Structure-Activity Relationship Studies Reveal New Astemizole Analogues Active against *Plasmodium falciparum* In Vitro

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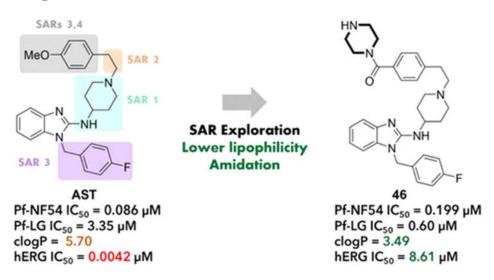
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Graphical abstract



Abstract

In the context of drug repositioning and expanding the existing structure-activity relationship around astemizole (AST), a new series of analogues were designed, synthesized, and evaluated for their antiplasmodium activity. Among 46 analogues tested, compounds **21**, **30**, and **33** displayed high activities against asexual blood stage parasites ($PfNF54 IC_{50} = 0.025-0.043 \mu M$), whereas amide compound **46** additionally showed activity against late-stage gametocytes (stage IV/V; $PfLG IC_{50} = 0.6 \pm 0.1 \mu M$) and 860-fold higher selectivity over hERG (**46**, SI = 43) compared to AST. Several analogues displaying high solubility (Sol > 100 μM) and low cytoxicity in the Chinese hamster ovary (SI > 148) cell line have also been identified.

KEYWORDS: Astemizole; *Plasmodium falciparum*; repositioning; human ether-a-go-go-related gene

Malaria is a life-threatening disease caused by *Plasmodium* parasites transmitted to humans via *Plasmodium*-infected *Anopheles* mosquitoes during a blood meal. Among the five human malaria species known, *P. falciparum* (*Pf*) is the most virulent and highly prevalent in the African region. ^(1–3) According to the 2020 WHO World Malaria Report, an estimated 229 million cases of malaria were recorded worldwide in 2019, with the sub-Saharan African region bearing the largest burden, accounting for 94% of the global incidence rate. ⁽³⁾ During that year, 409,000 deaths occurred, among which 67% were children under the age of 5, ⁽³⁾ maintaining the death rate of "one-child every 2 min" statistic.

The use of antimalarial drugs forms the cornerstone of both the prevention and treatment of the disease. Currently, the WHO recommends the use of the artemisinin-based combination therapy (ACT) regimen, which has provided both treatment and protection to millions over the last few decades. ^(4,5) However, the continuous evidence of the emergence of resistance in countries of the Greater Mekong Subregion (GMS) and more recently in the African Great Lakes region, particularly Rwanda, poses a serious threat to the current regimen. ^(6,7) Therefore, the need to utilize various approaches to discover and develop new and structurally diverse antimalarials that offer both chemoprotection and prevent reinfection via multistage activity is warranted.

Chong and co-workers identified the antimalarial properties of the antihistamine drug astemizole (AST) and its principal metabolite desmethylastemizole (DM-AST) via a high-throughput screen (HTS) of 2687 existing drugs. Both AST and DM-AST showed growth inhibition of chloroquine-sensitive (CQ-S) and multi-drug-resistant (MDR) malaria parasites, which translated to parasitemia reduction in two in vivo mouse models of malaria. Furthermore, preliminary mode of action studies revealed that AST exhibits its antimalarial action by accumulating in the Pf food vacuole, thereby disrupting heme crystallization (8) However, the presence of additional mechanism(s) was not ruled out. AST has also previously been shown to possess liver-stage antimalarial activity (IC50 = 0.114 μ M) from a HTS screen of 5375 known drugs with diverse modes of actions using P. berghei (Pb) parasites. (9)

Notably, various researchers have used AST as a template to derive analogues with varying degrees of antiplasmodium activity. Musonda and co-workers explored the potential to

overcome Pf resistance to chloroquine (CQ) via hybridization with the AST pharmacophore. Front-runner compounds from this work demonstrated improved in vitro activity against the K1 CQ-R strain of Pf (IC₅₀ = 0.037–0.610 μ M) as well as high in vivo efficacy in the Pb mouse infection model of malaria. (10) Additionally, two recent publications separately conducted brief antiplasmodium structure–activity relationship (SAR) studies based on AST. (11,12) The summary of SAR studies reported by De Jonghe and co-workers showed that the most potent analogues from their series also demonstrated reduced hERG channel inhibition activity (SI = 110). (12) However, the cytotoxicity of these analogues was not reported, and the poor drug likeness evidenced by high lipophilicity (clogP = 7.7) and low solubility of the front-runner compound would present challenges in development.

