

Structural Optimization of Quinolon-4(1*H*)-imines as Dual-Stage Antimalarials: Toward Increased Potency and Metabolic Stability

Ana S. Ressurreição,^{*,†,∞} Daniel Gonçalves,^{†,∞} Ana R. Siteo,[†] Inês S. Albuquerque,[‡] Jiri Gut,^{||} Ana Góis,[‡] Lídia M. Gonçalves,[†] Maria R. Bronze,[†] Thomas Hanscheid,[‡] Giancarlo A. Biagini,[⊥] Philip J. Rosenthal,^{||} Miguel Prudêncio,[‡] Paul O'Neill,[#] Maria M. Mota,[‡] Francisca Lopes,[†] and Rui Moreira[†]

[†]Research Institute for Medicines and Pharmaceutical Sciences (iMed.UL), Faculty of Pharmacy, University of Lisbon, Av. Prof. Gama Pinto, 1649-019 Lisbon, Portugal

[‡]Instituto de Medicina Molecular, Faculty of Medicine, University of Lisbon, Av. Prof. Egas Moniz, 1649-028 Lisbon, Portugal

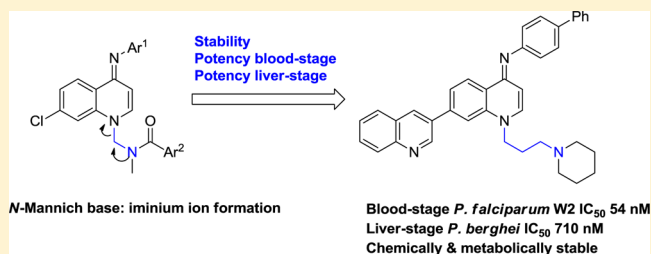
^{||}Department of Medicine, San Francisco General Hospital, University of California, Box 0811, San Francisco, California 94143, United States

[⊥]Liverpool School of Tropical Medicine, Pembroke Place, Liverpool, L3 5QA, U.K.

[#]Department of Chemistry, University of Liverpool, Liverpool, L69 3BX, U.K.

S Supporting Information

ABSTRACT: Discovery of novel effective and safe antimalarials has been traditionally focused on targeting erythrocytic parasite stages that cause clinical symptoms. However, elimination of malaria parasites from the human population will be facilitated by intervention at different life-cycle stages of the parasite, including the obligatory developmental phase in the liver, which precedes the erythrocytic stage. We have previously reported that *N*-Mannich-based quinolon-4(1*H*)-imines are potent antiplasmodial agents but present several stability liabilities. We now report our efforts to optimize quinolon-4(1*H*)-imines as dual-stage antiplasmodial agents



endowed with chemical and metabolic stability. We report compounds active against both the erythrocytic and exoerythrocytic forms of malaria parasites, such as the quinolon-4(1*H*)-imine **5p** (IC₅₀ values of 54 and 710 nM against the erythrocytic and exoerythrocytic forms), which constitute excellent starting points for further lead optimization as dual-stage antimalarials.

■ INTRODUCTION

Malaria remains a major global public health problem, affecting over 200 million people mainly in tropical and subtropical regions and with an estimated death toll of nearly 800 000 individuals in 2011.^{1,2} Five species of the genus *Plasmodium* cause infection in humans. Of these, *P. falciparum* and *P. vivax* account for more than 95% of malaria cases, with *P. falciparum* responsible for most of the deaths caused by malaria every year. The rapid evolution and spread of parasite resistance to most available antimalarial drugs threaten the use of these agents to treat and prevent malaria.^{3,4} To address the need for new and safe drugs for treating or preventing malaria, large phenotypic high-throughput screening campaigns^{5,6} have been performed to discover novel chemotypes acting against erythrocytic stage parasites, which are responsible for the clinical features of malaria. However, global control and eventual eradication of malaria will benefit from action against multiple parasite life-cycle stages.^{7,8}

Malaria parasites undergo an asymptomatic, obligatory developmental phase in the liver, which precedes the formation of the erythrocytic stage.⁹ In addition, *P. vivax* and *P. ovale* infections also generate liver forms, called hypnozoites, that are

not eliminated by standard therapy, persist in the liver for long periods, and upon reactivation, are responsible for relapses of malaria.¹⁰ Thus, the liver stage of infection offers an important target for intervention in addition to erythrocytic stages. Indeed, eradication of liver-stage parasites can provide causal prophylaxis, with elimination of parasites before the development of symptomatic infection.¹¹ Drugs active against hypnozoites will also facilitate malaria elimination campaigns.^{10,12} Primaquine (**1**, Figure 1), an 8-aminoquinoline (8-AQ), acts against liver forms of all *Plasmodium* species and is the only available antimalarial that acts against *P. vivax* and *P. ovale* hypnozoites, thereby preventing relapses after treatment for malaria. However, primaquine has significant side effects, including hemolysis in patients with glucose 6-phosphate dehydrogenase deficiency, a common genetic abnormality in malaria-endemic regions.¹³ The liver stage of infection has been underexploited as an antimalarial target because of the poorly understood biology of *Plasmodium* liver stages and technical difficulties in studying them.^{11,14} Only recently have systematic

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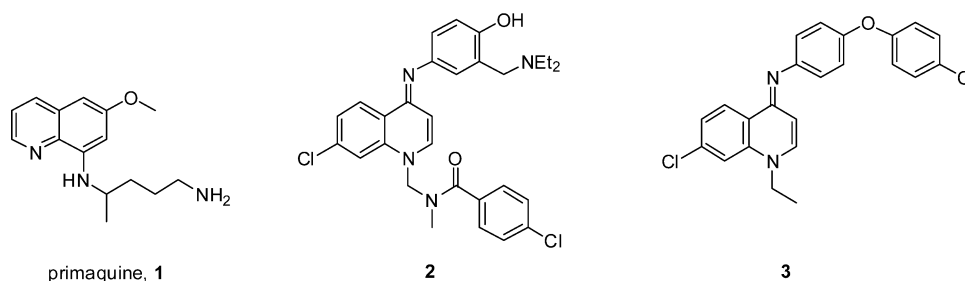
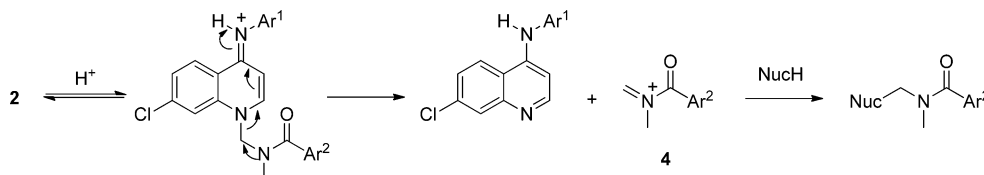


Figure 1. Structures of primaquine (1), a *N*-Mannich-based quinolon-4(1*H*)-imine (2), and a *N*-alkyl-quinolon-4(1*H*)-imine (3).

Scheme 1. Mechanism for the Acid-Catalyzed Hydrolysis of *N*-Mannich-Based Quinolon-4(1*H*)-imine, 2



efforts toward the identification of novel scaffolds active against *Plasmodium* liver stages, including phenotypic approaches, resulted in the discovery of potent compounds.^{15–17}

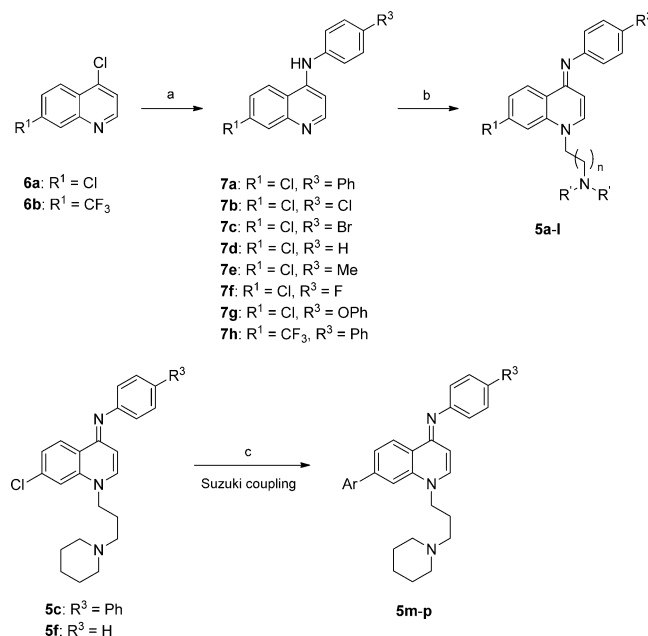
We previously reported that *N*-Mannich-based quinolon-4(1*H*)-imines derived from amodiaquine (e.g., 2, Figure 1) are potent antiplasmodial agents, with low nanomolar IC_{50} values against the chloroquine-resistant *P. falciparum* Dd2 strain.¹⁸ Quinolon-4(1*H*)-imines 2 were shown to hydrolyze to amodiaquine in aqueous solutions, particularly at low pH values, thus raising the possibility of these compounds acting as chemically activated prodrugs. The mechanism of hydrolysis probably involves protonation of the 4-imino nitrogen atom, followed by alkyl C–N bond scission via an S_N1 mechanism, leading to the formation of amodiaquine and the corresponding secondary amide (Scheme 1).¹⁸ This mechanism is similar to that reported for the acid-catalyzed hydrolysis of *N*-acyloxymethyl^{19–21} and *N*-amidomethyl prodrugs²² and involves the formation of an iminium ion intermediate, 4, which can alkylate biomolecules, potentially contributing to toxicity.²³ More recently, we have disclosed a library of *N*-alkyl-quinolon-4(1*H*)-imines (e.g., 3, Figure 1), lacking the *N*-Mannich base moiety, as potent antiplasmodial agents that target the liver stage of infection.²⁴ However, these compounds were shown to be degraded in rat liver microsomes with half-lives of ~ 4 h. We now report our strategy to improve stability toward aqueous buffers and microsomal enzymes by introducing an aminoalkyl side chain at the N-1 position while simultaneously exploiting the quinolon-4(1*H*)-imine scaffold to optimize antiplasmodial activity (5, Figure 2). The studies herein presented led to the discovery of dual-acting analogues active against both blood

and liver stages of parasites that are chemically and metabolically stable and lack apparent cytotoxicity.

