

# Comparative *Ex Vivo* Activity of Novel Endoperoxides in Multidrug-Resistant *Plasmodium falciparum* and *P. vivax*

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**The declining efficacy of artemisinin derivatives against *Plasmodium falciparum* highlights the urgent need to identify alternative highly potent compounds for the treatment of malaria. In Papua Indonesia, where multidrug resistance has been documented against both *P. falciparum* and *P. vivax* malaria, comparative *ex vivo* antimalarial activity against *Plasmodium* isolates was assessed for the artemisinin derivatives artesunate (AS) and dihydroartemisinin (DHA), the synthetic peroxides OZ277 and OZ439, the semisynthetic 10-alkylaminoartemisinin derivatives artemisone and artemiside, and the conventional antimalarial drugs chloroquine (CQ), amodiaquine (AQ), and piperaquine (PIP). *Ex vivo* drug susceptibility was assessed in 46 field isolates (25 *P. falciparum* and 21 *P. vivax*). The novel endoperoxide compounds exhibited potent *ex vivo* activity against both species, but significant differences in intrinsic activity were observed. Compared to AS and its active metabolite DHA, all the novel compounds showed lower or equal 50% inhibitory concentrations (IC<sub>50</sub>s) in both species (median IC<sub>50</sub>s between 1.9 and 3.6 nM in *P. falciparum* and 0.7 and 4.6 nM in *P. vivax*). The antiplasmodial activity of novel endoperoxides showed different cross-susceptibility patterns in the two *Plasmodium* species: whereas their *ex vivo* activity correlated positively with CQ, PIP, AS, and DHA in *P. falciparum*, the same was not apparent in *P. vivax*. The current study demonstrates for the first time potent activity of novel endoperoxides against drug-resistant *P. vivax*. The high activity against drug-resistant strains of both *Plasmodium* species confirms these compounds to be promising candidates for future artemisinin-based combination therapy (ACT) regimens in regions of coendemicity.**

Approximately 3.3 billion people (i.e., almost 50% of the world's population) are at risk of malaria with two *Plasmodium* species responsible for the majority of infections: *P. falciparum* and *P. vivax* (6, 7, 43). Traditionally, malaria control and research efforts have focused on *P. falciparum*, the dominant species in Africa. However, outside Africa, *P. falciparum* almost invariably coexists with *P. vivax* (7), with both species inflicting significant morbidity, particularly in infants and pregnant women (18, 27).

Chloroquine (CQ)-resistant *P. falciparum* is already well established, with emerging evidence that susceptibility to CQ in *P. vivax* is also declining across much of the world in which *vivax* is endemic. This combined threat is driving the investigation of alternative schizonticidal treatment regimens, such as artemisinin-based combination therapy (ACT), for deployment against both *P. falciparum* and *P. vivax* (29). ACTs have become the mainstay of antimalarial chemotherapy, adopted in more than 100 countries worldwide (42). This huge demand for artemisinin and its derivatives relies on isolation from the plant source *Artemisia annua* and is vulnerable to harvest and production costs and intermittent supply (2, 11). Of particular concern are recent reports of prolonged *in vivo* parasite clearance times following ACTs on the Thai-Cambodian border (4, 21), highlighting the possibility of emerging artemisinin resistance. These constraints have driven new efforts to develop novel semisynthetic artemisinins or fully synthetic peroxides, some of which are currently in phase I and II clinical trials. The fully synthetic peroxides include the 1,2,4-trioxolanes OZ277 and OZ439 (also known as ozonides) (3), and the

semisynthetic artemisinins include the 10-alkylamino-artemisinin derivative artemisone and its thiomorpholine precursor, artemiside (8). OZ277 (also known as RBx11160 or arterolane) was the first ozonide to enter clinical trials, is currently evaluated in combination with piperaquine (PIP) (25), and has obtained approval for use in India. The novel ozonide OZ439 has a slower elimination than OZ277 and is able to cure *Plasmodium berghei* infections in mice with a single oral administration (3). Artemisone in combination with mefloquine, amodiaquine, or clindamycin cures *P. falciparum* infections in *Aotus* monkeys (24) and is effective against murine cerebral malaria (38). Preliminary phase II data demonstrated that artemisone is curative at one-third the dose level of artesunate (AS) when used in patients with nonsevere malaria (19).

