

LSHTM Research Online

O'Neill, PM; Amewu, RK; Charman, SA; Sabbani, S; Gnädig, NF; Straimer, J; Fidock, DA; Shore, ER; Roberts, NL; Wong, MH; +38 more... Hong, WD; Pidathala, C; Riley, C; Murphy, B; Aljayyoussi, G; Gamo, FJ; Sanz, L; Rodrigues, J; Cortes, CG; Herreros, E; Angulo-Barturén, I; Jiménez-Díaz, MB; Bazaga, SF; Martínez-Martínez, MS; Campo, B; Sharma, R; Ryan, E; Shackleford, DM; Campbell, S; Smith, DA; Wirjanata, G; Noviyanti, R; Price, RN; Marfurt, J; Palmer, MJ; Copple, IM; Mercer, AE; Ruecker, A; Delves, MJ; Sinden, RE; Siegl, P; Davies, J; Rochford, R; Kocken, CHM; Zeeman, AM; Nixon, GL; Biagini, GA; Ward, SA; (2017) A tetraoxane-based antimalarial drug candidate that overcomes PfK13-C580Y dependent artemisinin resistance. Nature communications, 8. p. 15159. ISSN 2041-1723 DOI: https://doi.org/10.1038/ncomms15159

Downloaded from: http://researchonline.lshtm.ac.uk/4648924/

DOI: https://doi.org/10.1038/ncomms15159

Usage Guidelines:

Please refer to usage guidelines at https://researchonline.lshtm.ac.uk/policies.html or alternatively contact researchonline@lshtm.ac.uk.

Available under license: http://creativecommons.org/licenses/by/2.5/



ARTICLE

Received 2 Sep 2016 | Accepted 2 Mar 2017 | Published 24 May 2017

DOI: 10.1038/ncomms15159

OPEN

A tetraoxane-based antimalarial drug candidate that overcomes PfK13-C580Y dependent artemisinin resistance

Paul M. O'Neill^{1,2}, Richard K. Amewu^{1,†}, Susan A. Charman³, Sunil Sabbani¹, Nina F. Gnädig⁴, Judith Straimer⁴, David A. Fidock^{4,5}, Emma R. Shore¹, Natalie L. Roberts¹, Michael H.-L. Wong¹, W. David Hong¹, Chandrakala Pidathala¹, Chris Riley¹, Ben Murphy¹, Ghaith Aljayyoussi⁶, Francisco Javier Gamo⁷, Laura Sanz⁷, Janneth Rodrigues⁷, Carolina Gonzalez Cortes⁷, Esperanza Herreros⁷, Iñigo Angulo-Barturén⁷, María Belén Jiménez-Díaz⁷, Santiago Ferrer Bazaga⁷, María Santos Martínez-Martínez⁷, Brice Campo⁸, Raman Sharma⁶, Eileen Ryan³, David M. Shackleford³, Simon Campbell⁸, Dennis A. Smith⁸, Grennady Wirjanata⁹, Rintis Noviyanti¹⁰, Ric N. Price^{9,11}, Jutta Marfurt⁹, Michael J. Palmer⁸, Ian M. Copple², Amy E. Mercer², Andrea Ruecker¹², Michael J. Delves¹², Robert E. Sinden^{12,13}, Peter Siegl⁸, Jill Davies^{6,†}, Rosemary Rochford¹⁴, Clemens H.M. Kocken¹⁵, Anne-Marie Zeeman¹⁵, Gemma L. Nixon¹, Giancarlo A. Biagini⁶ & Stephen A. Ward⁶

K13 gene mutations are a primary marker of artemisinin resistance in *Plasmodium falciparum* malaria that threatens the long-term clinical utility of artemisinin-based combination therapies, the cornerstone of modern day malaria treatment. Here we describe a multinational drug discovery programme that has delivered a synthetic tetraoxane-based molecule, E209, which meets key requirements of the Medicines for Malaria Venture drug candidate profiles. E209 has potent nanomolar inhibitory activity against multiple strains of *P. falciparum* and *P. vivax in vitro*, is efficacious against *P. falciparum* in *in vivo* rodent models, produces parasite reduction ratios equivalent to dihydroartemisinin and has pharmacokinetic and pharmacodynamic characteristics compatible with a single-dose cure. *In vitro* studies with transgenic parasites expressing variant forms of K13 show no cross-resistance with the C580Y mutation, the primary variant observed in Southeast Asia. E209 is a superior next generation endoperoxide with combined pharmacokinetic and pharmacodynamic features that overcome the liabilities of artemisinin derivatives.