RESULTS AND DISCUSSION

Chemistry. The general synthetic approach to quinolon-4(1*H*)-imines 5a–p is presented in Scheme 2. Briefly, 4-

Scheme 2. Synthesis of Quinolon-4(1*H*)-imines 5a–p^a



^aReagents and conditions: (a) R^3PhNH_2 , TEA, MeOH, rt, 24 h; (b) chloroalkylamine, NaH, DMF, rt, 5–12 h; (c) $PdCl_2(Ph_3P)_2$, *t*-Bu-Xphos, Na_2CO_3 1 M, boronic acid, dioxane, 100 °C, o/n.

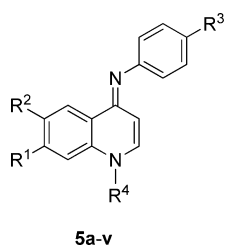


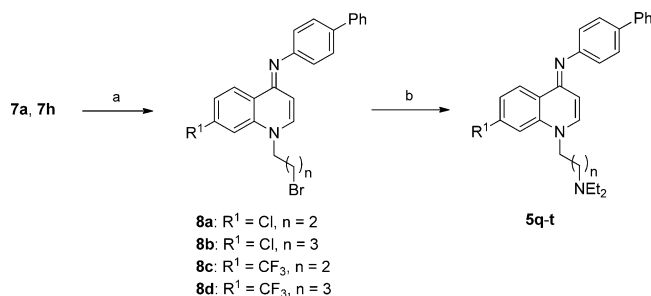
Figure 2. General structure for quinolon-4(1*H*)-imines 5 reported in this study. See Table 1 for structures 5a–v.

chloroquinolines 6a and 6b were reacted with the appropriate anilines in methanol and triethylamine to form the corresponding 4-anilinoquinolines 7a–h in very good to excellent yields. Intermediates 7 were subsequently transformed into the target compounds 5a–l in moderate to good yields by reaction with the appropriate chloroalkylamines in DMF in the presence of sodium hydride. X-ray analysis of compound 5c confirmed that the stereochemistry of the C=N bond corresponds to the *E*

isomer, in line with other reports.²⁵ Compounds **5m–p** were prepared from quinolon-4(1*H*)-imines **5c** and **5f** using Suzuki coupling with the appropriate boronic acids or pinacol esters. Surprisingly, the product isolated from the reaction of **5c** with 2-aminopyridine-5-boronic acid pinacol ester matches the C–N coupling product **5o**, as confirmed by mono- and bidimensional NMR analysis (see Supporting Information for NMR spectra and complete peak assignment).

Synthesis of compounds **5q–t** required alkylation of intermediates **7a** and **7h** with 1,3-dibromopropane or 1,4-dibromobutane to give the corresponding bromoalkyl-quinolon-4(1*H*)-imines **8a–d**, which were subsequently converted into the compounds **5q–t** by reaction with diethylamine in dichloromethane (Scheme 3). The route to 6-substituted

Scheme 3. Synthesis of Quinolon-4(1*H*)-imines **5q–t^a**



^aReagents and conditions: (a) Br(CH₂)₃Br or Br(CH₂)₄Br, NaH, DMF, rt, 5 h; (b) Et₂NH, THF, rt, o/n.

quinolon-4(1*H*)-imines **5u** and **5v** began with the synthesis of quinolines **6c** and **6d**, following reported procedures.^{26,27} Briefly, Gould–Jacobs protocol was applied to the appropriate anilines to form the 6-substituted quinolones **10a** and **10b** (Scheme 4). Hydrolysis of the ethyl ester followed by decarboxylation afforded their analogues **11a** and **11b**, which were converted into the corresponding 4-chloroquinolines by reaction with POCl₃. Reaction of **6c** and **6d** with the appropriate aniline followed by alkylation and reduction (as

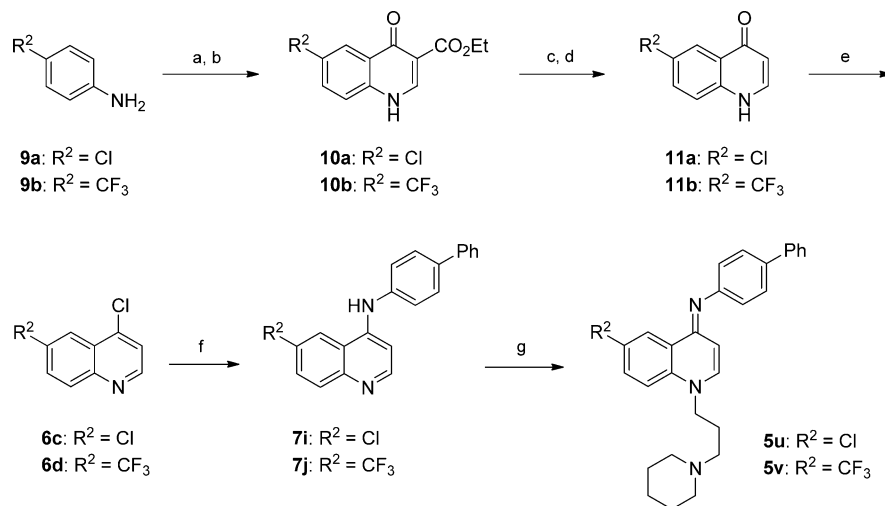
described before) provided the quinolon-4(1*H*)-imines **5u** and **5v** in good yields.

Activity against the Erythrocytic Stage of Infection and Cytotoxicity. Compounds **5a–v** were screened for activity against the erythrocytic stage of chloroquine-resistant *P. falciparum* W2 infection and for cytotoxicity against HEK293T mammalian cells (Table 1). Quinolon-4(1*H*)-imines **5** inhibited the growth of parasites with IC₅₀ values ranging from 50 to 520 nM while displaying negligible cytotoxicity, with EC₅₀ values against cultured mammalian cells ranging from 7 to ≥100 μM. This result indicates that the quinolon-4(1*H*)-imine scaffold, when attached to an alkylamino side chain through the N-1 nitrogen atom, provides potent antiplasmodial compounds comparable in potency to their *N*-Mannich-base counterparts, **2**,¹⁸ and are approximately 10-fold more potent than the previously reported *N*-alkyl analogues, **3** (IC₅₀ values ranging from 0.54 to 5.88 μM).²⁴ In general, most of the compounds **5a–v** presented selectivity indices (SI = EC₅₀(HEK293T)/IC₅₀(W2)) higher than 300, indicating that quinolon-4(1*H*)-imines are selective and nontoxic.

Inspection of the activity data against erythrocytic *P. falciparum* presented in Table 1 reveals that compounds with propyl or butyl linkers between the N-1 nitrogen atom of the quinolon-4(1*H*)-imine and the distal acyclic basic moiety are significantly more potent than those with the ethyl linker (e.g., **5q**, **5r** vs **5a**, and **5s**, **5t** vs **5j**). This difference is not observed for compounds containing a distal cyclic basic moiety (e.g., **5b** and **5c**). However, a reasonable correlation ($R^2 = 0.76$) was observed between the pIC₅₀ values for the 7-chloro and 7-trifluoromethyl series, suggesting that both series present similar structural requirements regarding the alkylamino side chain at the N-1 nitrogen (Figure 3).

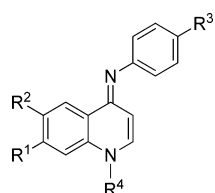
The effect of the electronic and hydrophobic properties of substituents in the quinolon-4(1*H*)-imine scaffold in the activity data in Table 1 was analyzed using the Hansch QSAR method. A good correlation ($R^2 = 0.82$) was found between the pIC₅₀ values and the Hansch π values for the substituents at the imine moiety, indicating that more lipophilic substituents improve activity against blood-stage parasites (Figure 4). In contrast, the electronic properties of the

Scheme 4. Synthesis of Quinolon-4(1*H*)-imines **5u and **5v**^a**



^aReagents and conditions: (a) (2-ethoxymethylene)malonate, 110 °C, 2 h; (b) Ph₂O, reflux, 1 h; (c) NaOH 2 M, MeOH, reflux, 18 h; (d) Ph₂O, reflux, 1 h; (e) POCl₃, 140 °C, 4 h; (f) 4-aminobiphenyl, TEA, MeOH, rt, 24 h; (g) 1-(3-chloropropyl)piperidine, NaH, DMF, rt, 12 h.

Table 1. Antiplasmodial Activity of Quinolon-4(1*H*)-imines **5a–v** and **3** against Blood and Liver Stages of Infection (*P. falciparum* W2 and *P. berghei*, respectively) and Their Cytotoxicity against Human Embryonic Kidney HEK293T Cells



Compd	R ¹	R ²	R ³	R ⁴	IC ₅₀ (nM)		EC ₅₀ (μM)	SI
					Blood-stage	Liver-stage		
5a	Cl	H	Ph		498	ND	37.1	74
5b	Cl	H	Ph		111	3460	>100	>900
5c	Cl	H	Ph		144	ND	41.8	290
5d	Cl	H	Cl		259	2696	>100	>386
5e	Cl	H	Br		74.0	3010	>100	>1351
5f	Cl	H	H		273	ND	>100	>366
5g	Cl	H	Me		331	ND	>100	>302
5h	Cl	H	F		369	4783	>100	>271
5i	Cl	H	OPh		160	ND	30.6	191
5j	CF ₃	H	Ph		525	4614	38.5	73
5k	CF ₃	H	Ph		226	ND	>100	>442
5l	CF ₃	H	Ph		360	ND	56.8	158
5m		H	H		216	1385	>100	>463
5n		H	Ph		55.4	2070	7.63	138
5o		H	Ph		357	440	7.34	21
5p		H	Ph		53.7	710	7.68	143
5q	Cl	H	Ph		151	ND	>100	>662
5r	Cl	H	Ph		67.6	2653	>100	>1479
5s	CF ₃	H	Ph		144	ND	>100	>694
5t	CF ₃	H	Ph		50.3	ND	>100	>1988
5u	H	Cl	Ph		99.7	ND	21.8	219
5v	H	CF ₃	Ph		631	ND	23.7	38
3					1090 ^a	87.0 ^a	--	--
CQ					186	ND	ND	--
PQ					ND	7500 ¹⁶	ND	--

^aFrom ref 24. ND: not determined; compounds not active at 1 μM. SI: selectivity index = EC₅₀(HEK293T)/IC₅₀(W2)).