Outside Africa, the high levels of malaria coendemicity and the emergence of multidrug resistance in both species have led to calls for a unified policy of ACT for both *P. falciparum* and *P. vivax* (5), an approach now endorsed by Indonesia, Papua New Guinea, the Solomon Islands, and Vanuatu. The success of such a policy is

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**TABLE 1** Baseline characteristics of isolates for which the *ex vivo* assay was accomplished

Baseline characteristic	<i>P. falciparum</i>	<i>P. vivax</i>
No. of isolates reaching harvest/total no. of isolates (%)	23/25 (92)	18/21 (90)
Median (range) delay from venipuncture to start of culture (min)	140 (80–246)	102.5 (77–180)
Median (range) duration of assay (h)	46 (32–56)	47 (41–48)
Geometric mean (95% CI <sup>a</sup> ) parasitemia (asexual parasites/ $\mu$ l)	17,291 (11,212–26,666)	11,110 (6,925–17,823)
Median initial % (range) of parasites at ring stage	100 (100–100)	85 (59–97)
Mean (95% CI) schizont count at harvest	36 (16–80)	34 (20–50)

<sup>a</sup> CI, confidence interval.

dependent upon confirmation of antimalarial activity of established and novel endoperoxide drugs against clinical isolates of both *P. falciparum* and *P. vivax*, particularly in areas of known multidrug resistance. The aim of the current study was to investigate the *ex vivo* drug susceptibility profiles of novel endoperoxide agents and compare these with those of established antimalarials.

**MATERIALS AND METHODS**

**Compounds.** The antimalarial drugs chloroquine (CQ), amodiaquine (AQ; Sigma-Aldrich, Australia), piperazine (PIP; Ranbaxy Lab. Ltd., Gurgaon, India), artesunate (AS; Guilin Pharmaceutical Co., Ltd., Guangxi, China), and dihydroartemisinin (DHA; Chongqing Holley Wuling Mountain Pharmaceutical Co., Ltd., China) and the experimental compounds OZ277, OZ439 (MMV), artemisone, and artemiside (The Hong Kong University of Science and Technology) were prepared as 10 mM stock solutions in dimethyl sulfoxide (DMSO). Drug plates were then pre-dosed by diluting the compounds in 50% ethanol, followed by lyophilization, and stored at 4°C. Drug plates were quality controlled by assessing drug response profiles in the CQ-resistant and -sensitive laboratory strains K1 and FC27, respectively, before and after completion of the study.

**Field location and sample collection.** *Plasmodium* species isolates were collected from patients with malaria attending outpatient clinics in Timika, Papua Province, Indonesia, a region where multidrug-resistant strains of *P. vivax* and *P. falciparum* are highly prevalent (10, 30, 31). Patients with symptomatic malaria were recruited into the study if they

were singly infected with *P. falciparum* or *P. vivax* and had a parasitemia of between 2,000  $\mu$ l<sup>-1</sup> and 80,000  $\mu$ l<sup>-1</sup>. After written informed consent was obtained, venous blood (5 ml) was collected by venipuncture. Host white blood cells (WBC) were removed by using CF 11 cellulose, and packed infected red blood cells (IRBC) were used for the *ex vivo* drug susceptibility assay.

**Ex vivo drug susceptibility assay.** *Plasmodium* drug susceptibility was measured using a protocol modified from the WHO microtest as described previously (17, 31). Two hundred microliters of a 2% hematocrit blood medium mixture (BMM), consisting of RPMI 1640 medium plus 10% AB<sup>+</sup> human serum (*P. falciparum*) or McCoy’s 5A medium plus 20% AB<sup>+</sup> human serum (*P. vivax*), was added to each well of pre-dosed drug plates containing 11 serial concentrations (2-fold dilutions) of the antimalarials (maximum concentration shown in parentheses) CQ (2,992 nM), AQ (80 nM), PIP (769 nM), AS (25 nM), DHA (34 nM), OZ277 (34 nM), OZ439 (34 nM), artemisone (24 nM), and artemiside (26 nM). A candle jar was used to mature the parasites at 37.0°C for 32 to 56 h. Incubation was stopped when >40% of ring stage parasites had reached the mature schizont stage (i.e.,  $\geq$ 4 distinct nuclei per parasite) in the drug-free control well.