Our group previously showed the potential of AST and its analogues to display multistage activity against *Plasmodium* parasites following their evaluation against asexual blood stage ($PfNF54\ IC_{50} = 0.033-1.9\ \mu M$), liver stage ($Pb\ IC_{50} = 0.210\ \mu M$), and late-stage gametocytes (stage IV/V; $PfLG\ IC_{50} = 1.9-4.1\ \mu M$). Additionally, we showed intracellular inhibition of hemozoin formation by AST analogues within the parasite, which supported Chong and coworkers' initial findings of AST's interference with the heme detoxification pathway. (13) These results justified further investigation of the AST scaffold toward its optimization for malaria. In the present work, several structural analogues of AST were designed with the goal of improving antiplasmodium activity and enhancing solubility while utilizing known strategies to circumvent the human ether-a-go-go-related gene (hERG) potassium (K⁺) channel inhibition liability associated with AST.

The SAR buildup involved modifications of various parts of AST categorized in four regions (Figure 1). Among the strategies to reduce hERG affinity and improve aqueous solubility, we utilized reducing lipophilicity (clogP), removal of aromaticity, attenuating the pK_a of the basic nitrogen, as well as discrete modifications around the AST scaffold.

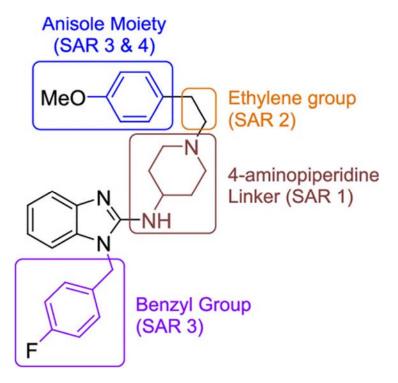


Figure 1. Summary of designed SAR around AST.

SAR 1 involved the replacement of the 4-aminopiperidine linker with open chain diamine alkyl moieties (Figure 1), whereas SAR 2 was designed to reduce the pK_a of the basic piperidine nitrogen via incorporation of a carbonyl (C=O) group at the α - and β -position on the ethylene group. SAR 3 was focused on replacing the 4-OMe and 4-F groups of the anisole and benzyl groups, respectively, with representative substituents from each of the four quadrants of the Craig plot. In the benzyl moiety, truncation, replacement with methyl, incorporation of heteroatom (N), methylation of the methylene bridge, and CN/F disubstitution were also explored in SAR 3. Toward reducing hERG affinity, replacement of the 4-OMe group in the anisole moiety with amides was explored in SAR 4 (Figure 1). Similarly, the anisole group was replaced with both aliphatic and aromatic heterocycles (SAR 4) to expand the SAR and interrogate the significance of the phenyl group.

The synthesis of SAR 1 and 2 analogues commenced with the coupling of commercially available 2-chloro-1-(4-fluorobenzyl)-1*H*-benzo[*d*]imidazole (Scheme 1) with various diamines, resulting in intermediates **1a**–**c** and **5**. In SAR 1, nucleophilic substitution reactions between amines **1a**–**c** and 4-methoxyphenethyl bromide in the presence of triethylamine (NEt₃) in *N*,*N*-dimethylformamide (DMF) afforded target compounds **2**–**4**. For SAR 2, deprotection of **5** followed by coupling of the resulting free amine (**6**) to commercially sourced 2-bromo-1-(4-methoxyphenyl)ethan-1-one and 2-(4-methoxyphenyl)acetic acid resulted in compounds **7** and **8**, respectively. (13)

Scheme 1.