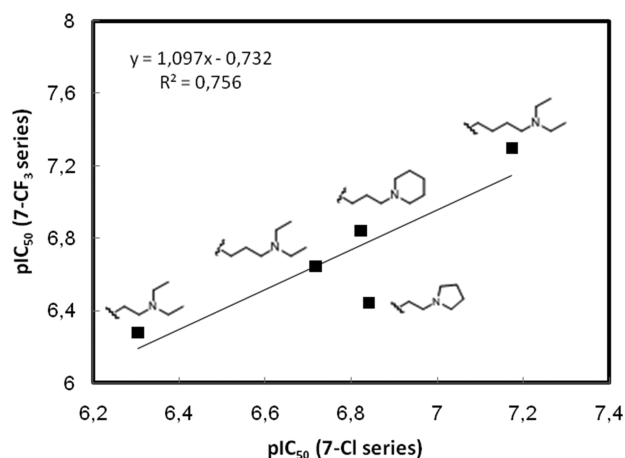


Figure 3. Plot of pIC_{50} against *P. falciparum* W2 for the 7- CF_3 series (5j–l,s,t) versus pIC_{50} for the 7-Cl series (5a–c,q,r). The data are from Table 1

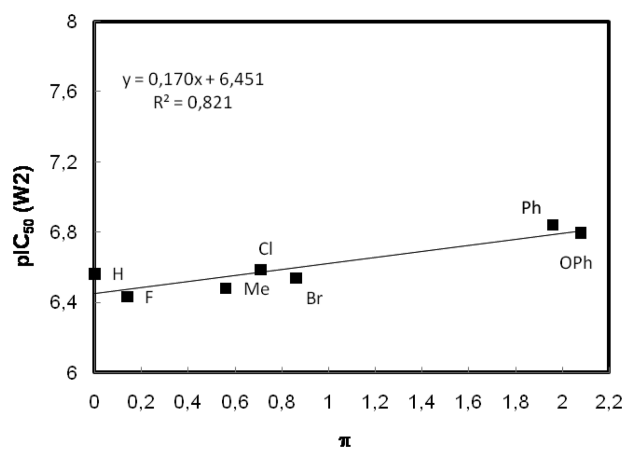


Figure 4. Plot of the pIC_{50} of the quinolon-4(1H)-imines 5c–i (*P. falciparum* W2 strain) versus the Hansch π values of the substituents in para-position of their phenylimine moiety.

substituents at the imine moiety do not affect the antiplasmodial activity (Figure 5). A moderate correlation ($R^2 = 0.70$) was observed between the pIC_{50} values and the Hammett σ values²⁸ for substituents in positions C-6 and C-7 of the quinolonimine moiety. Selection of Hammett constants was defined relative to the imine function, and thus, σ_{para} values were used for substituents at C-7 while σ_{meta} values were used for substituents at C-6. As shown in Figure 6, electron-withdrawing substituents at C-6 and C-7 are detrimental to activity of quinolon-4(1H)-imines against erythrocytic parasites. This effect probably reflects a decrease in the basicity of the quinolon-4(1H)-imine, which can be considered as a vinylogous cyclic amidine²⁹ with subsequent reduction of accumulation in the acidic digestive vacuole (DV) of the parasite, the site of action of aminoquinoline antimalarials. There is strong evidence that pH trapping plays a role in the activity of aminoquinoline antimalarials such as chloroquine.³⁰ Interestingly, derivative 5o, which contains a 2-pyridylamine substituent at C-7, is equipotent to its 7- CF_3 counterpart 5l, a result that might be ascribed to the protonation of the pyridine nitrogen atom within the parasite DV. The protonated 2-

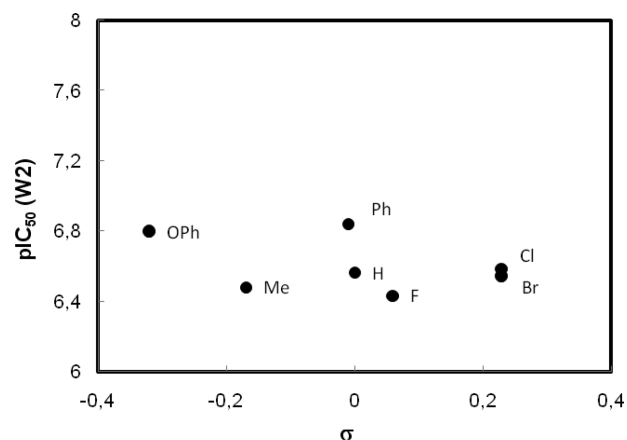


Figure 5. Plot of the pIC_{50} of the quinolon-4(1H)-imines 5c–i (*P. falciparum* W2 strain) versus the Hammett σ values²⁸ of the substituents in para-position of their phenylimine moiety.

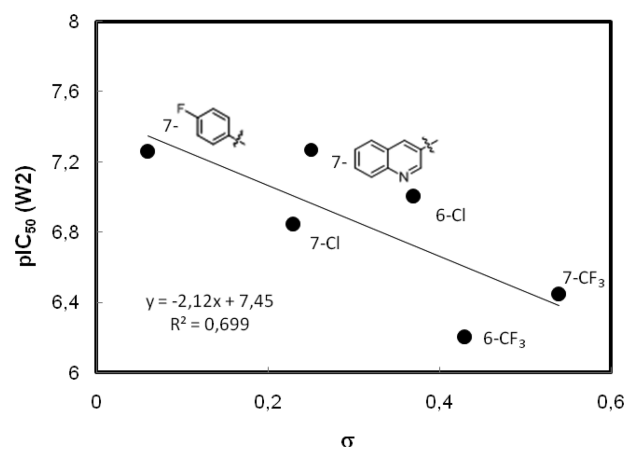


Figure 6. Plot of the pIC_{50} of the quinolon-4(1H)-imines 5c–i (*P. falciparum* W2 strain) versus the Hammett σ values²⁸ for substituents in positions C-6 and C-7 of the quinolonimine moiety.

pyridylamine is most likely a strong electron-withdrawing substituent, as suggested by the apparent σ_{ortho} value of 3.21 for the $=NH^+$ moiety (compared to the value of 0.56 for $=N$ group).²⁹

Inhibition of Heme Polymerization. *Plasmodium* parasites dispose the free, toxic ferriprotoporphyrin IX (FP) that results from digestion of host erythrocyte hemoglobin by crystallizing it into insoluble hemozoin crystals.^{31,32} Recent studies provided robust evidence that chloroquine and other quinoline antimalarials inhibit hemozoin formation by binding to free FP or to the surface of the growing hemozoin crystal.^{33,34} Since compounds 5 display a quinoline-related scaffold and two basic sites (the pK_a values for the quinolon-4(1H)-imine and distal amine groups are approximately 6–7 and 8–9, respectively³⁵), which might lead to accumulation in the acidic DV, we investigated whether they could also inhibit hemozoin formation. Selected quinolon-4(1H)-imines 5 were screened using a novel, simple, and efficient in vitro method based on the formation of hemozoin-like crystals.³⁶ The data presented in Table 2 reveal that the tested compounds 5 are either inactive or poor inhibitors of hemozoin formation when

Table 2. Inhibition of Hemozoin-like Crystal Formation by Selected Quinolone-4(1H)-imines **5 (Middle Column), Chloroquine and Gentamicin (Negative Control), and Comparison to IC₅₀ Values on Erythrocytic Stage (Right Column)^a**

compd	MIC (μM)	IC ₅₀ (W2) (nM)
5d	NI ^b	259
5f	NI ^b	273
5h	NI ^b	369
5i	500	160
5j	NI ^b	525
5k	NI ^b	226
5n	500	55.4
5u	1000	99.7
5v	1000	631
chloroquine	125	186
gentamicin	NI ^b	ND ^c

^aThe inhibition of hemozoin-like crystals formation was tested in the presence of 0–1000 μM of each compound. ^bNI: no inhibition of hemozoin-like crystal formation. ^cND: not determined.

compared to chloroquine, thus indicating that this is not the main mechanism of action of **5** against erythrocytic parasites.

