



Research paper

Evaluation of 7-arylamino-pyrazolo[1,5-*a*]pyrimidines as anti-*Plasmodium falciparum*, antimalarial, and *Pf*-dihydroorotate dehydrogenase inhibitors



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ABSTRACT

Malaria remains one of the most serious global infectious diseases. An important target for antimalarial chemotherapy is the enzyme dihydroorotate dehydrogenase from *Plasmodium falciparum* (*Pf*DHODH), which is responsible for the conversion of dihydroorotate to orotate in the *de novo* pyrimidine biosynthetic pathway. In this study, we have designed and synthesized fifteen 7-arylpyrazolo[1,5-*a*]pyrimidine derivatives using ring bioisosteric replacement and molecular hybridization of functional groups based on the highly active 5-methyl-*N*-(naphthalen-2-yl)-2-(trifluoromethyl)-[1,2,4]triazolo[1,5-*a*]pyrimidin-7-amine. The compounds were tested against *Plasmodium falciparum*, as antimalarials in mice with *P. berghei*, and as inhibitors of *Pf*DHODH. Thirteen compounds were found to be active against *P. falciparum*, with IC₅₀ values ranging from 1.2 ± 0.3 to 92 ± 26 μM in the anti-HRP2 and hypoxanthine assays. Four compounds showed the highest selective index (SI), which is a ratio between cytotoxicity and activity *in vitro*. The inhibition of *Pf*DHODH showed that compound **30** (R₂ = CH₃; R₅ = CF₃; Ar = 7-β-naphthyl) displayed higher and selective inhibitory activity, with IC₅₀ = 0.16 ± 0.01 μM, followed by **25** (R₂ = CH₃; R₅ = CH₃; Ar = 7-β-Naphthyl) and **19** (R₂ = CF₃; R₅ = CF₃; Ar = 7-β-naphthyl), with IC₅₀ = 4 ± 1 μM and 6 ± 1 μM, respectively. The trifluoromethyl group at the 2- or 5-positions of the pyrazolo[1,5-*a*]pyrimidine ring led to increased drug activity. The docking results agreed with the values obtained from enzymatic assays.

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1. Introduction

Among the current devastating infectious diseases, malaria is the third leading cause of death in the world after tuberculosis and

AIDS [1]. According to epidemiological data, it is present in more than 90 countries [2,3]. This means that approximately half of the world's population is exposed to this disease, which has resulted in approximately 198 million cases of malaria in 2013, leading to 584,000 deaths [4].

The emergence of *Plasmodium falciparum* multidrug resistance to chloroquine (CQ) and mefloquine (MQ) [5,6] and the confirmation of cases of artemisinin derivatives resistance [7] justifies the need for research to expand the range of therapeutic targets [8].

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The enzyme dihydroorotate dehydrogenase (DHODH) has been investigated as a pharmacological target for new therapeutic agents to treat cancer, rheumatoid arthritis and parasitic diseases such as malaria, Chagas disease and leishmaniasis [9–12].

The DHODH enzyme participates in the fourth step of the *de novo* pyrimidine biosynthesis pathway. This flavoenzyme catalyses the stereospecific oxidation of (S)-dihydroorotate (DHO) to orotate based on the ping-pong mechanism of catalysis [13]. In the first half-reaction comprising the oxidation of dihydroorotate to orotate, electrons are transferred to the flavin mononucleotide moiety (FMN) that is reduced to dihydroflavin mononucleotide (FMNH₂). After dissociation of orotate from the enzyme, FMNH₂ is regenerated by an appropriate electron receptor [13].

There are two classes of DHODHs that are classified based on the difference in their amino acid sequences and the preference for electron acceptors. Both human (*HsDHODH*) and *P. falciparum* (*PfDHODH*) enzymes belong to class 2, that use ubiquinone as a terminal oxidant agent and are located in the inner mitochondrial membrane [10–12].

HsDHODH inhibitors are not clinically significant and have demonstrated high potency as anti-inflammatory and anti-proliferative activity [14]. Unlike human cells, which can acquire pyrimidine bases through pathway reuse, the *P. falciparum de novo* pyrimidine biosynthetic pathway is essential for the maintenance of life. The inhibition of *PfDHODH* has been proved to be an attractive strategy to search for new substances with antiparasitological activity [9,10,12]. Synthetic compounds have been identified as inhibitors of class 2 DHODH enzymes (Fig. 1) such as brequinar and its analogues **1** [15]. Some triazolopyrimidine compounds such as 5-methyl-*N*-(naphthalen-2-yl)-[1,2,4]triazolo[1,5-*a*]pyrimidin-7-amine (**JZ8**; **2**) and **3**, and *N*-phenylbenzamide **4** were discovered by Phillips and co-workers through enzyme-based high-throughput screening [11,16,17]. Other compounds, such as **5** and **6**, were obtained through pharmacophoric group modification [18,19].