^aReagents and conditions: (a) ethane-1,2-diamine (for **1a**) or propane-1,3-diamine (for **1b**), 120 °C, 8–12 h (**1a**; 85%, and **1b**; 92%); 1,3-diaminopropan-2-ol, EtOH, 79 °C, 96 h (**1c**; 50%); (b) 4-methoxyphenethyl bromide, Et₃N, DMF, 22 °C, 10 h (15–62%); (c) ethyl 4-aminopiperidine-1-carboxylate, 170 °C, 4 h (98%); (d) 48% aq HBr, 120 °C, 3 h (85%); (e) 2-bromo-1-(4-methoxyphenyl)ethan-1-one, Et₃N, DCM, 22 °C (7; 10%); 2-(4-methoxyphenyl)acetic acid, T₃P, *i*-Pr₂NEt, DMF, 0–22 °C, 3 h (**8**; 55%).

Compounds containing modifications in the benzyl group (12–26, SAR 3) and the anisole moiety (27–53, SAR 4) were synthesized, as shown in Scheme 2. The synthesis of respective

alkylating agents (9a-g) required to deliver target compounds in Scheme 2 is described in Supporting Information (Scheme S1).

Scheme 2.

^aReagents and conditions: (a) (i) alkylating agent, K₂CO₃, acetone, 18–23 °C, 2–5 h (60–98%); (ii) ethyl 4-aminopiperidine-1-carboxylate or *tert*-butyl 4-aminopiperidine-1-carboxylate, Et₃N, 150 °C, 2–12 h (58–89%); (b) 48% aq HBr, 120 °C, 3 h or TFA, DCM, 20 °C, 3 h, Amberlyst-A21, DCM, 3 h (58–98%); (c) **9a** (for **12**–**14**), or alkylating agent (for **27–30**) or **9c** (for **31**) or **9d** (for **33** and **34**), K₂CO₃, MeCN, 80–85 °C (32–89%); (d) **9f** (for **23**) or respective alkylating agent, K₂CO₃, DMF, 70 °C (73–93%); (e) (i) 2 M KOH, MeOH, 80 °C, 4 h; (ii) 2 N HCl, 0–4 °C, pH 2 (90–96%); (f) 2-(piperazinyl)ethanol (for **39**) and **9g** (for **40**), MsCl, Et₃N, DCM, 0–45 °C, 4 h (**39**: 65%; **40**: 46%); (g) amine, *i*-Pr₂NEt, T₃P, DCM, 23°C, 4–24 h (15–74%).

Briefly, alkylation of 2-chloro-1*H*-benzo[*d*]imidazole at N-1, followed by coupling with ethyl 4-aminopiperidine-1-carboxylate or *tert*-butyl 4-aminopiperidine-1-carboxylate furnished intermediates **10a–c**. Following acidic deprotection, free amines (**11a–c**) were coupled with 4-(2-bromoethyl)benzonitrile (**9a**, Scheme S1) to afford **12–14** in good yields. ⁽¹³⁾ Treatment of **12** with various benzyl bromides or bromomethylpyridines afforded the target compounds **15–17** and **19–26** (Scheme 2A), ⁽¹¹⁾ whereas intermediates **27–31** and **33–40** (Scheme 2C) were accessed using **6** or **11c** via nucleophilic substitution (S_N2) with various commercially available or previously synthesized alkylating agents (**9c,d** and **9f,g**, Scheme S1). Carboxylic acid derivatives **18** and **32** were prepared via hydrolysis of methyl esters **17** and **31**, respectively, whereas piperazine compound **41** was obtained from **40** by TFA N-Boc deprotection (Scheme 2A–C).

Compounds 43–53 were obtained via S_N2 coupling of either 6 or 11c with amide derivatives (42a–j) of 4-(2-bromoethyl)benzoic acid prepared using propanephosphonic acid anhydride (T₃P)-mediated acid–amine coupling (Scheme 2D).