Activity against the Mitochondrial Electron Transport Chain. Quinolone-4(1H)-imines **5** are structurally related to dihydroacridinedione **12** (WR 243251, Figure 7), a potent blood schizonticidal agent.^{37,38} Mechanistic studies suggested that dihydroacridinedione **12** targets the parasite mitochondrial electron transport chain (mtETC),³⁹ which is responsible for maintaining a constant pool of ubiquinone for the last step of pyrimidine biosynthesis, catalyzed by dihydroorotate dehydrogenase (DHODH).⁴⁰ The ketone hydrolysis product, a potential metabolite of **12**, appeared to target the Q_o quinol oxidation site of the cytochrome *bc*₁ complex, an essential component of the mtETC.⁴¹ A similar mode of action has been reported for the closely related class of antiparasitic 4(1H)-quinolones^{16,42,43} and 4(1H)-pyridones⁴⁴ (e.g., **13** and **14**, respectively). In order to investigate whether the antiparasitic mode of action of quinolone-4(1H)-imines involved inhibition of the parasite's *bc*₁ complex, we screened compounds **5b**, **5c**, **5l**, and **5r** using a phenotypic assay based on a *P. falciparum* 3D7 cell line transfected with yeast DHODH.⁴⁵ This cell line, which uses fumarate instead of ubiquinone as the final electron acceptor, has been shown previously to be resistant to mtETC inhibitors.⁴⁶ The results presented in Table 3 indicate that **5b**, **5c**, **5l**, and **5r** display similar antiparasitic activity against both 3D7-γDHODH and parental 3D7 strains, suggesting that inhibition of the *bc*₁ complex is not the primary antiparasitic mode of action for quinolone-4(1H)-imines **5**. This result was further supported by the observation that compound **5r** did not

Table 3. Growth Inhibition of Parental (3D7) and Transgenic (3D7-γDHODH) *P. falciparum* Strains with Quinolone-4(1H)-imines **5b, **5c**, **5l**, and **5r****

<i>P. falciparum</i> strain	IC ₅₀ ± SD (nM)			
	5b	5c	5l	5r
3D7	177 ± 14	73.7 ± 2.9	136 ± 19	64.7 ± 1.9
3D7-γDHODH	186 ± 42	52.3 ± 3.5	195 ± 82	68.3 ± 1.7

inhibit *bc*₁ complex activity at ≤10 μM (measured as ubiquinol/cyt c reductase), from *P. falciparum* cell-free extracts.

Activity against the Liver Stage of Infection. Quinolone-4(1H)-imines **5** were also evaluated for their antiparasitic activity against liver stages of the rodent parasite *P. berghei* and for cytotoxicity to Huh-7 human hepatoma cells, employing methods that were previously reported.⁴⁷ Primaquine was also included (1 and 10 μM) as a positive control for liver stage activity. Remarkably, nine compounds were more potent than primaquine and were nontoxic to Huh-7 human hepatoma cells at concentrations up to 5 μM. These compounds were then submitted to a dose–effect study, and the resulting IC₅₀ values for the inhibition of hepatic *P. berghei* infection ranged from 0.44 to 4.6 μM, which compare favorably with the IC₅₀ value of 7.5 μM determined for primaquine (Table 1). Notably, the most active compounds **5o** and **5p** (IC₅₀ values of 0.44 and 0.71 μM, respectively), which have heterocyclic moieties in position C-7 of the quinolinimine scaffold, were at least 10-fold more potent than primaquine, establishing these novel quinolinimine derivatives as promising dual-stage antimalarials. Currently, the mechanism of action for quinolone-4(1H)-imines **5o** and **5p** is unknown, but in contrast to their quinolone counterparts (e.g., decoquinone, **13**), it is unlikely that they target the parasite mtETC,^{16,42,45} as described in the previous section.

Metabolic and Chemical Stabilities. The microsomal stability of a selected set of quinolone-4(1H)-imines was determined using rat liver microsomes. Inspection of Table 4

Table 4. Results from the in Vitro Metabolism Studies Using Rat Liver Microsomes^a

compd	% remaining	compd	% remaining
5a	68.0	5l	65.0
5i	73.0	5r	62.7
5j	80.4	5t	81.5
5k	79.0	5u	75.0

^aData present the percentage of remaining compound after 6 h of incubation with microsomes in the presence of the NADPH-regenerating system.

reveals that *N*-alkylaminequinolone-4(1H)-imines **5** display excellent metabolic stability, with most of the compounds

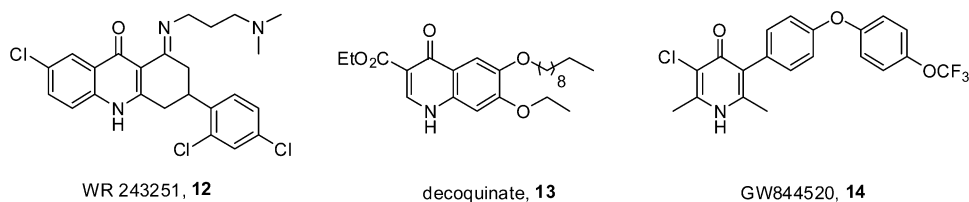


Figure 7. Chemical structures of *Plasmodium falciparum* *bc*₁ complex inhibitors containing dihydroacridinedione (**12**), 4(1H)-quinolone (**13**), and 4(1H)-pyridone (**14**) cores.

presenting negligible degradation after 6 h of incubation, in contrast with the previously reported *N*-alkyl counterparts (e.g., **3**, Figure 1), which were degraded in rat liver microsomes with half-lives of ~4 h.²⁴ These results also suggest that the rate of degradation is not significantly affected by the length of the aminoalkyl chain (e.g., **5j** vs **5t**) or by the nature of substituents at the quinolon-4(1*H*)-imine moiety (e.g., **5r** vs **5t**). A similar result was observed when acyclic terminal amines were replaced by cyclic amines (e.g., **5j** vs **5k**). None of the selected compounds showed any degradation in the control experiments (with the NADPH-regenerating system absent), suggesting that there was no major noncofactor dependent metabolism contributing to their degradation in liver microsomes. Furthermore, compound **5u** showed no significant degradation when incubated in pH 7.4 and 0.3 aqueous buffers at 37 °C for 10 days, suggesting that the *N*-alkylamine-quinolon-4(1*H*)-imines are stable compounds, in contrast to their *N*-Mannich-base counterparts.¹⁸

The MS scan data from the incubation mixtures of compound **5u** in microsomes were screened for the presence of fragments corresponding to the putative metabolites (see Supporting Information). The reaction profile of **5u** revealed the formation of the corresponding quinoline as the major metabolite, suggesting that *N*-dealkylation is the predominant pathway.

CONCLUSION

We report that quinolon-4(1*H*)-imines **5** bearing an alkylamine side-chain at N-1 are a new family of efficient dual-stage antiplasmodial agents endowed with chemical and metabolic stability. Compounds **5** inhibited the development of intra-erythrocytic forms of *P. falciparum* with IC₅₀ values in the nanomolar range and were associated with low cytotoxicity against HEK293T mammalian cells. The length of the side chain between N-1 and the distal amine seems to be critical for activity, with side chains with three and four carbon atoms being most active. Lipophilic substituents in the iminoaryl moiety improve potency, while strong electron-withdrawing substituents at C-6 and C-7 of the quinolonimine moiety were detrimental to activity against the erythrocytic stage parasites. In spite of structural resemblance of compounds **5** to antimalarial 4-aminoquinolines, which includes the presence of two basic groups that can lead to concentration in the acidic DV of the parasite, the primary mechanism of action of quinolon-4(1*H*)-imines **5** does not appear to involve inhibition of heme crystallization. In addition, a phenotypic assay based on a *P. falciparum* cell line transfected with cytoplasmic yeast DHODH suggested that inhibition of the *bc*₁ complex is not the primary antimalarial mode of action for quinolon-4(1*H*)-imines **5**. Additional studies on the mechanism of action of new quinolon-4(1*H*)-imines are underway.

Quinolon-4(1*H*)-imines **5** were also active against the liver stage of infection. Derivative **5p** was the most promising compound, with IC₅₀ values of 54 and 710 nM against the erythrocytic and liver stages of infection, respectively, with minimal toxicity against human cells. Importantly, compounds **5** displayed considerable stability toward hepatic enzymes and aqueous buffers, supporting the conclusion that extending the linker between N-1 and the nitrogen atom is an effective approach to remove the source of reactivity of their *N*-Mannich-base and *N*-alkyl counterparts, **2** and **3**, respectively. Overall, our study reveals that quinolon-4(1*H*)-imines such as

5p are a good starting point to exploit for lead optimization toward developing dual-stage antimalarials.

EXPERIMENTAL SECTION

Chemistry. All manipulations requiring anhydrous conditions were carried out in flame-dried glassware, with magnetic stirring and under a nitrogen atmosphere. Chemicals and anhydrous solvents were obtained from commercial sources (Sigma-Aldrich or Alfa Aesar) and were used as received. 4,6-Dichloroquinoline (**6c**) and 4-chloro-6-(trifluoromethyl)quinolone (**6d**) were prepared according to literature procedures, and their analytical data were in agreement with those already published^{26,27} (see Supporting Information). Analytical TLC plates, aluminum sheets with silica gel F250 (Merck), were visualized by UV light (254 or 366 nm) or stained with KMnO₄ or with a solution of ninhydrin in ethanol. Flash chromatography was performed on Kieselgel 60 GF₂₅₄ silica (Merck) of 0.040–0.063 mm.

Melting points were determined on a Kofler melting point apparatus. NMR spectra were recorded on a Bruker 400 Ultra-Shield (¹H, 400.13 MHz; ¹³C, 100.61 MHz). Data are reported as follows: chemical shift, multiplicity (s, singlet; d, doublet; t, triplet; dd, doublet of doublets; dt, doublet of triplets; q, quartet; m, multiplet; br, broad), coupling constants (*J* in Hz), integration, and assignment. Chemical shifts are given in parts per million (ppm) with the solvent signal as internal standard [CDCl₃ δ (¹H) 7.26 ppm and δ (¹³C) 77.2 ppm, and CD₃OD δ (¹H) 4.87 ppm and δ (¹³C) 49.0 ppm]. High resolution mass spectrometry (HRMS) was performed on a Bruker MicrOTOF equipped with ESI ion source from the Mass Spectrometry and Proteomics Unity of the University of Santiago de Compostela, Spain. To determine the purity of the final compounds (**5a–v**), HPLC experiments were conducted using an ELITE LaChrom VWR HITACHI equipment (pump + controller L-2130, UV detector L-2400) and a column LiChroCART, RP-18e, 5 μm. All quinolon-4(1*H*)-imines tested in biological assays were ≥95% pure. Conditions and retention times are described in the Supporting Information.