¹Department of Chemistry, University of Liverpool, Liverpool L69 7ZD, UK. ²Department of Pharmacology, School of Biomedical Sciences, MRC Centre for Drug Safety Science, University of Liverpool, Liverpool L69 3GE UK. ³ Centre for Drug Candidate Optimisation, Monash Institute of Pharmaceutical Sciences, Monash University, 381 Royal Parade, Parkville, Victoria 3052, Australia. ⁴ Department of Microbiology and Immunology, Columbia University College of Physicians and Surgeons, HHSC 1502, 701 W. 169th Street, New York, New York 10032, USA. ⁵ Division of Infectious Diseases, Department of Medicine, Columbia University Medical Center, HHSC 1502, 701 W. 168th Street, New York, New York 10032, USA. ⁶ Research Centre for Drugs and Diagnostics, Liverpool School of Tropical Medicine, Pembroke Place, Liverpool L3 SQA, UK. ⁷ Tres Cantos Medicines Development Campus, DDW, GlaxoSmithKline, Severo Ochoa 2, 28760 Tres Cantos, Spain. ⁸ Medicines for Malaria Venture, ICC, Route de Pré-Bois 20, P.O. Box 1826, 1215 Geneva, Switzerland. ⁹ Global and Tropical Health Division, Menzies School of Health Research, Charles Darwin University, P.O. Box 1826, 1215 Geneva, Switzerland. ⁹ Global and Tropical Health Division, Menzies School of Health Research, Charles Darwin University, P.O. Box 1826, 1215 Geneva, Switzerland. ⁹ Global and Tropical Health Division, Menzies School of Health Research, Charles Darwin University, P.O. Box 14096, Casuarina, Darwin, Northern Territory 0811, Australia. ¹⁰ Eijkman Institute for Molecular Biology, JI. Diponegoro 69, 10430 Jakarta, Indonesia. ¹¹ Nuffield Department of Clinical Medicine, Centre for Tropical Medicine and Global Health, University of Oxford, Old Road Campus, Roosevelt Drive, Oxford OX3 7FZ, UK. ¹² Department of Life Sciences, Imperial College London, South Kensington, London SW7 2AZ, UK. ¹⁴ Department of Immunology and Microbiology, University of Colorado, Aurora Colorado, CO 80045, USA. ¹⁵ Department of Parasitology, Biomedical Primate Research Centre, P. O.

he control and elimination of malaria requires effective treatment strategies and the universal availability of artemisinin-based combination therapies $(ACTs)^{1,2}$. For over a decade, research groups have sought to replace the semisynthetic artemisinin components (Fig. 1a, **1a-1c**) of ACTs with a fully synthetic alternative to reduce uncertainties regarding cost and supply and to improve overall pharmacological qualities, most notably short clinical half-lives. Several pharmacophores have been explored³ including the tetraoxanes (Fig. 1b, **3a** and **3b**).

Recent years, however, have witnessed the emergence of parasite 'resistance' to artemisinin, first documented in western Cambodia^{4,5}. Malaria patients from this region were reported to have a delayed parasite clearance following either artesunate monotherapy or an ACT. The delayed parasite clearance phenotype is not resistance as defined by the WHO^{6,7} and does not necessarily lead to treatment failure⁸. However, it gives rise to concerns of a reduction in the therapeutic lifespan of the most effective, currently registered anti-malarials and the only class that offers rapid parasite biomass reduction. Furthermore, reduced artemisinin efficacy places greater selective pressure on the ACT partner drugs, increasing the risk of multidrug resistance to emerge. Indeed, recent results in Cambodia now document increasing rates of clinical treatment failure in patients treated with the first-line combination dihydroartemisinin plus piperaquine, because of the emergence of resistance to both agents⁹

Here we describe the development of a fully synthetic tetraoxane analogue, E209, which displays nanomolar efficacy against multiple strains of *P. falciparum* and *P. vivax* with a tailored combination of pharmacodynamic (PD) and pharmacokinetic (PK) properties compatible with a single-dose cure even against a backdrop of the K13-dependent resistance mechanism in operation in S.E. Asia⁸.

Results

Optimization of the tetraoxane series. The tetraoxane core **3a** is differentiated from 1,2,4-trioxolanes **2a** and 1,2,4-trioxanes as it is achiral, has greater inherent thermodynamic stability and potent *in vitro* antimalarial activity, either as a simple unsubstituted analogue, for example, **3a** (Fig. 1) or related derivatives. Recent theoretical calculations based on stereoelectronic analysis suggest that the remarkable thermodynamic stability of tetraoxanes can be attributed to a stereoelectronic 'double anomeric effect' which stabilizes the six-membered ring system¹⁰.

