

# Antimalarial activity of novel imidazolidinedione derivatives

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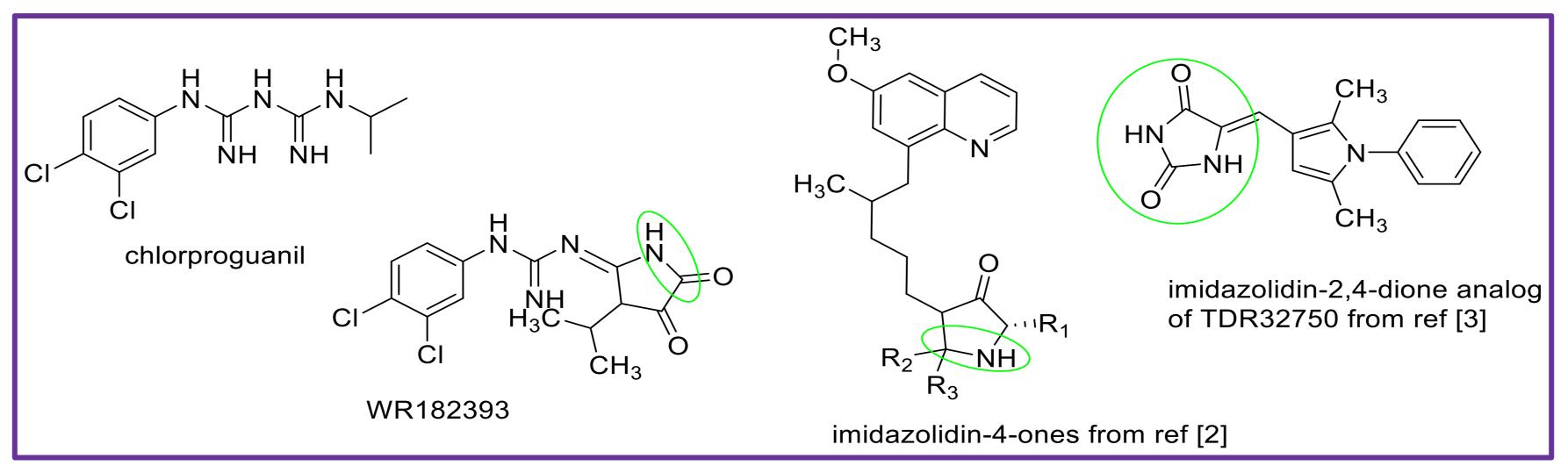
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## Background

Malaria is caused by protozoan parasites of the species *Plasmodium*. Malaria parasites have developed resistance to chemotherapeutic agents such as chloroquine (CQ), mefloquine, and sulfadoxine/pyrimethamine. Therefore, an urgent need exists to develop new classes of antimalarial drugs that operate by novel mechanisms of action. Chlorproguanil is highly active against primary exoerythrocytic forms of *P. falciparum* and *P. vivax*. Moreover, a cyclic dicarboxamide derivative of chlorproguanil, compound WR182393 (2-guanidinoimidazolidinedione derivative) was found to completely eliminate malaria parasites from the body [1]. Several peptide and amino acid derivatives of primaquine and other 8-aminoquinoline antimalarials have been synthesized with the aim of reducing the metabolic oxidative deamination pathway, as well as to reduce toxicity of the parent drug. Moreover, imidazolidin-4-ones prepared from amino acid derivatives of primaquine exhibit potent gametocytocidal activity against *P. berghei* [2]. Systematic structure-activity relationship studies undertaken on a hit compound, TDR32750 (4-oxo-4,5-dihydro-1H-pyrrole-3carboxylate derivative), with the aim of improving antiparasitic activity, revealed that replacement of the 4-oxo-4,5-dihydro-1H-pyrrole core on the imidazolidin-2,4-dione gave a similar level of activity against *P. falciparum* [3].