General Procedure A for *N*-Alkylated Derivatives (5a–l**, **5u**, **5v**).** To a solution of the corresponding quinolin-4-amine in anhydrous DMF (3.0 mL/mmol), under N₂ atmosphere, was slowly added NaH (4.0 equiv, 40% dispersion in oil) at room temperature. The mixture was stirred until evolution of hydrogen ceased (10–60 min). The desired alkylamine chloride reagent (1.5 equiv) and NaI (3.0 equiv) were added, and the mixture was stirred at 80 °C for 5–12 h. The mixture was then concentrated under reduced pressure, diluted with DCM, and washed with brine. The aqueous phase was extracted with DCM (3 × 50 mL), and the combined organic phases were dried over Na₂SO₄. The crude product was then purified by flash chromatography (DCM/MeOH/TEA, 9:0.9:0.1) and recrystallized from different systems (e.g., DCM/MeOH) to afford the final pure product.

General Procedure B for *N*-Alkylated Derivatives (5q–t**).** To a solution of the corresponding quinolin-4-amine in anhydrous DMF (3 mL/mmol), under N₂ atmosphere, was slowly added NaH (4 equiv, 40% dispersion in oil) at room temperature. The mixture was stirred until evolution of hydrogen ceased (10–60 min). The desired alkyl dibromide reagent (1.5 equiv) was then added, and the mixture was stirred at room temperature. The mixture was then concentrated under reduced pressure, diluted with DCM, and washed with brine. The aqueous phase was extracted with DCM (3 × 50 mL), and the combined organic phases were dried over Na₂SO₄. The resulting product and NaI (1.5 equiv) were then dissolved in DEA (15 mL/mmol) and stirred at room temperature for 78 h. The mixture was concentrated under reduced pressure and the crude product was purified by flash chromatography (DCM/MeOH/Et₃N, 9:0.9:0.1) and recrystallized from different systems (e.g., DCM/MeOH) to afford the final pure product.

General Procedure for Suzuki Coupling (5m–p**).** To a solution of the aryl halide in dry 1,4-dioxane at room temperature was consecutively added PdCl₂(Ph₃P)₂ (0.1 equiv), *t*-Bu-Xphos (0.1 equiv), Na₂CO₃ 1 M (3 equiv), and boronic acid (1.2 equiv). After degassing for 5 min, the resulting mixture was heated at 100 °C under

N₂ atmosphere overnight. The mixture was then cooled to room temperature, diluted with DCM, and filtered through a pad of Celite. The filtrate was concentrated under reduce pressure and the crude product was purified by thin layer chromatography (DCM/MeOH/Et₃N, 95:4:1) to afford the corresponding 7-substituted quinolin-4-imines.

(E)-N-(7-Chloro-1-(2-(diethylamino)ethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5a). According to general procedure A, the product was obtained as a yellow solid (37 mg, 11%). Mp 115–117 °C. ¹H NMR (CD₃OD) δ 8.68 (d, J = 8.8 Hz, 1H), 8.43 (d, J = 7.6 Hz, 1H), 8.37 (s, 1H), 7.90–7.86 (m, 3H), 7.72–7.70 (m, 2H), 7.58–7.56 (m, 2H), 7.52–7.49 (m, 2H), 7.43–7.41 (m, 1H), 6.98 (d, J = 7.2 Hz, 1H), 4.66 (t, J = 5.6 Hz, 2H), 2.90 (t, J = 5.6 Hz, 2H), 2.58–2.53 (m, 4H), 0.87–0.83 (m, 6H). ¹³C NMR (CD₃OD) δ 154.3, 148.4, 140.8, 139.3, 135.9, 128.7, 128.3, 127.7, 127.6, 126.6, 125.7, 125.5, 117.9, 99.9, 53.3, 51.2, 46.9, 10.9. HRMS (ESI) *m/z* calcd for [C₂₇H₂₉ClN₃]⁺, 430.2045 [M + H⁺]; found, 430.2032.

(E)-N-(7-Chloro-1-(2-(pyrrolidin-1-yl)ethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5b). According to general procedure A, the product was obtained as a yellow solid (49 mg, 39%). Mp 221–224 °C. ¹H NMR (CD₃OD) δ 8.56 (d, J = 9.0 Hz, 1H), 7.71–7.55 (m, 4H), 7.45 (t, J = 7.7 Hz, 2H), 7.38–7.21 (m, 3H), 7.02 (d, J = 8.4 Hz, 2H), 6.92 (d, J = 8.0 Hz, 1H), 6.02 (d, J = 8.0 Hz, 1H), 3.99 (t, J = 7.1 Hz, 2H), 2.86 (t, J = 7.1 Hz, 2H), 2.60 (d, J = 5.3 Hz, 4H), 1.91–1.73 (m, 4H). ¹³C NMR (CD₃OD) δ 154.6, 152.4, 141.6, 140.2, 140.0, 137.4, 135.3, 129.0, 128.4, 128.2, 127.0, 126.9, 124.3, 124.0, 122.1, 114.4, 101.7, 54.9, 54.3, 51.8, 23.9. HRMS (ESI) *m/z* calcd for [C₂₇H₂₇ClN₃]⁺, 428.1888 [M + H⁺]; found, 428.1898.

(E)-N-(7-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5c). According to general procedure A, the product was obtained as a yellow solid (50 mg, 48%). Mp 176–178 °C. ¹H NMR (CDCl₃) δ 8.56 (d, J = 8.7 Hz, 1H), 7.67–7.57 (m, 4H), 7.46 (t, J = 7.7 Hz, 2H), 7.35–7.20 (m, 3H), 7.06–6.95 (m, 3H), 6.00 (d, J = 8.0 Hz, 1H), 3.98 (t, J = 6.6 Hz, 2H), 2.37 (br, 4H), 2.31 (t, J = 6.5 Hz, 2H), 1.97–1.91 (m, 2H), 1.65–1.56 (m, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 154.4, 152.2, 141.3, 140.0, 137.1, 134.9, 128.7, 128.0, 127.9, 126.9, 127.0, 127.0, 124.1, 123.6, 121.8, 114.4, 101.0, 54.9, 54.5, 49.6, 26.1, 25.6, 24.5. HRMS (ESI) *m/z* calcd for [C₂₉H₃₁ClN₃]⁺, 456.2201 [M + H⁺]; found, 456.2199.

(E)-4-Chloro-N-(7-chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)aniline (5d). According to general procedure A, the product was obtained as a yellow solid (205 mg, 34%). Mp 157–159 °C. ¹H NMR (CDCl₃) δ 8.55 (d, J = 8.7 Hz, 1H), 7.36 (s, 1H), 7.33–7.20 (m, 3H), 7.06 (d, J = 8.0 Hz, 1H), 6.96–6.83 (m, 2H), 5.90 (d, J = 8.0 Hz, 1H), 4.00 (t, J = 6.6 Hz, 2H), 2.36 (br, 4H), 2.30 (t, J = 6.5 Hz, 2H), 1.93 (dt, J = 13.1, 6.5 Hz, 2H), 1.62 (dt, J = 10.9, 5.5 Hz, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 154.8, 148.9, 140.4, 139.8, 137.4, 129.3, 128.0, 127.3, 123.9, 123.4, 123.0, 114.5, 100.6, 54.8, 54.5, 49.7, 26.1, 25.5, 24.5. HRMS (ESI) *m/z* calcd for [C₂₃H₂₆Cl₂N₃]⁺, 414.1498 [M + H⁺]; found, 414.1497.

(E)-4-Bromo-N-(7-chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)aniline (5e). According to general procedure A, the product was obtained as a yellow solid (133 mg, 29%). Mp 165–167 °C. ¹H NMR (CDCl₃) δ 8.55 (d, J = 8.7 Hz, 1H), 7.49–7.40 (m, 2H), 7.36 (d, J = 1.8 Hz, 1H), 7.31–7.21 (m, 1H), 7.06 (d, J = 8.0 Hz, 1H), 6.85 (t, J = 5.7 Hz, 2H), 5.90 (d, J = 8.0 Hz, 1H), 4.00 (t, J = 6.5 Hz, 2H), 2.36 (br, 4H), 2.29 (t, J = 6.5 Hz, 2H), 1.96–1.91 (m, 2H), 1.69–1.56 (m, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 154.7, 140.5, 139.8, 137.4, 132.3, 128.0, 123.9, 123.5, 114.9, 114.6, 100.6, 54.8, 54.5, 49.7, 26.1, 25.5, 24.5. HRMS (ESI) *m/z* calcd for [C₂₃H₂₆BrClN₃]⁺, 460.0973 [M + H⁺]; found, 460.0965.

(E)-N-(7-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)aniline (5f). According to general procedure A, the product was obtained as a yellow solid (133 mg, 40%). Mp 137–139 °C. ¹H NMR (CDCl₃) δ 8.55 (d, J = 8.7 Hz, 1H), 7.41–7.32 (m, 3H), 7.24 (dd, J = 8.7, 1.9 Hz, 1H), 7.09–7.03 (m, 1H), 7.02–6.87 (m, 3H), 5.91 (d, J = 8.0 Hz, 1H), 3.97 (t, J = 6.5 Hz, 2H), 2.39 (br, 4H), 2.29 (t, J = 6.5 Hz, 2H), 2.00–1.83 (m, 2H), 1.69–1.54 (m, 4H), 1.46 (br, 2H). ¹³C NMR (CDCl₃) δ 154.4, 140.0, 137.0, 129.3, 127.9, 123.8, 123.6, 122.2, 121.5, 114.4, 100.9, 54.8, 54.5, 49.5, 26.1, 25.5, 24.5.

HRMS (ESI) *m/z* calcd for [C₂₃H₂₇ClN₃]⁺, 380.1888 [M + H⁺]; found, 380.1881.