All synthesized compounds were evaluated for their solubility and antiplasmodium activity against the chloroquine-sensitive (NF54) strain of Pf. Furthermore, selected compounds with IC₅₀ < 0.700 μ M against PfNF54 were evaluated against the PfK1 (MDR) strain to assess potential for cross-resistance with known antimalarials (i.e., CQ). We previously reported AST analogues exhibiting signatures attributable to possible interaction with the parasitic heme detoxification machinery; (13) for this reason, a representative set of analogues from this series were also assessed for their ability to block β -hematin (β -H) formation, as a surrogate for the inhibition of heme detoxification by the parasite.

Potency was reduced by 4–8-fold when the 4-aminopiperidine group in AST was replaced with an open chain diamine (compounds 2–4; Table 1) in SAR 1.

We next investigated the effect of reducing the basicity of piperidine nitrogen. Activity was lost in both β -keto and amide analogues 7 ($PfNF54\ IC_{50} = 3.18\ \mu M$, Table 1) and 8 ($PfNF54\ IC_{50} = 2.89\ \mu M$) compared to AST. This observation was consistent with previously reported results by De Jonghe and co-workers, which showed loss of activity in urea and sulfonamide analogues. (12) This may suggest the significance of piperidine nitrogen basicity toward antiplasmodium activity. Solubility was generally high in all open chain amine-containing analogues (2–4) and acylated compound 7 (Sol > 190 μM). However, amide derivative 8 displayed significantly lower solubility (Sol = 60 μM) albeit higher than that of AST. All analogues in this SAR exhibited low βH inhibition (IC₅₀ > 1000 μM) with the exception of hydroxy compound 4 (IC₅₀ = 94.6 μM , Table S1, Supporting Information).

Table 1. In Vitro Antiplasmodium Activity and Solubility of AST Analogues in SARs 1 and 2 (ND, Not Determined)

Compound	X	Y	Pf IC ₅₀ (μM) ^{a,b} NF54 K1		RI^c	Sol. ^d (μM)
AST	, M,	`	0.086	0.370	4.30	40
2	, N _ H		0.691	0.787	1.14	190
3	`.NN		0.330	0.677	2.05	200
4) H OH H		0.706	1.406	3.46	200
7	N'		3.180	ND	-	200
8	, N		2.890	ND	-	60

^aMean from $n \ge 2$ independent experiments with sensitive (NF54) and multi-drug-resistant (K1) strains of Pf. ^bChloroquine (NF54 IC₅₀ = 0.004 μM; K1 IC₅₀ = 0.14 μM) was used as a reference drug for activity. ^cResistance index = [PfK1 IC₅₀/PfNF54 IC₅₀]. ^dSolubility, determined using turbidimetric method at pH 7.4. Hydrocortisone (>200 μM) and reserpine (<10 μM) were used as control drugs.

For the exploration of the benzyl group SAR (SAR 3), the 4-aminopiperidine was retained due to its significance toward potency. However, the 4-OMe group in the anisole moiety was replaced with the 4-CN group (Table 2). This switch was driven by the lipophilicity-lowering effects and resilience toward metabolism associated with the cyano group compared to methoxy, (14,15) There was a general loss in antiplasmodium potency across the SAR, with only three analogues (24: $PfNF54 \text{ IC}_{50} = 0.071 \mu\text{M}$; 21: $PfNF54 \text{ IC}_{50} = 0.025 \mu\text{M}$; and 25, $PfNF54 \text{ IC}_{50} = 0.108 \mu\text{M}$) out of 15, showing activity higher or comparable to that of AST. Removal of the benzyl group resulted in \sim 7-fold loss of activity (12: $PfNF54 \text{ IC}_{50} = 0.579$ uM) compared to that of AST, whereas replacement with a methyl group significantly diminished the activity (13: $PfNF54 \text{ IC}_{50} = 5.08 \mu\text{M}$). The weak activity of 12 was found to contradict that of a previously reported debenzylated analogue (compound 5c: IC₅₀ = 0.047 μM, Kumar and co-workers) (13) in which removal of the benzyl group with retention of the 4-OMe group in the anisole moiety increased the potency by 2-fold when compared to AST. This observation may, however, suggest the possibility to significantly optimize antiplasmodium potency in debenzylated analogues by exploring substitutions for the 4-OMe group in the anisole moiety.