In 2012, we evaluated a series of 2-(trifluoromethyl)-[1,2,4]triazolo[1,5-*a*]pyrimidines against *P. falciparum* that were designed based on bioisosteric replacement of functional groups in the

antimalarial compounds mefloquine and amodiaquine (**11**) [20]. We observed that a trifluoromethyl group at the 2-position of the triazolopyrimidine ring plays an important role in antiparasitological activity (Fig. 2).

Following the good results obtained with 2-(trifluoromethyl)-[1,2,4]triazolo[1,5-*a*]pyrimidines [20], in this study, we planned to use the ring bioisosteric replacement (green) of triazolopyrimidine **10** (Fig. 3) to obtain 7-arylamino-pyrazolo[1,5-*a*]pyrimidines. Molecular hybridization (blue) of 7-β-naphthylamino of **10** was also used. Different arylamines (blue) were incorporated into the structure to investigate the importance of the substituent at the 7-position of the pyrazolopyrimidine nucleus. Investigation of the impact of the substituents CF₃ and CH₃ (red) in the 2- or 5-positions of the pyrazolo[1,5-*a*]pyrimidine scaffold was prioritized. These compounds were evaluated *in vitro* against *P. falciparum*, *in vivo* as antimalarial, and *in vitro* as *PfDHODH* inhibitors.

In addition, a molecular docking study was performed to evaluate the possible binding mode of the 7-arylamino-pyrazolo[1,5-*a*]pyrimidine compounds to *PfDHODH* [21].

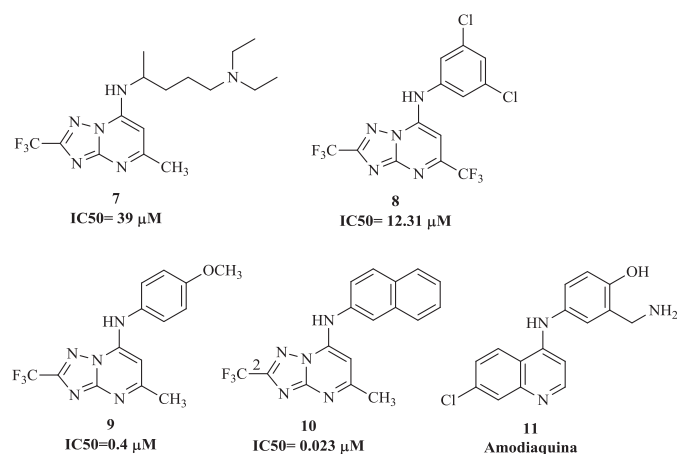


Fig. 2. 2-(trifluoromethyl)-[1,2,4]triazolo[1,5-*a*]pyrimidines **7–10** reported as having antiparasitological activity.

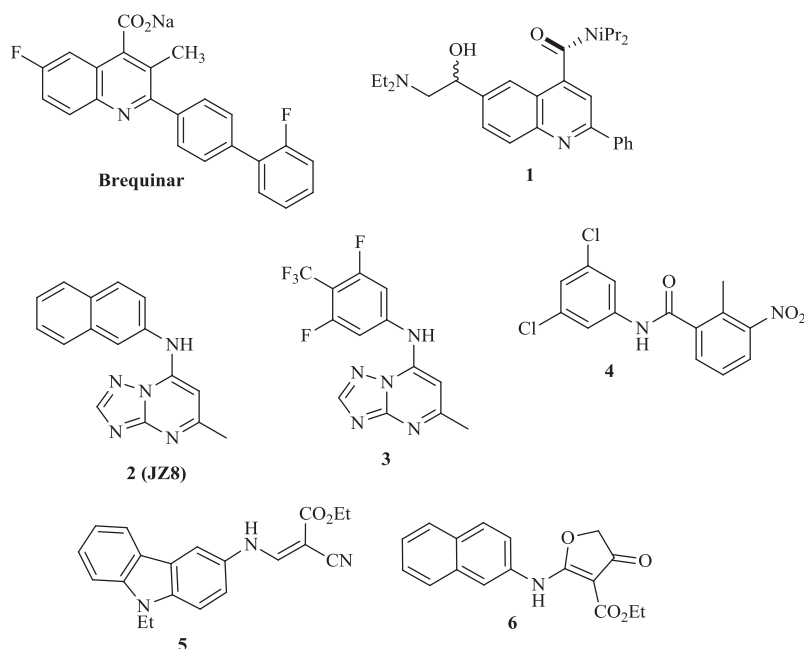


Fig. 1. Brequinar and compounds **1–6** as *PfDHODH* inhibitors.

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