(E)-N-(7-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)-4-methylaniline (5g). According to general procedure A, the product was obtained as a yellow solid (30 mg, 20%). Mp 164–166 °C. ¹H NMR (CDCl₃) δ 8.59 (d, J = 8.4 Hz, 1H), 7.33 (d, J = 1.7 Hz, 1H), 7.24 (dd, J = 8.4, 1.7 Hz, 1H), 7.16 (d, J = 7.9 Hz, 2H), 6.99 (d, J = 7.9 Hz, 1H), 6.87 (d, J = 8.0 Hz, 2H), 5.96 (d, J = 8.0 Hz, 1H), 3.98 (t, J = 6.6 Hz, 2H), 2.36 (br, 4H), 2.36 (br, 3H), 2.29 (t, J = 6.5 Hz, 2H), 1.95–1.89 (m, 2H), 1.67–1.54 (m, 4H), 1.46 (br, 2H). ¹³C NMR (CDCl₃) δ 154.4, 140.0, 139.9, 137.1, 131.7, 130.0, 128.0, 123.7, 121.4, 114.4, 100.9, 54.8, 54.5, 49.6, 26.1, 25.5, 24.4, 20.9. HRMS (ESI) *m/z* calcd for [C₂₄H₂₉ClN₃]⁺, 394.2045 [M + H⁺]; found, 394.2040.

(E)-N-(7-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)-4-fluoroaniline (5h). According to general procedure A, the product was obtained as a yellow solid (38 mg, 15%). Mp 123–125 °C. ¹H NMR (CDCl₃) δ 8.55 (d, J = 8.6 Hz, 1H), 7.34 (s, 1H), 7.32–7.18 (m, 1H), 7.06–7.02 (m, 3H), 6.89 (dd, J = 7.6, 5.2 Hz, 2H), 5.90 (d, J = 7.9 Hz, 1H), 3.99 (t, J = 6.4 Hz, 2H), 2.36 (br, 4H), 2.30 (t, J = 6.4 Hz, 2H), 2.04–1.85 (m, 2H), 1.64–1.59 (m, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 160.0, 157.6, 154.9, 140.2, 139.9, 137.2, 127.9, 123.8, 123.4, 122.6, 115.7, 114.5, 100.6, 54.8, 54.5, 49.6, 26.1, 25.5, 24.5. HRMS (ESI) *m/z* calcd for [C₂₃H₂₆ClFN₃]⁺, 398.1794 [M + H⁺]; found, 398.1788.

(E)-N-(7-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)-4-phenoxyaniline (5i). According to general procedure A, the product was obtained as a yellow solid (42 mg, 21%). Mp 154–156 °C. ¹H NMR (CDCl₃) δ 8.84 (d, J = 8.7 Hz, 1H), 7.49 (s, 1H), 7.45–7.31 (m, 4H), 7.17–6.95 (m, 7H), 6.19 (d, J = 7.7 Hz, 1H), 4.15 (t, J = 6.2 Hz, 2H), 2.36 (br, 4H), 2.30 (t, J = 6.2 Hz, 2H), 2.05–1.89 (m, 2H), 1.69–1.55 (m, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 157.8, 154.8, 153.2, 141.8, 139.7, 138.3, 129.7, 128.4, 124.9, 123.9, 122.9, 122.0, 118.3, 115.0, 100.7, 54.7, 54.5, 50.4, 26.1, 25.6, 24.4. HRMS (ESI) *m/z* calcd for [C₂₉H₃₁ClN₃O]⁺, 472.2150 [M + H⁺]; found, 472.2159.

(E)-N-(1-(2-(Diethylamino)ethyl)-7-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5j). According to general procedure A, the product was obtained as a yellow solid (97 mg, 21%). Mp 148–150 °C. ¹H NMR (CD₃OD) δ 8.94 (d, J = 8.8 Hz, 1H), 8.71 (d, J = 7.6 Hz, 1H), 8.55 (s, 1H), 8.15 (d, J = 8.8 Hz, 1H), 7.89–7.87 (m, 2H), 7.72–7.70 (m, 2H), 7.66–7.62 (m, 2H), 7.52–7.48 (m, 2H), 7.42–7.40 (m, 1H), 7.11 (d, J = 7.6 Hz, 1H), 5.29 (br, 2H), 3.54 (br, 2H), 3.19 (br, 2H), 1.26 (br, 6H). ¹³C NMR (CD₃OD) δ 155.7, 148.8, 141.3, 139.6, 135.5, 128.7, 128.4, 127.6, 126.6, 126.2, 125.6, 123.0, 120.9, 115.6, 101.7, 49.9, 48.1, 47.0, 18.8. HRMS (ESI) *m/z* calcd for [C₂₈H₂₉F₃N₃]⁺, 464.2308 [M + H⁺]; found, 464.2315.

(E)-N-(1-(2-(Pyrrolidin-1-yl)ethyl)-7-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5k). According to general procedure A, the product was obtained as a yellow solid (37 mg, 30%). Mp 238–240 °C. ¹H NMR (CD₃OD) δ 8.92 (d, J = 8.8 Hz, 1H), 8.72 (d, J = 7.2 Hz, 1H), 8.55 (s, 1H), 8.13 (d, J = 8.8 Hz, 1H), 7.87 (d, J = 8.4 Hz, 2H), 7.70 (d, J = 7.6 Hz, 2H), 7.63 (d, J = 8.4 Hz, 2H), 7.49 (t, J = 7.6 Hz, 2H), 7.40 (t, J = 7.6 Hz, 1H), 7.09 (d, J = 7.2 Hz, 1H), 4.99 (t, J = 6.4 Hz, 2H), 3.39 (br, 2H), 3.01 (br, 4H), 1.95 (m, 4H). ¹³C NMR (CD₃OD) δ 155.6, 148.6, 141.5, 139.6, 138.3, 135.7, 135.3, 128.7, 128.3, 127.6, 126.6, 126.1, 125.6, 124.6, 122.9, 115.8, 101.6, 54.0, 53.2, 52.0, 23.0. HRMS (ESI) *m/z* calcd for [C₂₈H₂₇F₃N₃]⁺, 462.2152 [M + H⁺]; found, 462.2146.

(E)-N-(1-(3-(Piperidin-1-yl)propyl)-7-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5l). According to general procedure A, the product was obtained as a yellow solid (33 mg, 21%). Mp 174–177 °C. ¹H NMR (CDCl₃) δ 8.74 (d, J = 8.3 Hz, 1H), 7.68–7.58 (m, 4H), 7.56–7.49 (m, 2H), 7.49–7.41 (m, 2H), 7.36–7.30 (m, 1H), 7.12–6.99 (m, 3H), 6.05 (d, J = 8.0 Hz, 1H), 4.06 (t, J = 6.5 Hz, 2H), 2.36 (br, 2H), 2.32 (t, J = 6.5 Hz, 2H), 1.98–1.92 (m, 2H), 1.64–1.55 (m, 4H), 1.47 (br, 2H). ¹³C NMR (CDCl₃) δ 154.2, 152.0, 141.2, 140.4, 138.2, 135.1, 132.7, 132.4, 128.7, 128.0, 127.6, 126.7, 126.6, 125.3, 123.0, 122.6, 121.6, 119.3, 111.8, 111.8,

101.2, 54.9, 54.5, 49.6, 26.1, 25.4, 24.5. HRMS (ESI) m/z calcd for $[C_{30}H_{31}F_3N_3]^+$, 490.2465 $[M + H]^+$; found, 490.2465.

(E)-N-(7-(4-Fluorophenyl)-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)aniline (5m). Yellow solid (32 mg, 80%). Mp 132–134 °C. 1H NMR (CD_3OD) δ 8.68 (d, $J = 8.6$ Hz, 1H), 8.49 (d, $J = 7.2$ Hz, 1H), 8.28 (s, 1H), 8.10 (d, $J = 8.6$ Hz, 1H), 7.97–7.89 (m, 2H), 7.62–7.54 (m, 2H), 7.50–7.42 (m, 3H), 7.31 (t, $J = 8.4$ Hz, 2H), 6.84 (d, $J = 7.2$ Hz, 1H), 4.74 (t, $J = 6.8$ Hz, 2H), 2.65–2.30 (m, 6H), 2.24–2.15 (m, 2H), 1.57–1.48 (m, 4H), 1.43 (br, 2H). ^{13}C NMR (CD_3OD) δ 165.1, 156.9, 148.6, 147.7, 140.4, 138.4, 136.1, 131.4, 131.1, 129.3, 127.7, 126.7, 125.9, 118.9, 117.3, 116.7, 101.4, 56.1, 55.4, 54.0, 26.7, 26.3, 24.8. HRMS (ESI) m/z calcd for $[C_{29}H_{31}FN_3]^+$, 440.2497 $[M + H]^+$; found, 440.2498.

(E)-N-(7-(4-Fluorophenyl)-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)-[1,1'-biphenyl]-4-amine (5n). Yellow solid (52 mg, 76%). Mp 158–160 °C. 1H NMR (CD_3OD) δ 8.73 (d, $J = 8.8$ Hz, 1H), 8.54 (d, $J = 7.4$ Hz, 1H), 8.33 (s, 1H), 8.15 (d, $J = 8.8$ Hz, 1H), 8.00–7.93 (m, 2H), 7.87 (d, $J = 8.5$ Hz, 2H), 7.71 (d, $J = 7.6$ Hz, 2H), 7.58 (d, $J = 8.5$ Hz, 2H), 7.50 (t, $J = 7.6$ Hz, 2H), 7.44–7.32 (m, 3H), 6.98 (d, $J = 7.4$ Hz, 1H), 4.78 (t, $J = 7.0$ Hz, 2H), 2.52–2.32 (m, 6H), 2.26–2.16 (m, 2H), 1.58–1.50 (m, 4H), 1.46 (br, 2H). ^{13}C NMR (CD_3OD) δ 165.4, 157.1, 148.8, 148.0, 142.6, 141.4, 140.7, 137.8, 136.4, 131.4, 130.4, 130.0, 129.2, 128.2, 128.0, 127.2, 126.2, 119.2, 117.5, 116.9, 101.8, 56.1, 55.4, 54.1, 26.7, 26.2, 24.7. HRMS (ESI) m/z calcd for $[C_{35}H_{35}FN_3]^+$, 516.2810 $[M + H]^+$; found, 516.2800.