Thick blood films made from each well were stained with 5% Giemsa solution for 30 min and examined microscopically. The number of schizonts per 200 asexual stage parasites was determined for each drug concentration and normalized to that of the control well. The dose-response data were analyzed using nonlinear regression analysis (WinNonLn 4.1; Pharsight Corporation), and the 50% inhibitory concentration (IC<sub>50</sub>) was derived using an inhibitory sigmoid maximum effect (*E*<sub>max</sub>) model. IC<sub>50</sub>

**TABLE 2** Overall *ex vivo* sensitivity for each drug according to the species tested

Drug	Indicated values for each group of isolates					
	Median IC <sub>50</sub> (nM) in each <i>P. falciparum</i> lab line <sup>a</sup>		<i>P. falciparum</i> clinical field isolates		<i>P. vivax</i> clinical field isolates	
	FC27 (CQ <sup>s</sup> )	K1 (CQ <sup>r</sup> )	n (%) <sup>c</sup>	Median IC <sub>50</sub> (range) in nM and <i>P</i> value <sup>b</sup>	n (%) <sup>c</sup>	Median IC <sub>50</sub> (range) in nM and <i>P</i> value <sup>b</sup>
Chloroquine	39.9	168.7	23 (100)	91.4 (37.3–147.7)	18 (100) <sup>d</sup>	48.3 (12.9–143.4)
Amodiaquine	36.2	21.7	23 (100)	11.8 (5.6–22.8)	18 (100)	15.9 (7.1–25.8)
Piperazine	64.2	50.2	23 (100)	16.4 (4.0–43.6)	18 (100)	14.6 (4.0–28.9)
Artesunate	15.4	10.3	22 (96) <sup>e</sup>	4.5 (0.3–9.8)	15 (83) <sup>e</sup>	4.0 (1.6–6.3)
DHA	10.0	9.6	22 (96) <sup>f</sup>	6.4 (2.2–11.1), <i>P</i> = 0.013	16 (89) <sup>e</sup>	4.9 (2.8–11.5), <i>P</i> = 0.013
OZ277	10.9	20.0	23 (100)	3.6 (0.1–18.3), <i>P</i> = 0.884	16 (89) <sup>e</sup>	4.6 (1.7–16.3), <i>P</i> = 0.088
OZ439	11.1	18.6	23 (100)	2.1 (0.4–10.5), <i>P</i> = 0.012	16 (89) <sup>e</sup>	4.5 (1.3–10.3), <i>P</i> = 0.306
Artemisone	1.9	2.0	22 (96) <sup>e</sup>	1.9 (0.1–5.7), <i>P</i> < 0.001	15 (83) <sup>e</sup>	0.7 (0.3–4.1), <i>P</i> < 0.001
Artemiside	4.0	4.3	22 (96) <sup>e</sup>	2.5 (0.2–15.0), <i>P</i> = 0.592	15 (83) <sup>e</sup>	3.0 (0.5–14.4), <i>P</i> = 0.460

<sup>a</sup> Mean IC<sub>50</sub>s (derived from 2 independent experiments) were assessed by *in vitro* schizont maturation quantified by microscopy. CQ<sup>s</sup>, chloroquine-sensitive laboratory strain; CQ<sup>r</sup>, chloroquine-resistant laboratory strain.

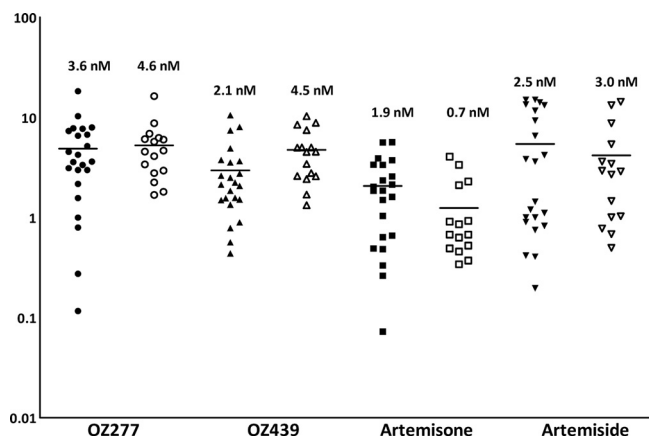
<sup>b</sup> *P* values are from a comparison with values for artesunate (Wilcoxon signed-rank test).

<sup>c</sup> Total number of assays with acceptable IC<sub>50</sub>s (percentage of samples which fulfilled the criteria for a successful culture).

<sup>d</sup> One *P. vivax* isolate was excluded from analysis (harvested at <30 h).