Table 2. In Vitro Antiplasmodium and Solubility of AST Analogues in SAR 3 (ND, Not Determined)

Compound	R	R_1	Pf IC ₅₀ NF54	(μM) ^{a,b} K1	RIc	Sol ^d (µM)
12	Н	-	0.579	ND	-	120
13	Me	-	5.080	ND	-	100
14		CN	0.156	0.176	1.1	90
15		OMe	0.624	ND	-	50
16		SO ₂ Me	0.485	ND	-	40
17	`————R ₁	CO_2Me	0.591	ND	-	120
18		CO ₂ H	>6	ND	-	>200
19		CF ₃	0.253	0.384	1.5	20
20		CH ₃	0.565	ND	-	80
21	N -	CF ₃	0.025	0.082	3.3	20
22	R ₁	CH ₃	0.261	ND	-	50
23		-	0.156	0.157	1.0	120
24	CN E		0.071	0.211	2.9	40
25	E_CN		0.108	ND	-	20
26	NC F		0.185	4.013	21.7	50

^aMean from $n \ge 2$ independent experiments with sensitive (NF54) and multi-drug-resistant (K1) strains of *Pf*. ^bChloroquine (NF54 IC₅₀ = 0.004 μM; K1 IC₅₀ = 0.14 μM) was used as a reference drug for activity. ^cResistance index = [*Pf*K1 IC₅₀/*Pf*NF54 IC₅₀]. ^dSolubility, determined using turbidimetric method at pH 7.4. Hydrocortisone (>200 μM) and reserpine (<10 μM) were used as control drugs.

CN/F disubstitution in the benzyl group was generally tolerated, although potency was lowered by ~2-fold when the CN group was in the *ortho*-position (**26**, *Pf*NF54 IC₅₀ = 0.185 μ M) compared to that of *ortho*-fluorine (**25**, *Pf*NF54 IC₅₀ = 0.108 μ M). However, CN/F disubstitution preferentially favored *para*-CN and *meta*-F regioisomerism (**24**, *Pf*NF54 IC₅₀ = 0.071 μ M, Table 2). Craig plot substitution tends to generally favor electron-withdrawing groups (EWGs) for antiplasmodium potency. However, among the groups tested, only the hydrophilic CN (**14**; *Pf*NF54 IC₅₀ < 0.156 μ M) and hydrophobic CF₃ (**19**; *Pf*NF54 IC₅₀ < 0.253 μ M) groups are tolerated. Pyridyl analogues **21** and **22** were more potent than their corresponding phenyl (**19** and **20**) match pairs, with the 4-trifluoromethyl-3-pyridyl analogue

(21, PfNF54 IC₅₀ = 0.025 μ M) showing the highest potency in the series (Table 2). With the exception of 26 (PfK1 IC₅₀ = 4.0 μ M, RI = 21.7), PfK1 data suggested the absence of cross-resistance with existing antimalarial drugs reflected in the low resistance indices (SI < 5).

Consistently, analogues bearing water-solubilizing groups in SAR 3 (17 and 18) or devoid of the benzyl aromatic ring (12 and 13) exhibited moderate to high solubility (100–200 μ M) compared to those containing lipophilic groups such as the F group (19, 21, and 24–26; Sol < 100 μ M). Methylation of the benzyl methylene linker was designed to discourage aromatic π – π interactions as a strategy to improve solubility. This effect was indeed observed as the solubility of 23 (Sol = 120 μ M) increased compared to that with 14 (Sol = 90 μ M) with retention of potency (PfNF54 IC₅₀ = 0.156 μ M). Three out of eight tested compounds (14, 19, and 25) displayed β -H inhibition activities of IC₅₀ < 100 μ M (Table S1).

