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# Novel diaryl ureas with efficacy in a mouse model of malaria

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Traditional treatments for *Plasmodium falciparum* malaria such as chloroquine and pyrimethamine-sulfadoxine have in many areas succumbed to drug resistance, and evidence now suggests emerging resistance to the new first-line artemisinins in Western Cambodia.<sup>1</sup> Infection is rampant, resulting in ~650,000 deaths per year.<sup>2</sup> Clearly, a need exists for novel therapies that are orally bioavailable, economical, and safe and whose activities are not compromised by existing resistance.<sup>3-5</sup>

Our laboratories have been interested in fatty acid biosynthesis in both *Plasmodium spp.* and *Mycobacterium tuberculosis.* Initial investigations into the mycobacterial target of the frontline drug isoniazid (INH) led to the discovery that the INH-NAD adduct binds the enoyl acyl carrier protein reductase InhA (also known as FabI or ENR).<sup>6</sup> This sparked efforts by us and others to study triclosan-inspired small molecule inhibitors of InhA that do not require KatG activation of INH for *M. tuberculosis*,<sup>7</sup> and that inhibit the *P. falciparum* homolog (PfFabI, previously termed PfENR) in biochemical assays with purified enzyme.<sup>8-10</sup> Interestingly,

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the plasmodial FabI is not essential for the intra-erythrocytic stage of the parasite and instead is required for normal progression of the liver-stage of infection.<sup>11, 12</sup>

Efforts to optimize the antimalarial efficacy of small molecule triclosan analogs began with the hypothesis that the phenol could be replaced by organic functionality capable of maintaining some of the hydrogen-bonding interactions that have been shown crystallographically to be important for enzyme inhibition.<sup>13</sup> The key interactions in this case involve the PfFabI Tyr-277 phenol and the 2'-OH of the ribose unit of the bound NAD+ co-factor. The compounds (Table 1) were prepared via adaptation of standard synthetic methods common to diaryl ether assembly<sup>14</sup> and feature a variety of 1-substituents, ranging from an ether and a nitrile to carboxylic acid derivatives, amines, sulfonamides, and ureas. Triclosan methyl ether was prepared by methylation of the parent with methyl iodide in the presence of potassium carbonate to afford **1**. The synthesis of triclosan 1-substituted analogs relied significantly on the preparation of anilines 5a and 5b (Scheme 1). The anilines were synthesized beginning with the coupling of the respective o-chloronitrobenzene with 2.4dichlorophenol in DMSO. The resulting diaryl ethers underwent reduction to afford anilines 5a and b. Aniline 5a was carried in three straightforward steps, involving diazotization, the Sandmeyer reaction, and the Rosenmund-von Braun reaction,<sup>15</sup> to benzonitrile 2. The 1cyano triclosan 2 was then hydrolyzed to afford amide 4 and carboxylic acid 3. Reduction of 2 with lithium aluminum hydride afforded aminomethyl 6. Utilization of Ciganek's methylorganocerium reagent,<sup>16</sup> generated *in situ* from cerium(III) chloride and one equivalent of methyllithium reduced 2 to a,a-dimethylamine 7. Aniline 5a provided a starting point for the synthesis of a limited set of sulfonamide and amide analogs 8 - 11 via reaction with a sulfonyl chloride, anhydride or acid chloride under traditional conditions (Scheme 2). Two routes to ureas were devised, involving either direct reaction of aniline 5a or **5b** with a commercial isocyanate or activation with triphosgene followed by mixing with a commercial amine. Two simple urea analogs 12 and 13 were initially made.

While none of the phenol replacements displayed significant activity ( $IC_{50} > 10 \mu M$ ) in a PfFabI inhibition assay,<sup>13</sup> both aminomethyl **6** and diaryl urea **13** demonstrated efficacy in growth inhibition assays<sup>17</sup> with cultured 3D7 (drug-sensitive) and Dd2 (resistant to chloroquine and pyrimethamine-sulfadoxine) *P. falciparum* strains. Molecular modeling studies rationalized how the loss of hydrogen-bonding (perhaps through both donating and accepting) upon replacement of the 1-OH may lead to abrogation of potent binding to, and hence reduced inhibition of PfFabI. Chemical inspection of **6** and **13** led to the prioritization of the diaryl urea for optimization based on its superior whole-cell efficacy and the potential to readily explore a range of aryl substituents, not belonging to the diaryl ether subunit.

A set of follow-up diaryl ureas (Scheme 2; Table 2) was prepared to probe the structureactivity relationship pertinent to growth inhibition of *in vitro*-cultured *P. falciparum*, regardless of molecular target. It is clear from a select subset of analogs that the substitution pattern on the non-diaryl ether aryl moiety affected antimalarial activity against both strains of *P. falciparum*. In particular, the 3-position favored electron-withdrawing groups such as trifluoromethyl or cyano. The 4-position also preferred electron-withdrawing groups such as cyano, nitro, fluoro, and chloro. This was exemplified in the most potent compounds in this series: **18** (3-CF<sub>3</sub>-4-ClC<sub>6</sub>H<sub>3</sub>), **21** (3-CF<sub>3</sub>-4-NO<sub>2</sub>C<sub>6</sub>H<sub>3</sub>), and **24** (3,4-diCNC<sub>6</sub>H<sub>3</sub>).