<sup>e</sup> Insufficient BMM mixture to test all compounds in 3 *P. vivax* isolates and 1 *P. falciparum* isolate.

<sup>f</sup> No IC<sub>50</sub> estimate of DHA for 1 *P. falciparum* isolate (highly sensitive, model not possible).



**FIG 1** *Ex vivo* drug susceptibility (median  $IC_{50}$ s) of the synthetic ozonides OZ277 and OZ439 and the semisynthetic endoperoxides artemisone and artemiside in *P. falciparum* (filled symbols) and *P. vivax* (open symbols) clinical field isolates.

*ex vivo* data were used from predicted curves only where the  $E_{max}$  and  $E_0$  were within 15% of 100 and 0, respectively.

**Data analysis.** Analysis was performed using STATA software (version 10.1; Stata Corp., College Station, TX). The Mann-Whitney U test, Wilcoxon signed-rank test, and Spearman rank correlation were used for nonparametric comparisons and correlations. Since previous studies have highlighted several potential confounders of *ex vivo/in vitro* assessment of drug susceptibility (13, 31), we used robust multiple regression analysis on log-transformed data to control for the effect of potential confounders on *ex vivo* drug response, including the initial parasitemia, the proportion of parasites at ring stage at the start of the assay, and the duration of the assay.

**Ethical approval.** Ethical approval for this project was obtained from the Human Research Ethics Committee of the Northern Territory Department of Health & Families and Menzies School of Health Research (HREC 2010-1396), Darwin, Australia, and the Eijkman Institute Research Ethics Commission (EIREC-47), Jakarta, Indonesia, in September 2010.

## RESULTS

**Antimalarial susceptibility.** *Ex vivo* drug susceptibility was assessed in field isolates from 46 patients presenting with single-species infections of *P. falciparum* ( $n = 25$ ) or *P. vivax* ( $n = 21$ ). Adequate growth for harvest was achieved in 92% (23/25) of *P.*

*falciparum* and 90% (19/21) of *P. vivax* isolates. Baseline characteristics of the isolates processed are presented in Table 1. Median  $IC_{50}$ s for all isolates which fulfilled criteria for successful culture for the two species and for the *P. falciparum* laboratory strains FC27 and K1 are depicted in Table 2.

Drug susceptibility to CQ was significantly lower in *P. falciparum* isolates (median  $IC_{50}$ , 91.4 nM; range, 37.3 to 147.7 nM) than in *P. vivax* isolates (median, 48.3 nM; range, 12.9 to 143.4 nM;  $P < 0.001$ ) (Fig. 1), whereas OZ439 exhibited higher  $IC_{50}$ s for *P. vivax* (4.5 nM versus 2.1 nM;  $P = 0.013$ ) (Fig. 1). Previous studies have highlighted a difference in the stage-specific activities of CQ and AQ in *P. vivax*. Stratified analysis of *P. vivax* isolates with  $\geq 80\%$  and 60 to 80% of parasites at ring stage at the start of the assay showed a slight but significant difference in activity for AS (median  $IC_{50}$  of 3.5 nM in isolates with  $\geq 80\%$  rings and 4.9 nM in isolates with 60 to 80% rings;  $P = 0.037$ ) and DHA (median  $IC_{50}$  of 4.8 nM in isolates with  $\geq 80\%$  rings and 6.0 nM in isolates with 60 to 80% rings;  $P = 0.051$ ) but no stage-specific difference in *ex vivo* susceptibility to the other compounds (Table 3).

A within-species comparison with AS revealed significantly lower  $IC_{50}$ s for artemisone in both species (median AS  $IC_{50}$  of 4.5 nM versus a median artemisone  $IC_{50}$  of 1.9 nM [ $P < 0.001$ ] for *P. falciparum* and 4.0 nM versus 0.7 nM, respectively [ $P < 0.001$ ], for *P. vivax*). The median  $IC_{50}$  for OZ439 was also significantly lower than that for AS, but this was apparent only in *P. falciparum* isolates (median, 2.1 nM;  $P = 0.012$ ).  $IC_{50}$ s for DHA were significantly higher than those for AS in both species (median DHA  $IC_{50}$  of 6.4 nM [ $P = 0.013$ ] for *P. falciparum* and 4.9 nM [ $P = 0.013$ ] for *P. vivax*) (Table 2). The trends in drug susceptibility compared to the AS reference drug remained after controlling for the initial stage composition of the parasite at the start of the assay.