We next explored various substitutions for the 4-OMe group in the anisole moiety (Table 3). Similar to benzyl group SAR (SAR 3), antiplasmodium activity is generally favored by substitution with EWGs. This is exemplified by the enhanced potency of CF₃-substituted **30** ($PfNF54 IC_{50} = 0.035 \mu M$). Notably, some EWGs produce activities superior to those of others. For instance, ester analogues **33** ($PfNF54 IC_{50} = 0.043 \mu M$) and **34** ($PfNF54 IC_{50} = 0.067 \mu M$) showed higher activity than did the methyl ester analogue **31** ($PfNF54 IC_{50} = 0.204 \mu M$). On the other hand, activity is lost in the carboxylic acid derivative (**32**, $PfNF54 IC_{50} > 6 \mu M$). Furthermore, ethyl ester **33** bearing a 4-CN-benzyl group retains comparable potency in the MDR strain PfK1 (RI = 1.21) compared to the 4-F-benzyl match pair **34**, in which the resistance index is higher (RI = 2.83). The low solubility and susceptibility of the ester structural motif to metabolism (Table 3) pre-empted the further exploration of bioisosteres.

Amides are utilized as ester group bioisosteres; additionally, amidation is a known strategy for reducing hERG affinity and improving aqueous solubility. (16,17) Therefore, we accordingly prepared and evaluated amides at the 4-position in the lateral phenyl group while maintaining either a 4-F or 4-CN group in the benzyl portion (Table 3). However, these AST amide analogues 43–53 produced lower activities ($PfNF54 IC_{50} = 0.199-1.91 \mu M$) compared to those of the ester match pairs (33 and 34) and AST, although the desired increase in solubility was achieved in some of the amides, especially those that contained water-solubilizing groups (43–46).

Table 3. In Vitro Antiplasmodium and solubility of AST Analogues in SAR 2 (ND, Not Determined)

		- (N N N	\sim_{R_1}			
Compound	R	R-()-	— ¡¡		(μΜ) ^{a,b} Κ1	RI ^c	Sol. ^d (µM)
27			Н	0.163	0.170	1.04	40
28			SO ₂ Me	0.238	4.592		80
29		{	CH ₃	0.357	ND	-	40
30	CN		CF ₃	0.035	0.103	2.94	20
31			CO ₂ Me	0.204	ND	-	80
32			CO ₂ H	>6	ND	-	120
33			CO ₂ Et	0.043	0.052	1.21	40
34	F		CO ₂ Et	0.067	0.19	2.83	65
35		NN	-	4.152	ND	-	20
36		N	-	1.386	ND	-	20
37		N	-	1.248	ND	-	<5
38	CN	N_O	-	>6	ND	-	10
39		N_N-8 0	-	>6	ND	-	40
40		NN-Boc	-	1.542	ND	-	<10
41		N_NH	-	1.677	ND	-	20
43			N\$—ОН	0.640	ND	-	150
44	F	R ₂	NOH	1.810	ND	-	200
45			NOH	1.613	ND	-	72
46			NNH	0.199	1.136	5.71	200
47			NH	0.318	ND		30

N_NH	0.199	1.136	5.71	200
NH	0.318	ND	-	30
N	1.912	ND	-	ND
N—<	0.295	ND	-	<5
NH	0.314	ND	-	ND
NH OH	0.759	ND	-	50
NH O-	0.755	ND	-	100
N	0.249	ND	-	5

^aMean from $n \ge 2$ independent experiments with sensitive (NF54) and multi-drug-resistant (K1) strains of Pf. ^bChloroquine (NF54 IC₅₀ = 0.004 μM; K1 IC₅₀ = 0.14 μM) was used as a reference drug for activity. ^cResistance index = [PfK1 IC₅₀/PfNF54 IC₅₀]. ^dSolubility, determined using turbidimetric method at pH 7.4. Hydrocortisone (>200 μM) and reserpine (<10 μM) were used as control drugs.