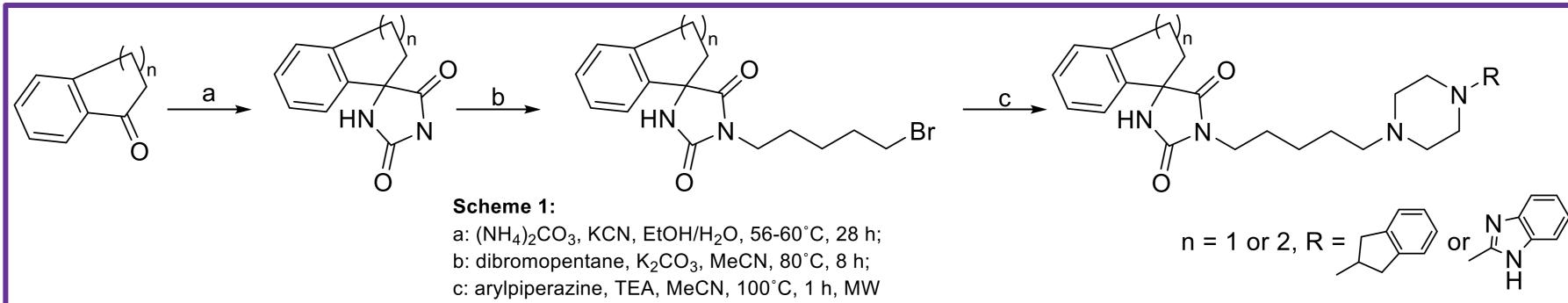


### Objective

The aim of the presented study was the evaluation of antimalarial activity of novel imidazolidinedione derivatives.

## Chemistry

The starting imidazolidinedione rings were prepared from appropriate ketones, 3,4-dihydro-naphthalen-1(2*H*)-one and 2,3-dihydro-1*H*-inden-1-one, using the Bucherer-Bergs reaction, with modifications described by Goodson *et al*. The intermediate products of imidazolidinedione were obtained through an alkylation reaction with 1,5-dibromopentane. In the final step, the intermediate products were coupled with an arylpiperazine moiety to derive the desired compounds (Acz 100-101 and Acz 108-109).



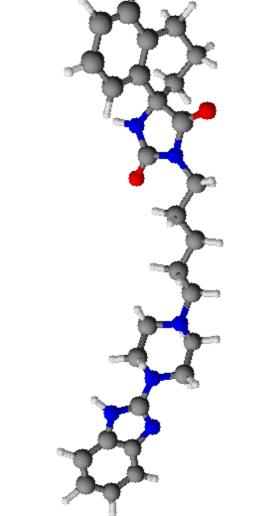
## Methods

*P. falciparum* cultures were established according to Trager and Jensen, with slight modifications [4]. Parasite growth was determined spectrophotometrically (OD<sub>650</sub>) by measuring the activity of parasite lactate dehydrogenase (pLDH), according to a modified version of the method of Makler [5]. Antimalarial activity was expressed as 50% inhibitory concentrations (IC<sub>50</sub>).

### Results

Compound	<i>P. falciparum</i> IC <sub>50</sub> (ng/ml)	
	D10	W2
A <sub>CZ</sub> 100	6202.00 ± 892.09	2424.15 ± 255.34
A <sub>cz</sub> 101	8269.85 ± 736.88	4782.00 ± 97.44
A <sub>CZ</sub> 108	9659.70 ± 140.86	4346.50 ± 659.31
A <sub>cz</sub> 109	> 10000	5648.07 ± 1946.84
CQ	$7.64 \pm 1.70$	102.67 ± 27.49

*In vitro* antimalarial activity of Acz compounds against the D10 (CQ-sensitive) and W2 (CQ-resistant) strains of *P. falciparum*.



3D-structure of the most active compound, Acz 100

### Conclusion

The synthetized derivatives, possessing enhanced antimalarial activity against the CQ-resistant strain of *P. falciparum*, are promising antimalarial drug candidates. The results also indicate the need for development of appropriate lipid delivery systems due to the highly hydrophobic nature of these active compounds.

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