**Cross-susceptibility patterns.** Cross-susceptibility patterns were assessed using the Spearman correlation analysis (Table 4). The initial parasitemia was identified as a weak confounding factor for the *ex vivo* drug response to OZ277 in both species (*P. falciparum*,  $r_s = 0.568$ ,  $P = 0.021$ ; *P. vivax*,  $r_s = 0.394$ ,  $P = 0.036$ ) and artemisone and artemiside in *P. vivax* ( $r_s = 0.617$ ,  $P = 0.005$ , and  $r_s = 0.481$ ,  $P = 0.035$ , respectively), whereas this was not apparent for either the duration of assay or the initial parasite stage. In multivariate analysis, correlation patterns did not change substantially after controlling for these confounding

**TABLE 3** Overall *ex vivo* sensitivity for *P. vivax* with  $\geq 80\%$  and  $< 80\%$  of parasites at ring stage at the start of the assay

Drug	Indicated values for <i>P. vivax</i> isolates with:			
	$\geq 80\%$ rings		60–80% rings	
	$n$ (%) <sup>a</sup>	Median $IC_{50}$ (range) in nM and $P$ value <sup>b</sup>	$n$ (%) <sup>a</sup>	Median $IC_{50}$ (range) in nM and $P$ value <sup>b</sup>
Chloroquine	11/11 (100)	45.7 (12.9–143.4)	7/7 (100)	50.8 (23.6–68.6)
Amodiaquine	11/11 (100)	17.1 (7.1–25.8)	7/7 (100)	13.4 (7.5–17.0)
Piperaquine	11/11 (100)	13.1 (4.0–28.7)	7/7 (100)	21.9 (5.2–28.9)
Artesunate	10/11 (91)	3.5 (1.6–6.2)	5/7 (71)	4.9 (3.6–6.3)
DHA	10/11 (91)	4.8 (2.8–6.8), $P = 0.029$	6/7 (86)	6.0 (4.3–11.5), $P = 0.225$
OZ277	10/11 (91)	5.2 (1.7–8.7), $P = 0.037$	6/7 (86)	4.0 (1.8–16.3), $P = 0.500$
OZ439	10/11 (91)	3.7 (1.3–8.8), $P = 0.285$	6/7 (86)	4.8 (2.6–10.3), $P = 0.892$
Artemisone	10/11 (91)	0.6 (0.3–3.4), $P = 0.005$	5/7 (71)	0.9 (0.6–4.1), $P = 0.043$
Artemiside	10/11 (91)	2.8 (0.7–14.4), $P = 0.386$	5/7 (71)	2.9 (0.5–13.3), $P = 0.893$

<sup>a</sup> Total number of assays with acceptable  $IC_{50}$ s/total number of assays (percentage).

<sup>b</sup>  $P$  values are from a comparison with values for artesunate (Wilcoxon signed-rank test).

**TABLE 4** Correlation coefficients for *ex vivo* antimalarial susceptibilities in *P. falciparum* and *P. vivax* field isolates