(E)-4-([1,1'-Biphenyl]-4-ylimino)-1-(3-(piperidin-1-yl)propyl)-N-(pyridin-2-yl)-1,4-dihydroquinolin-7-amine (5o). Yellow solid (16 mg, 24%). Mp 172–174 °C. 1H NMR (CD_3OD) δ 9.21 (s, 1H), 8.46 (d, $J = 9.3$ Hz, 1H), 8.40 (d, $J = 4.3$ Hz, 1H), 8.36 (d, $J = 7.4$ Hz, 1H), 7.82 (d, $J = 8.4$ Hz, 2H), 7.77–7.66 (m, 4H), 7.53 (d, $J = 8.4$ Hz, 2H), 7.48 (t, $J = 7.6$ Hz, 2H), 7.39 (t, $J = 7.6$ Hz, 1H), 7.11–6.95 (m, 2H), 6.81 (d, $J = 7.4$ Hz, 1H), 4.58 (t, $J = 7.0$ Hz, 2H), 2.73–2.58 (m, 2H), 2.55 (br, 4H), 2.39–2.23 (m, 2H), 1.66–1.54 (m, 4H), 1.48 (br, 2H). ^{13}C NMR (CD_3OD) δ 156.4, 155.7, 148.9, 148.5, 147.3, 141.9, 141.9, 141.2, 139.2, 137.8, 130.1, 129.6, 128.9, 128.0, 126.9, 125.6, 120.6, 118.1, 113.9, 113.2, 102.7, 100.3, 56.5, 55.3, 54.3, 26.2, 25.8, 24.8. HRMS (ESI) m/z calcd for $[C_{34}H_{36}N_5]^+$, 514.2965 $[M + H]^+$; found, 514.2966.

(E)-N-(1'-(3-(Piperidin-1-yl)propyl)-[3,7'-biquinolin]-4'(1'H)-ylidene)-[1,1'-biphenyl]-4-amine (5p). Orange solid (69 mg, 62%). Mp 152–154 °C. 1H NMR ($CDCl_3$) 9.23 (d, $J = 2.1$ Hz, 1H), 8.86 (d, $J = 7.1$ Hz, 1H), 8.40 (d, $J = 2.1$ Hz, 1H), 8.18 (d, $J = 8.4$ Hz, 1H), 7.94 (d, $J = 7.9$ Hz, 1H), 7.77 (t, $J = 8.4$ Hz, 1H), 7.70–7.58 (m, 7H), 7.43 (t, $J = 7.6$ Hz, 2H), 7.31 (t, $J = 7.6$ Hz, 1H), 7.18 (d, $J = 7.5$ Hz, 1H), 7.11 (d, $J = 8.0$ Hz, 2H), 6.12 (d, $J = 7.5$ Hz, 1H), 4.16 (t, $J = 6.3$ Hz, 2H), 2.42–2.26 (m, 6H), 2.05–1.97 (m, 2H), 1.55–1.47 (m, 4H), 1.44–1.34 (br, 2H). ^{13}C NMR ($CDCl_3$) δ 155.0, 149.7, 147.8, 141.2, 140.9, 140.9, 139.6, 135.4, 135.2, 134.0, 133.2, 130.0, 129.4, 128.8, 128.2, 128.0, 128.0, 127.7, 127.4, 126.8, 126.7, 124.6, 123.0, 122.4, 113.5, 100.8, 55.0, 54.9, 50.1, 26.1, 25.7, 24.5. HRMS (ESI) m/z calcd for $[C_{38}H_{37}N_4]^+$, 549.3013 $[M + H]^+$; found, 549.3003.

(E)-N-(7-Chloro-1-(3-(diethylamino)propyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5q). According to general procedure B, the product was obtained as a yellow solid (16 mg, 8%). Mp 95–98 °C. 1H NMR ($CDCl_3$) δ 8.68 (d, $J = 8.9$ Hz, 1H), 7.73–7.52 (m, 4H), 7.52–7.41 (m, 3H), 7.41–7.22 (m, 2H), 7.17–7.05 (m, 3H), 6.09 (d, $J = 7.7$ Hz, 1H), 4.02 (t, $J = 6.6$ Hz, 2H), 2.56 (dd, $J = 14.1, 7.0$ Hz, 4H), 2.47 (t, $J = 6.2$ Hz, 2H), 1.98–1.84 (m, 2H), 1.10–0.94 (m, 6H). ^{13}C NMR ($CDCl_3$) δ 154.6, 141.4, 140.6, 139.7, 137.6, 128.7, 128.2, 128.0, 126.7, 124.0, 122.2, 119.9, 114.6, 101.0, 50.3, 49.3, 46.4, 26.2, 11.5. HRMS (ESI) m/z calcd for $[C_{28}H_{31}ClN_3]^+$, 444.2201 $[M + H]^+$; found, 444.2200.

(E)-N-(7-Chloro-1-(4-(diethylamino)butyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5r). According to general procedure B, the product was obtained as a yellow solid (34 mg, 12%). Mp 108–110 °C. 1H NMR ($CDCl_3$) δ 8.59 (d, $J = 8.7$ Hz, 1H), 7.72–7.57 (m, 4H), 7.52–7.40 (m, 2H), 7.37–7.33 (m, 2H), 7.29–7.24 (m, 2H), 7.04 (d, $J = 8.3$ Hz, 2H), 6.93 (d, $J = 8.0$ Hz, 1H), 6.03 (d, $J = 8.0$ Hz, 1H),

3.99–3.83 (m, 2H), 2.55 (q, $J = 7.1$ Hz, 4H), 2.48 (t, $J = 7.1$ Hz, 2H), 1.84 (dt, $J = 15.0, 7.6$ Hz, 2H), 1.63–1.52 (m, 2H), 1.12–0.97 (m, 6H). ^{13}C NMR ($CDCl_3$) δ 154.4, 151.6, 141.2, 139.7, 137.2, 135.1, 132.7, 128.7, 128.1, 127.9, 126.7, 126.6, 123.8, 121.9, 114.5, 101.2, 52.4, 52.1, 46.8, 26.4, 24.4, 11.6. HRMS (ESI) m/z calcd for $[C_{29}H_{33}ClN_3]^+$, 458.2358 $[M + H]^+$; found, 458.2360.

(E)-N-(1-(3-(Diethylamino)propyl)-7-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5s). According to general procedure B, the product was obtained as a yellow solid (34 mg, 4%). Mp 97–99 °C. 1H NMR ($CDCl_3$) δ 9.28 (d, $J = 8.8$ Hz, 1H), 7.98 (s, 1H), 7.77–7.71 (m, 2H), 7.63–7.58 (m, 4H), 7.47–7.38 (m, 2H), 7.36–7.31 (m, 3H), 6.63 (d, $J = 7.2$ Hz, 1H), 4.43 (t, $J = 6.8$ Hz, 2H), 2.68–2.67 (m, 4H), 2.61 (t, $J = 6.4$ Hz, 2H), 2.08 (t, $J = 6.4$ Hz, 2H), 1.09–1.05 (m, 6H). ^{13}C NMR ($CDCl_3$) δ 154.8, 146.4, 140.1, 139.6, 138.0, 129.0, 128.4, 127.8, 126.8, 124.9, 124.3, 122.8, 121.8, 114.2, 101.6, 52.1, 49.0, 46.7, 26.1, 10.3. HRMS (ESI) m/z calcd for $[C_{29}H_{31}F_3N_3]^+$, 478.2465 $[M + H]^+$; found, 478.2470.

(E)-N-(1-(4-(Diethylamino)butyl)-7-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5t). According to general procedure B, the product was obtained as a yellow solid (74 mg, 15%). Mp 135–137 °C. 1H NMR ($CDCl_3$) δ 8.74 (d, $J = 8.7$ Hz, 1H), 7.79–7.57 (m, 4H), 7.55–7.49 (m, 2H), 7.45 (t, $J = 7.6$ Hz, 2H), 7.33 (t, $J = 6.8$ Hz, 1H), 7.03 (d, $J = 8.3$ Hz, 2H), 6.96 (d, $J = 8.0$ Hz, 1H), 6.05 (d, $J = 8.0$ Hz, 1H), 3.97 (t, $J = 7.3$ Hz, 2H), 2.54 (dd, $J = 14.2, 7.1$ Hz, 4H), 2.48 (t, $J = 7.1$ Hz, 2H), 1.85 (dt, $J = 14.6, 7.3$ Hz, 2H), 1.59 (dd, $J = 14.2, 7.1$ Hz, 2H), 1.04 (t, $J = 7.1$ Hz, 6H). ^{13}C NMR ($CDCl_3$) δ 154.1, 151.9, 141.2, 139.8, 138.5, 135.2, 132.7, 128.7, 128.0, 127.6, 126.7, 126.6, 125.3, 121.6, 119.4, 111.7, 101.6, 52.4, 52.1, 46.8, 26.5, 24.5, 11.5. HRMS (ESI) m/z calcd for $[C_{30}H_{33}F_3N_3]^+$, 492.2621 $[M + H]^+$; found, 492.2629.

(E)-N-(6-Chloro-1-(3-(piperidin-1-yl)propyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5u). According to general procedure A, the product was obtained as a yellow solid (128 mg, 43%). Mp 48–50 °C. 1H NMR ($CDCl_3$) δ 8.66 (s, 1H), 7.70–7.52 (m, 4H), 7.53–7.40 (m, 3H), 7.38–7.24 (m, 2H), 7.12–6.98 (m, 3H), 6.03 (d, $J = 7.8$ Hz, 1H), 4.02 (t, $J = 6.4$ Hz, 2H), 2.36 (br, 4H), 2.30 (t, $J = 6.4$ Hz, 2H), 1.93 (dd, $J = 12.6, 6.4$ Hz, 2H), 1.67–1.53 (m, 4H), 1.47 (br, 2H). ^{13}C NMR ($CDCl_3$) δ 154.2, 151.5, 141.2, 140.2, 137.5, 135.1, 131.0, 129.4, 128.7, 127.9, 126.7, 126.6, 126.5, 125.8, 121.9, 116.4, 100.6, 54.8, 54.5, 49.8, 26.1, 25.6, 24.4. HRMS (ESI) m/z calcd for $[C_{29}H_{31}ClN_3]^+$, 456.2201 $[M + H]^+$; found, 456.2207.