The *in vivo* activity of a subset of the urea derivatives was assessed utilizing the *P. berghei* rodent malaria model (Table 2).<sup>18</sup> Briefly, ~5 weeks old CD-1 mice (Charles River Laboratories) were infected intraperitoneally with 10<sup>6</sup> *P. berghei* (KBG-173 line) parasitized red blood cells on day 0. Drug dosing (bid, divided dose) was initiated on day 3 and continued on days 4 and 5. Subcutaneous administration was achieved using a peanut oil suspension of the compound. Activity was defined by the fractional survival at day 31. The

infected control mice survived an average of 8 days whereas non-infected control mice survived the entire 31 days of the study. Compounds with comparatively lower *in vitro* activity, such as **14** – **17**, failed to exhibit an extension of survival beyond the control animals. **19**, exhibiting better whole-cell efficacy, allowed survival of 2 of 7 and 1 of 7 animals at 31 days post-infection, at the 128 mg/kg and 256 mg/kg doses respectively. Diaryl urea **18**, displaying the most potent whole-cell efficacy to date in this family, when dosed at 128 mg/kg, enabled extension of survival of the infected mice beyond the control. Promisingly, dosing at 256 mg/kg of **18** demonstrated 4 of 7 animals surviving 31 days post-infection. While many factors contribute to the *in vivo* efficacy of a small molecule antimalarial, it is clear that the diaryl ureas' ability to inhibit the growth of *P. falciparum in vitro* was correlated with their efficacy in an *in vivo P. berghei* mouse model of infection.

It is interesting to note that phenoxy-substituted ureas have been previously reported as potent inhibitors of *Plasmodium spp.*, based on solely in vitro data. These include the compound WR268961 (Supplementary Data Figure S1),<sup>19</sup> where the urea linkage is parawith respect to the oxygen of the diaryl ether unit instead of ortho- as in 13 - 25. WR268961 abrogated parasite growth with an EC<sub>50</sub> = 87 nM (W2 strain) and 460 nM (D6 strain), and modestly inhibited the *P. falciparum* cysteine protease plasmepsin 2 (PfPM2;  $IC_{50} = 17$ μM). The diaryl ureas presented herein, however, do not appear to significantly target the plasmepsins as they equally inhibit the growth of both wild type and knockouts of PfPM1 through 4 (See Supplementary Data Table S1), attained via a genetic disruption methodology in the Dd2 background.<sup>20</sup> GlaxoSmithKline disclosed the whole-cell efficacy of screening hit TCMDC-139010 (Supplementary Data Figure S1; XC<sub>50</sub> = 930 nM vs. 3D7 strain), but without information concerning the biochemical target.<sup>4</sup> We also reported in 2005 the preparation of triclosan-based 4'-ureas (Supplementary Data Figure S1) that were less potent against cultured parasite (EC<sub>50</sub> values of ~100  $\mu$ M) than 12 – 25, but exhibited IC<sub>50</sub> values of ~100 nM against purified PfFabI assayed in vitro.<sup>14</sup> More generally, the diaryl urea class of small molecules has been previously reported in the literature to exhibit potent efficacy against cultured parasites<sup>21-25</sup> but without definitive biological target identification. This chemotype was also found amongst hits against P. falciparum dihydroorotate dehydrogenase<sup>26</sup> that lacked whole-cell activity.

In order to more quantitatively compare the diaryl ureas reported herein with those disclosed in the literature as antimalarials, we leveraged a total of thirty-four diaryl ureas generated in our laboratories (Supplementary Data Table 1) to generate a common features pharmacophore (Accelrys Discovery Studio 2.5.5) with five hydrophobic features and two hydrogen bond acceptors. Figure 1 depicts the top 3 active compounds that mapped to it. This pharmacophore is also able to select 23 compounds out of the 451 with the diaryl urea substructure present in the GSK library of antimalarial hits<sup>4</sup> (up to 100 conformers per molecule generated using the FAST algorithm in CAESAR). Slight variants on the diaryl urea scaffold, such as TCMDC-140251 (Supplementary Data Figure S1;  $XC_{50} = 190$  nM against 3D7 strain) containing aryl and 1-indolinyl moieties, mapped well to the model. Interestingly, TCMDC-139010 exhibited a poor fit because it lacked three of the hydrophobic features. The pharmacophore differs from that constructed by Zhang et al. which contained 2-3 hydrophobic features and 2-3 hydrogen bond acceptors.<sup>25</sup> Not surprisingly, the 3 most active diaryl ureas from the paper by Zhang et al. failed to map to our pharmacophore; most likely, the 4-aminoquinaldine-derived diaryl ureas present different features than the triclosan-derived diaryl ureas. Distinctions in the arrangement of hydrophobic and hydrogen bonding features in the diaryl ureas from Zhang et al. and in this study may enable these molecules to target different proteins in *Plasmodium falciparum*. The pharmacophore developed in this study may be leveraged to search other databases (e.g. compound vendor libraries and approved drugs) to identify novel compounds with antimalarial activity.