Our initial medicinal chemistry focus was the preparation of variants of a simplified analogue of our previous candidate RKA 182 with more balanced ADME properties $(3b)^{11}$ (Supplementary Figs 1, 2, 3 and 4). Multiple series were prepared to modulate solubility, metabolic stability, pKa and log D (Supplementary Fig. 4) but in all cases, the amide linked analogues, although superior to the artemisinins, had relatively short half-lives and poor antimalarial activity in the Plasmodium berghei mouse model. We next prepared tetraoxane 4 as a direct comparator (Supplementary Fig. 4; Supplementary Table 1) to OZ439 (ref. 12), which is currently in phase 2 clinical trials. Although this analogue had equivalent in vitro potency to OZ439, it only achieved a 16.3 day mean survival time (MSD) at a single $1 \times 30 \text{ mg kg}^{-1}$ oral dose in *P. berghei* infected mice (Table 1). This molecule was also slightly less stable in microsomal studies in vitro with high clearance rates and a short half-life in rat PK studies.

Cognisant of the liabilities of the artemisinin class, single-dose survival time, intrinsic clearance and blood stability were used as the key parameters to drive SAR studies towards lead series progression (Supplementary Fig. 4; Supplementary Information). Representative lead compounds are shown in Supplementary Fig. 4 from which E209 and N205 were the clear front-runner molecules with respect to cures in the *P. berghei* model and initial DMPK assessments (Supplementary Table 1; Supplementary Fig. 5) and were selected for further profiling. *In vitro* metabolic stability assessments (Supplementary Table 2) and solubility assessments (Supplementary Table 3) demonstrated the superior properties of E209 and this compound was taken forward.

Following scale up synthesis (Supplementary Note 2; Supplementary Fig. 1), E209 was progressed to candidate selection through a series of chemical, parasitological, pharmacological and toxicological checkpoints as outlined below.

PD properties of E209. E209 retained potent in vitro efficacy (mean IC₅₀ range 2.9-14.0 nM) against a panel of 10 sensitive and multidrug-resistant P. falciparum parasite isolates from distinct geographical origins, with no observable cross-resistance in growth inhibition studies (Supplementary Table 4). Supplementary Table 4 represents a panel of parasite isolates used as part of a standard Medicines for Malaria Venture (MMV) in vitro evaluation and includes parasites that are universally sensitive to current drugs (HB3) and parasites resistant to chloroquine (as indicated), mefloquine (FCB, NF54), pyrimethamine and related DHFR inhibitors (K1, TMC90, V1/S and so on) and atovaquone (TMC90). In the case of E209, activity against all parasites is below the accepted cut off for resistance and it is clear from visual inspection that there is no relationship at all with chloroquine sensitivity. Cytotoxicity studies using HeLa cells demonstrated that E209 had a high in vitro therapeutic index of 200-1,000, with a median IC_{50} 2.85 (s.d. ± 0.99 μ M, n=3) (c.f. dihydroartemisinin cytotoxicity (median IC₅₀ 0.4 $(s.d. \pm 0.1 \,\mu M, n = 3)).$

Ex vivo P. falciparum and P. vivax drug sensitivity experiments were carried out in Papua Province in eastern Indonesia, an area with documented multidrug-resistant P. falciparum and CQresistant *P. vivax*, using a modified schizont maturation assay as described previously¹³⁻¹⁷. Ethical approval for the *ex vivo P*. falciparum and P. vivax efficacy data was obtained from the Human Research Ethics Committee of the NT Department of Health & Families and Menzies School of Health Research, Darwin, Australia (HREC 2010-1396) and the Eijkman Institute Research Ethics Commission, Jakarta, Indonesia (EIREC 47 and EIREC 67). E209 exhibited equipotent ex vivo activity against *P. vivax* (n = 13; median IC₅₀: 10.5 nM) and *P. falciparum* (n = 3; median IC₅₀: 15.7 nM) clinical field isolates from Papua, Indonesia (Mann–Whitney U test, P = 0.570); Supplementary Table 5. With the exception of chloroquine in P. falciparum (Spearman rank correlation, P = 0.042), there was no correlation between the activity of E209 and any of the other anti-malarials assessed in either species ((amodiaquine (AQ), piperaquine (PIP), mefloquine (MFQ) and artesunate (AS) (Supplementary Table 6)).

E209 was screened for late stage gametocytocidal activity in a standard membrane feeding assay (SMFA) and transmissionblocking activity was determined in *Anopheles stephensi* mosquitoes. Late stage gametocytes were incubated with E209 for 24 h prior to mosquito feeding and the IC₅₀, (determined by the reduction in oocyst mean intensity) was 14.5 nM with a corresponding ~60 and 90% transmission-blocking activity (defined as the percentage of mosquitoes with no observed oocysts) recorded at 100 nM and 1 μ M, respectively (Supplementary Fig. 6). The transmission reducing activity (TRA) shown by E209 is consistent with that shown for the endoperoxide class as reported in recent SMFA-based studies^{18,19}.



