumefantrine. Indeed, studies of the effects on malaria of treatment of HIV with lopinavir-ritonavir are under way. More broadly, as the use of antiretroviral therapy is increasing in countries where malaria is endemic, our results highlight the importance of evaluating the antimalarial effects of antiretroviral drugs, both in laboratory settings and in clinical trials.

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