(E)-N-(1-(3-(Piperidin-1-yl)propyl)-6-(trifluoromethyl)quinolin-4(1H)-ylidene)biphenyl-4-amine (5v). According to general procedure A, the product was obtained as a yellow solid (47 mg, 12%). Mp 85–87 °C. 1H NMR ($CDCl_3$) δ 8.93 (s, 1H), 7.74 (d, $J = 9.0$ Hz, 1H), 7.70–7.56 (m, 4H), 7.52–7.39 (m, 3H), 7.39–7.26 (m, 1H), 7.04 (t, $J = 7.2$ Hz, 3H), 6.05 (d, $J = 8.0$ Hz, 1H), 4.04 (t, $J = 6.5$ Hz, 2H), 2.37 (br, 4H), 2.32 (t, $J = 6.5$ Hz, 2H), 2.02–1.88 (m, 2H), 1.66–1.54 (m, 4H), 1.47 (br, 2H). ^{13}C NMR ($CDCl_3$) δ 154.2, 151.8, 141.2, 140.2, 135.1, 128.7, 128.0, 126.7, 126.6, 125.5, 125.3, 124.2, 121.7, 115.2, 101.6, 54.8, 54.5, 49.7, 26.1, 25.4, 24.4. HRMS (ESI) m/z calcd for $[C_{30}H_{31}F_3N_3]^+$, 490.2465 $[M + H]^+$; found, 490.2473.

In Vitro Cytotoxicity. The cytotoxicity was assessed using general cell viability end point MTT (3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide). Briefly, the day before experiment, cells HEK293T (human embryonic kidney epithelial cell line, ATCC CRL-11268) were seeded in 96-well tissue culture plates, in RPMI 1640 culture medium supplemented with 10% fetal serum bovine, 100 units of penicillin G (sodium salt), 100 μ g/mL streptomycin sulfate, and 2 mM L-glutamine, at a concentration that allowed cells to grow exponentially during the time of the assay. Compounds to be tested were diluted in DMSO and then serially diluted in the culture medium. Compounds at different concentrations and DMSO were then added to the cells, which were incubated at 37 °C in humidified 5% CO_2 atmosphere. After 48 h, cell medium containing DMSO (for control cells) or tested compound solution (for test cells) was removed and replaced with fresh medium containing MTT dye. After 3 h of incubation the complete medium was removed and the intracellular formazan crystals were solubilized and extracted with DMSO. After 15 min at room temperature the absorbance was measured at 570 nm in a

microplate reader. The percentage of cell viability was determined for each concentration of tested compound as described previously. The concentration of a compound reflecting a 50% inhibition of cell viability (i.e., EC_{50}) was determined from the concentration–response curve, applying a nonlinear regression procedure to the concentration response data using GraphPad Prism software.

In Vitro Blood Stage Activity Assays. Human red blood cells infected with 1% ring-stage *P. falciparum* strains synchronized with 5% sorbitol were incubated with tested compounds in 96-well plates at 37 °C for 48 h in RPMI-1640 medium, supplemented with 25 mM HEPES, pH 7.4, 10% heat inactivated human serum (or 0.5% AlbuMAX, 2% human serum), and 100 μ M hypoxanthine under an atmosphere of 3% O₂, 5% CO₂, 91% N₂. After 48 h, the cells were fixed in 2% formaldehyde in PBS, transferred into PBS with 100 mM NH₄Cl, 0.1% Triton X-100, 1 nM YOYO-1, and then analyzed in a flow cytometer (FACSsort, Beckton Dickinson; excitation 488 nm, emission 520 nm). Values of IC_{50} were calculated using GraphPad Prism software.

Hemozoin-like Crystals Growth Inhibition. Inhibition of hemozoin-like crystals formation by tested compounds was assessed using the previously described in vitro method.³⁶ In short, a hemozoin-like crystals stock suspension was sonicated for 3 min and diluted in fresh broth medium to a final concentration of 2 μ M (heme equivalents) in the wells of a 96-well plate. Stock solutions of tested compounds were prepared at 25 mM in DMSO. Stock solutions of chloroquine (positive control) and gentamicin (negative control) were prepared at 100 mM in distilled water, and all were 0.22- μ m-filtered previous to being diluted to 0–1000 μ M final concentration in the wells. Plates were incubated at 37 °C in a 5% CO₂ atmosphere for 5 days to observe the presence or absence of crystal growth. All tests were performed in triplicate.

Growth Inhibition against Transgenic *P. falciparum*. 3D7-yDHOD-GFP, a transgenic derivative of *P. falciparum* 3D7 containing yeast dihydroorotate dehydrogenase (DHODH), was generated via the electroporation of purified pHHyDHOD-GFP plasmid into ring stages of *P. falciparum* using a Bio-Rad GenePulser as described previously.⁴⁶ *P. falciparum* cell extracts and measurement of cytochrome *c* reductase activity was performed as described previously.^{41,48}

In Vitro Liver Stage Activity Assays. Huh-7 cells, a human hepatoma cell line, were cultured in 1640 RPMI medium supplemented with 10% v/v fetal calf serum (FCS), 1% v/v nonessential amino acids, 1% v/v penicillin/streptomycin, 1% v/v glutamine, and 10 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES), pH 7, and maintained at 37 °C with 5% CO₂. Inhibition of liver stage infection was determined by measuring the luminescence intensity in Huh-7 cells infected with a firefly luciferase-expressing *P. berghei* line, PbGFP-Luc_{con}, as previously described.⁴⁷ Briefly, cells (12 \times 10³/well) were seeded in 96-well plates the day before drug treatment and infection. Tested compounds were prepared in the following way: 10 mM stock solutions were obtained by dissolving accurately weighed compounds in MeOH and dilutions subsequently made with medium to the desired concentration. Medium was replaced by fresh medium containing the appropriate concentration of each compound 1 h prior to infection. Sporozoites (10 000 spz/well), freshly obtained through disruption of salivary glands of infected female *Anopheles stephensi* mosquitos, were added to the wells 1 h after compound addition. Sporozoite addition was followed by centrifugation at 1700g for 5 min. At 24 h after infection, the medium was once more replaced by fresh medium containing the appropriate concentration of each compound. Inhibition of parasite development in the liver was measured 48 h after infection. The effect of the compounds on the viability of Huh-7 cells was assessed by the AlamarBlue assay (Invitrogen, U.K.), using the manufacturer's protocol. Nonlinear regression analysis was used to fit the normalized results of the dose–response curves, and IC_{50} values were determined using the SigmaPlot software.

Rat Microsomes Stability Assay. The viability of the rat microsomes was initially accessed by evaluating their CYP2E1-catalyzed *p*-nitrophenol hydroxylation capacity, applying a method-

ology described elsewhere.⁴⁹ Two parallel determinations in rat microsomes, with and without NADPH regenerating system, were performed for selected compounds. Compounds were incubated at 37 °C at 1 μ M in rat or mouse pooled liver microsomes (0.2 mg of protein/mL, BD Gentest, Discovery Labware Inc., Woburn, MA), suspended in 10 mM phosphate buffer solution (pH 7.4 at 25 °C). Metabolic reactions were initiated by the addition of a NADPH regenerating system A (31 mM NADP⁺, 66 mM glucose 6-phosphate, and 0.67 mM MgCl₂) and NADPH regenerating system B (BD Gentest, containing 40 U/mL glucose 6-phosphate dehydrogenase in 5 mM sodium citrate). At appropriate intervals, aliquots were removed and added to acetonitrile. The samples were mixed and centrifuged for 10 min at room temperature (Σ , 112g–4000g), and the supernatant fraction was analyzed by HPLC. Measurements were carried out in a LabChrom chromatograph coupled to a Merck Hitachi L-7100 pump, a Shimadzu SPD-SAV UV/vis spectrophotometric detector, and a Merck Hitachi 2500 integrator. A LichroCARTRP-18 (5 μ m, 250–254 mm) analytical column (Merck) was used with an acetonitrile/H₂O (25:75, v/v) eluent system. Elution was performed at a solvent flow rate of 1 mL/min. In the assay without NADPH generating system, 24 μ L of purified water was used instead of solutions A and B.

■ ASSOCIATED CONTENT

● Supporting Information

General procedure for 6- and 7-substituted quinolin-4-amines, structural data of intermediates 7a–j, X-ray analysis of compound 5c, NMR spectra of compound 5o, HPLC conditions and retention times of final compounds 5a–v, and LC–MS analysis of microsomal incubation mixtures of compound 5u. This material is available free of charge via the Internet at <http://pubs.acs.org>.

■ AUTHOR INFORMATION

Corresponding Author

*Phone: +351 217946400. E-mail: aressurreicao@ff.ul.pt.

Author Contributions

[∞]A.S.R. and D.G. contributed equally.

Notes

The authors declare no competing financial interest.

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■ ABBREVIATIONS USED

CQ, chloroquine; PQ, primaquine; DV, digestive vacuole; NMR, nuclear magnetic resonance; TEA, triethylamine; DEA, diethylamine; rt, room temperature; o/n, overnight; DMF, *N,N*-dimethylformamide; THF, tetrahydrofuran; DCM, dichloromethane; DMSO, dimethylsulfoxide; mtETC, mitochondrial electron transport chain; DHODH, dihydroorotate dehydrogenase; HPLC, high performance liquid chromatog-

raphy; LC–MS, liquid chromatography coupled to mass spectrometry

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