Novel drug and potential confounder or compared antimalarial drug	Correlation coefficient and <i>P</i> value for field isolates of <sup>a</sup> :			
	<i>P. falciparum</i>		<i>P. vivax</i>	
	Coefficient	<i>P</i> value	Coefficient	<i>P</i> value
<b>OZ277</b>				
Parasitemia	<b>0.568</b>	<b>0.021</b>	<b>0.394</b>	<b>0.036</b>
Duration of assay	-0.041	0.667	-0.080	0.537
Rings at start of the assay	NA	NA	0.053	0.641
Chloroquine	<b>0.736</b>	<b>0.032</b>	-0.050	0.176
Amodiaquine	0.397	0.181	-0.059	0.546
Piperaquine	<b>0.718</b>	<b>&lt;0.001</b>	-0.027	0.057
Artesunate	<b>0.291</b>	<b>0.002</b>	0.282	0.389
DHA	<b>0.581</b>	<b>0.014</b>	0.424	0.172
OZ439	<b>0.853</b>	<b>&lt;0.001</b>	0.456	0.325
Artemisone	<b>0.767</b>	<b>&lt;0.001</b>	<b>0.693</b>	<b>0.001</b>
Artemiside	<b>0.828</b>	<b>&lt;0.001</b>	<b>0.611</b>	<b>0.059</b>
<b>OZ439</b>				
Parasitemia	0.424	0.104	0.082	0.487
Duration of assay	-0.193	0.290	-0.020	0.854
Rings at start of the assay	NA	NA	-0.117	0.494
Chloroquine	<b>0.658</b>	<b>0.018</b>	<b>0.427</b>	<b>0.014</b>
Amodiaquine	<b>0.563</b>	<b>&lt;0.001</b>	0.100	0.145
Piperaquine	<b>0.653</b>	<b>0.004</b>	0.077	0.809
Artesunate	<b>0.486</b>	<b>0.019</b>	0.629	0.091
DHA	<b>0.665</b>	<b>0.008</b>	0.353	0.054
Artemisone	<b>0.693</b>	<b>&lt;0.001</b>	<b>0.550</b>	<b>0.031</b>
Artemiside	<b>0.809</b>	<b>&lt;0.001</b>	<b>0.500</b>	<b>0.052</b>
<b>Artemisone</b>				
Parasitemia	0.353	0.069	<b>0.617</b>	<b>0.005</b>
Duration of assay	-0.258	0.177	0.078	0.725
Rings at start of the assay	NA	NA	-0.268	0.504
Chloroquine	<b>0.682</b>	<b>0.016</b>	0.327	0.432
Amodiaquine	0.319	0.681	<b>0.325</b>	<b>0.010</b>
Piperaquine	<b>0.747</b>	<b>&lt;0.001</b>	0.600	0.528
Artesunate	<b>0.501</b>	<b>&lt;0.001</b>	0.593	0.452
DHA	<b>0.573</b>	<b>&lt;0.001</b>	0.621	0.273
Artemiside	0.461	0.308	<b>0.454</b>	<b>&lt;0.001</b>
<b>Artemiside</b>				
Parasitemia	0.393	0.133	<b>0.481</b>	<b>0.035</b>
Duration of assay	0.143	0.388	0.195	0.496
Rings at start of the assay	NA	NA	0.156	0.596
Chloroquine	<b>0.660</b>	<b>0.010</b>	-0.021	0.849
Amodiaquine	0.509	0.060	-0.157	0.216
Piperaquine	<b>0.565</b>	<b>0.038</b>	0.111	0.781
Artesunate	0.273	0.143	<b>0.489</b>	<b>0.028</b>
DHA	<b>0.570</b>	<b>0.055</b>	<b>0.711</b>	<b>0.012</b>

<sup>a</sup> Correlation coefficients were obtained by the Spearman rank correlation. *P* values were obtained by robust multiple regression analysis. Bold values indicate significant correlations. NA, not applicable.

factors (Table 4), but there was a difference in correlation patterns between the two *Plasmodium* species. In *P. falciparum* isolates, novel endoperoxide drug responses correlated significantly with those of AS, DHA, CQ, and PIP. While *ex vivo* drug response was correlated with the novel endoperoxides (ozonides and artemisone and artemiside) in *P. vivax* isolates, there was no correlation between these compounds and AS, DHA

(the exception being artemiside), CQ, or PIP in this species (Table 4).

**DISCUSSION**

The synthetic peroxides OZ277 and OZ439 and the semisynthetic 10-alkylamino-artemisinin derivatives artemisone and artemiside demonstrate potent *ex vivo* activity against both *Plasmodium* species. All of the novel compounds tested in our study showed IC<sub>50</sub>s lower than or equal to those of AS and DHA, with IC<sub>50</sub>s varying between 1.9 and 3.6 nM against *P. falciparum* and 0.7 and 4.6 nM in *P. vivax* (Table 2).

Significant interspecies differences in *ex vivo* drug responses were observed for CQ (the median CQ IC<sub>50</sub> is greater in *P. falciparum* than *P. vivax*) and OZ439 (the median OZ439 IC<sub>50</sub> is greater in *P. vivax* than *P. falciparum*). The differences observed in CQ IC<sub>50</sub>s may reflect the spectrum of acquired resistance for each species prevalent within the community, but this is unlikely to explain the difference for OZ439. Although intrinsic species variation in life cycle maturation and drug susceptibility inherent to the assay cannot be ruled out, the number of isolates assessed was relatively small, with a borderline level of significance for the OZ439 comparison (*P* = 0.013), and thus the result may represent a chance occurrence within the context of multiple comparisons.