Notably, secondary amides (47, 49–52) generally displayed activity higher than that of tertiary amides (43–45 and 48), although as a caveat, some of these tertiary amides (43–45) contain other structural features that may contribute to low activity. The 3-hydroxyazetidine amide (43, PfNF54 IC₅₀ = 0.640 μ M) displayed activity more than 2-fold higher than that of the pyrrolidine (44, PfNF54 IC₅₀ = 1.81 μ M) and piperidine (45, PfNF54 IC₅₀ = 1.61 μ M) analogues, both of which had comparable activities. The phenyl moiety was found to be crucial for potency (27, PfNF54 IC₅₀ = 0.163 μ M) as its replacement with either aromatic or aliphatic heterocycles led to a significant loss in potency (35–41, PfNF54 IC₅₀ > 1.2 μ M, Table 3) compared to that of both 27 and AST. Following the assessment of representative compounds from each compound class, 4-CF₃30, 4-CO₂Et 34, and methyl amide 47 displayed high β -H inhibition, with IC₅₀ values of <100 μ M (Table S1). Most analogues in this SAR exhibited low to moderate solubility (Sol = 10–100 μ M), with some amides showing high solubility.

To assess potential for transmission blocking, a selected number of compounds across the four SARs were evaluated against early gametocyte (EG, stages I–III) and late gametocyte (LG, stage IV/V) stages in a dual-point screen at 1.0 and 5.0 μ M (Table S2 in the Supporting Information). Compounds **2**, **7**, and **8** displayed high activity (>70% at 5 μ M, or >50% at 1 μ M), with IC50 values in the low micromolar range and stage specificity toward EGs, defined as compounds that show a ≥2-fold change in dual-point values between EG and LG. However, these compounds generally exhibited loss of activity following IC50 determination, a phenomenon that is not surprising and may be attributable to confounding factors in the assay such as freeze—thawing cycles. Comparatively, AST, **4**, and **46** showed activity and stage specificity toward LG, with IC50 values lower than that of active compounds on EG. Of these active compounds against the LG, piperazine amide **46** exhibited the highest potency (*Pf*LG IC50 = 0.6 ± 0.1 μ M, Table S2), indicating it as a potential transmission-blocking candidate.

Representative compounds and those with antiplasmodium activity of PfNF54 IC₅₀ < 0.2 μ M were assessed for cytotoxicity in the Chinese hamster ovary (CHO) cell line. Accordingly, structurally diverse analogues with favorable antiplasmodium activity were evaluated for their hERG inhibition using potassium channels expressed in the CHO cell line on an automated Q-patch clamp platform (Table 4). Additionally, microsomal metabolic stability of compounds displaying low cytotoxicity (SI > 100) were evaluated in human, rat, and mouse liver microsomes (HLMs, RLMs, and MLMs, respectively, Table S3, Supporting Information).

Table 4. In Vitro Cytotoxicity (CHO), hERG Channel Activity, and Microsomal Metabolic Stability Data of Selected Analogues

	cytotox	icity (CHO)	a hERGb		
compound	IC ₅₀ (μM)	SIc	IC ₅₀ (μM)	SIc	clogPd
AST	29.6	344	0.0042	0.049	5.70
3			0.41	1.24	4.65
12			0.09	0.16	3.80
14			0.05	0.32	4.70
16			0.11	0.23	4.07
18			4.16	>0.7	4.45
21	>50	>2000	0.97	38.8	4.83
24	47.4	667	0.05	0.70	4.85
30	5.36	153			5.71
32	12.4	289			5.22
34	42.3	631			5.69
46	29.5	148	8.61	43.3	3.49
verapamil			0.2-0.8		
emetine	0.95				

^aCHO: Chinese hamster ovary epithelial cell line. Mean value from n=3 independent experiments. Emetine (IC₅₀ = 0.033 μM) was used as the reference drug. ^bhERG: Human ether-a-go-go-related gene. Mean value from n=3 independent experiments. Verapamil (IC₅₀ = 0.56 ± 0.096 μM) as a reference. ^cSI: Selectivity index = [CHO IC₅₀/PfNF54 IC₅₀] or [hERG IC₅₀/PfNF54 IC₅₀]. ^dCalculated using StarDrop software v6.5-1-1.