A novel class of diaryl ureas derived initially from triclosan has been disclosed with regard to their potent *in vitro* efficacy against cultured drug-sensitive and drug-resistant strains of *P. falciparum*. Importantly, family members such as **18** demonstrate promising *in vivo* activity in a *P. berghei* mouse model of infection. Further investigation of the structure-activity relationship of these triclosan derivatives is necessary to further improve their antimalarial activity, in addition to their pharmacokinetic profiles. Biological studies are also important to determine the molecular target(s) of these potent compounds.

## **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### Figure 1.

Common features pharmacophore for triclosan-inspired diaryl ureas showing the top 3 active molecules aligned to the 5 hydrophobic features (cyan) and 2 hydrogen bond acceptors (green).



#### Scheme 1.

Synthesis of various 1-substituted triclosan derivatives. a) KOH, DMSO, 100 °C; b) H<sub>2</sub>, Ra-Ni, EtOH; c) *t*-BuONO, BF<sub>3</sub> Et<sub>2</sub>O, THF, 0 °C ; d) NaI, acetone; e) CuCN, DMF, 150 °C; f) NaOH, H<sub>2</sub>O<sub>2</sub>, EtOH, 50 °C; g) 6 N HCl<sub>(aq)</sub>, 2-methoxyethanol, 80 °C.





#### Scheme 2.

Synthesis of various 1-substituted triclosan amides, sulfonamides, and ureas. Reagents: (a) RSO<sub>2</sub>Cl, NEt<sub>3</sub>, DCM; (b) Ac<sub>2</sub>O or PhC(O)Cl and NEt<sub>3</sub>, DCM; (c) i. (Cl<sub>3</sub>CO)<sub>2</sub>CO, NEt<sub>3</sub>, DCM, -78 °C -> rt, ii. R<sup>2</sup>NH<sub>2</sub>; (d) R<sup>2</sup>NCO, tol.

#### Table 1

In vitro activities of triclosan derivatives with a 1-substituent against cultured P. falciparum strains.



Compound	R	$EC_{50}\ 3D7\ (\mu M)$	$EC_{50}Dd2(\mu M)$
triclosan	ОН	2.9	3.8
1	OCH <sub>3</sub>	100	76
2	CN	20	43
3	СООН	>150	>150
4	CONH <sub>2</sub>	73	60
5a	NH <sub>2</sub>	52	55
6	CH <sub>2</sub> NH <sub>2</sub>	2.3	4.3
7	C(CH <sub>3</sub> ) <sub>2</sub> NH <sub>2</sub>	7.2	1.8
8	NHSO <sub>2</sub> Me	38	60
9	NHSO <sub>2</sub> Ph	4.7	5.1
10	NHAc	33	48
11	NHBz	9.7	9.4
12	NHC(O)NH <sub>2</sub>	18	18
13	NHC(O)NHPh	0.73	1.2



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Compound	Ar	×	EC <sub>50</sub> 3D7 (µM)	ЕС <sub>50</sub> Dd2 (µM)	SC Dose (mg/kg)	F <sub>Survival</sub> @ 31d <sup>a</sup> ,b,c
14	1-naphthyl	$CF_3$	1.2	0.81	256	NES
15	2-naphthyl	$CF_3$	0.61	0.41	256	NES
16	4-CIPh	C	2.0	5.0	256	NES
17	3,4-Cl <sub>2</sub> Ph	C	1.9	2.5	256	NES
18	3-CF <sub>3</sub> -4-CIPh	$CF_3$	0.037	0.055	256	4/7
18					128	EXT
19	3-CF <sub>3</sub> -4-FPh	$CF_3$	0.13	0.15	256	1/7
19					128	2/7
20	3-CF <sub>3</sub> -4-OMePh	$CF_3$	0.18	0.18	$^{\mathrm{pq}}q$	nd
					pu	pu
21	3-CF <sub>3</sub> -4-NO <sub>2</sub> Ph	$\mathrm{CF}_3$	0.072	0.072	hd	nd
22	3-CF <sub>3</sub> -4-NH <sub>2</sub> Ph	$CF_3$	1.5	3.6	nd	nd
23	3-CF <sub>3</sub> -4-CNPh	$CF_3$	0.13	0.075	pu	nd
24	3,4-CN <sub>2</sub> Ph	$CF_3$	0.081	0.14	pu	nd
25	3,4-Cl <sub>2</sub> Ph	$CF_3$	0.20	0.20	hd	nd
<sup>a</sup> FSurvival = p	proportion of animals	; living	at day 31			
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<sup>C</sup>EXT = extension of survival beyond infected control animals (8 days), but no survival at day 31