A comparative analysis in *P. vivax* between isolates with >80% rings and 60 to 80% rings at the start of the assay revealed a significant, albeit modest, difference in IC<sub>50</sub>s for AS and DHA, with isolates containing 80% rings at the start of the assay appearing more susceptible. Stage specificity of drug action in *P. vivax* has been highlighted in previous studies for CQ (31, 33) in which *ex vivo* susceptibility increased by more than 20-fold for isolates predominantly at the trophozoite stage compared to the ring stage. The variation in drug susceptibility between parasites at the ring and trophozoite stages was considerably less for AS and DHA (1.5-fold) and was not apparent for any of the novel endoperoxide compounds (Table 2).

The IC<sub>50</sub> values for DHA were significantly higher than those for artesunate and, to various degrees, the other endoperoxides. This may represent either reduced stability of DHA in predosed drug plates (35) or differences in *in vitro* drug partition and metabolism (14, 16, 37). High variability and fluctuations in IC<sub>50</sub>s for artemisinin and its derivatives have been reported for both *P. falciparum* laboratory strains and field isolates (15, 34, 41) and reflect, at least in part, the application of nonstandardized *in vitro* assay systems, including differences in levels of parasitemia at the start of the assay, parasite synchrony, duration of the assay, and quantification method of the drug response (9, 12, 13). These variations were also apparent in the IC<sub>50</sub> estimates derived from laboratory-adapted *P. falciparum* strains in the current study, for which drug susceptibility was assessed by the schizont maturation test as opposed to a reinvasion assay. The median IC<sub>50</sub>s ranged from 1.9 to 15.4 nM and 2.0 to 20.0 nM in the CQ-sensitive and CQ-resistant *P. falciparum* culture-adapted strains FC27 and K1, respectively (Table 2).

The schizont maturation test has been used to quantify *P. falciparum* drug susceptibility for more than 30 years, although alternative, more robust methods are now widely used (1a). However, for *P. vivax*, the assay options remain limited, and in view of the marked stage specificity of drug action, microscopic quantification of parasite growth remains the preferred option (12a, 17, 31). Previous studies have demonstrated that IC<sub>50</sub> estimates de-

rived from schizont maturation tend to be higher than those derived by *in vitro* reinvasion assays with growth quantified by <sup>3</sup>H-hypoxanthine incorporation (1, 23, 36, 39).

Recent clinical, molecular, and *in vitro* studies have raised the specter of emerging drug resistance to the artemisinin derivatives, the focus of which appears to be on the western border of Cambodia. The hallmark of artemisinin tolerance is a delayed parasite clearance rate *in vivo*. However, the relationship between this delayed parasitological response and *in vitro* artemisinin drug susceptibility using established *in vitro* assays is unclear (4, 21, 22). This lack of *in vivo*-*in vitro* correlates has undermined the definition of artemisinin resistance and the *ex vivo* characterization of this process (20, 28, 40). As fears that artemisinin tolerance has already spread beyond Cambodia grow, there is an urgent need to develop *ex vivo* assay systems to characterize better the phenotypic response of *Plasmodium* to current and novel artemisinin derivatives. These will need to address stage-specific quantification of *in vitro*/*ex vivo* drug response and potentially the detection and quantification of dormant parasite stages.

Our study is one of the first to assess parasite susceptibility of a range of endoperoxide compounds, quantifying stage-specific growth between two *Plasmodium* species. In *P. falciparum*, the activity of the novel endoperoxides correlated with that of AS, DHA, CQ, and PIP, a phenomenon previously observed in studies of field isolates (2a, 14, 26, 31). The same trend was observed in *P. vivax* but did not reach statistical significance. Although this may simply reflect the small number of isolates tested and, thus, the low statistical power to detect a true difference, it may also represent species-specific differences in drug uptake and metabolism as well as the mode of action of these compounds.

The novel endoperoxides have a number of advantages, including better pharmacodynamic profiles, pharmacokinetic profiles amenable to single-dosing strategies, and feasibility of synthesis, over currently used artemisinin derivatives. Our study highlights their potent *ex vivo* activity against clinical field isolates of both *P. falciparum* and *P. vivax* in an area of known multidrug resistance. Together, these factors demonstrate that these novel compounds represent promising candidates for future ACT combination regimens that will have activity in areas of coendemicity. Development of *ex vivo* assays that robustly quantify *ex vivo*/*in vitro* growth dynamics will help to characterize the stage-specific activity of these compounds and facilitate assessment of their activity against clinical *Plasmodium* field isolates in areas in which the clinical efficacy of artemisinin is declining.

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