All tested compounds showed high selectivity (SI > 148) against CHO cells (Table 4). Compounds **21** (hERG IC₅₀ = 0.97 μ M) and **46** (hERG IC₅₀ = 8.61 μ M) displayed the lowest hERG channel inhibition activity with 202-fold and 1790-fold higher IC₅₀ values compared to those of AST (hERG IC₅₀ = 0.0042 μ M). However, the low hERG selectivity (SI < 43.3) of compounds remains a challenge that needs to be further addressed. Notably, carboxylic acid analogue **18** (hERG IC₅₀ = 4.16 μ M) also displayed low hERG activity potentially attributable to its zwitterionic furnishing properties; however, the presence of a carboxylic acid group is often associated with poor absorption and limited membrane permeability and may thus explain the equally observed weak whole-cell antiplasmodium activity (*Pf*NF54 IC₅₀ > 6 μ M).

Microsomal metabolic stability was generally low, especially in rodent species (M/RLM < 45% remaining and $CL_{int} > 67 \mu L \cdot min^{-1} \cdot mg^{-1}$, Table S2) compared to that of human-derived microsomes. Therefore, none of the compounds could be progressed for in vivo antimalarial evaluation in mice.

In conclusion, 46 analogues were synthesized and evaluated for their antiplasmodium activity and solubility. Overall, we highlight modifications that lead to improved antiplasmodium activity, solubility, and hERG channel inhibition activity. Cytotoxicity of tested compounds was low, although metabolic stability and marginal selectivity for antiplasmodium activity over hERG affinity still remains a challenge in the development of AST analogues as antimalarial agents. SAR explorations have demonstrated the significance of the 4-aminopiperidine linker and the phenethyl moiety and the importance of adequate basicity of the piperidine nitrogen. We further showed that F/CN disubstitution is tolerated in the benzyl moiety and that replacement of either 4-F or 4-OMe groups in AST with nitrile broadly produces equipotent analogues. Additionally, replacement of 4-F or 4-OMe groups in AST with EWGs such as CF₃ significantly improves the potency. From this study, incorporation of 4-CF₃-3-pyridyl moiety on the benzyl portion while retaining the 4-CN in the lateral phenyl group produced the most potent analogue (21, PfNF54 IC₅₀ = 0.025 μ M) with low hERG channel activity (hERG IC₅₀ = 0.97, SI = 38.8).

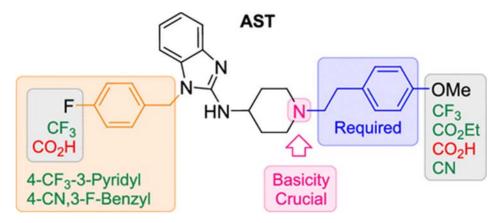


Figure 2. Summary of SAR trend around AST derived from this series.

Exploration of potential substituents at the 4-phenyl position of debenzylated analogue (12) is warranted, as these are envisaged to produce low molecular weight analogues with improved physicochemical and solubility profiles. Additionally, the potential for blood-stage/LG dual activity and the low hERG affinity of 46 presents an opportunity for further optimization. Furthermore, ester analogue 33 is amenable for exploration of other ester bioisosteric replacements to mitigate the underlying metabolic liability associated with the ester group.

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Abbreviations

AST astemizole

DMAST desmethylastemizole

hERG human ether-a-go-go-related gene

SI selectivity index

CQ chloroquine

AQ amodiaquine

CHO Chinese hamster ovary

HTS high-throughput screening

SAR structure–activity relationship

 $I_{\rm Kr}$ potassium ion current

ND not determined

IC₅₀ concentration of a drug that is required for 50% inhibition in vitro

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