Figure 1 | Artemisinins and synthetic peroxides. (a) Artemisinin and semi-synthetic analogues. (b) Comparison of tetraoxanes with trioxolane-based anti-malarials. MSD in days following a single oral dose of 30 mg kg⁻¹, Pf SCID (NOD-SCID IL- $2R\gamma c^{null}$ (NSG) mice engrafted with human erythrocytes and infected with *P. falciparum* strain 3D7^{0087/N9}).

E209 ($10 \times IC_{50}$) displayed fast killing kinetics (Supplementary Methods), with an *in vitro* parasite reduction ratio ($log_{10}PRR$) > 4.8 similar to artemisinin ($log_{10}PRR > 4.8$), and faster than CQ ($log_{10}PRR 4.5$), pyrimethamine ($log_{10}PRR 3.7$) and atovaquone ($log_{10}PRR 2.9$) (Fig. 2)²⁰.

Critical for the development of E209 and related structures is the evaluation of their activity against ART-resistant parasites, which has been defined as delayed parasite clearance in patients⁵. This phenotype has been associated clinically with a number of nonsynonymous SNPs in the propeller domain of the K13 protein, located on chromosome 13 (ref. 23). Notably, these mutations could not be associated with ART resistance using the standard in vitro 72h inhibitory assay, therefore a novel phenotypic assay was developed to assess ART resistance in vitro. The ring-stage survival assay (RSA_{0-3 h}) is based on the increased resistance of young ringstage parasites to a 6 h pulse of 700 nM dihydroartemisinin (DHA)^{21,22}. To investigate the impact of K13 mutations on E209 susceptibility, a series of gene-edited, otherwise isogenic parasite lines with either K13 mutant or wild-type alleles have been subjected to RSA_{0-3 h} assays using DHA as comparison²³. This allowed us to assess the potency of a pharmacological relevant dose of E209 against K13 mutant parasites and to identify potential cross-resistance phenotypes. In this study we focused on the most prevalent mutation C580Y and R539T, which was demonstrated to confer high levels of DHA resistance in vitro²³(Fig. 3).Generally, we observe higher

survival rates after exposure of young ring stages to E209 in both parasite backgrounds when compared to DHA. When subjected to a 700 nM dose for 6 h, we observe 4% survival of V1/S and 16% of Cam3.II parasites for E209 compared to <1% survival when pulsed with DHA. The highly prevalent *K13* mutation C580Y does not confer cross-resistance to E209. However, R539T shows a moderate but significant increase of parasite survival in both parasites lines tested. (Ring-stage survival assays, fold changes between K13 wild-type and K13-mutant lines and IC₅₀ data for DHA and E209 are recorded in Supplementary Tables 15 and 16; Supplementary Fig. 7).

In vivo experiments using *P. berghei* infected mice demonstrated that E209 when given orally resulted in complete parasite clearance in the Peters' standard 4-day suppressive test²⁴, with an estimated ED₅₀ of 4 mg kg⁻¹ after three doses. Importantly, E209 also achieves 66% cure rate after a single oral dose of 30 mg kg⁻¹ (Table 1). Efficacy of E209 against the human malaria parasite was tested in the immune-deficient NOD-SCID IL-2R_null mouse engrafted with human erythrocytes and infected with the *P. falciparum* strain 3D7^{0087/N9} (ref. 25). Complete parasite clearance was achievable with a single oral dose of E209 (30 mg kg⁻¹) with a calculated ED₉₀ of 11.6 mg kg⁻¹ and an AUC ED₉₀ of 1.2 µg h ml⁻¹ (Fig. 4). This compares with an ED₉₀ of 10 mg kg⁻¹ for artesunate following four consecutive daily doses²⁶. The maximum rate of parasite killing by E209 was achieved with blood exposure between 5 and 10 µg h ml⁻¹

Compound Number	Side Chain (R)	ClogP	PSA/Ų	IC ₅₀ (3D7) nM	<i>In vivo</i> (Pb) MSD (days) at 1 x 30 mg/kg
4 [†]		5.6	58.6	10 ± 1	16.3 (14, 21, 21)
5^{\dagger}	₹~~~N_O	5.9	58.6	7.5 ± 0.2	13 (13, 13, 13)
6†	NSO ₂ Me	5.6	86.7	3.5 ± 0.2	19.3 (14, 30,14)
7 [†]		5.9	86.7	3.7 ± 0.1	14 (14, 14, 14)
8†	N F F	6.1	49.3	8.5 ± 0.6	(8, 8, 8 (8)
E209 [†]	₹ N F	6.4	49.3	5.1 ± 0.81	25.0 (15, 30, 30) (2 cures)
OZ439 ¹²	See Figure 1b	5.5	49.3	5.1 ± 0.81	30 (30, 30, 30) (3 cures)

Table 1 | Calculated physicochemical* properties and in vitro and in vivo antimalarial activity profiles of selected tetraoxane derivatives.



(Fig. 4). Therefore, this range of exposure in human blood is predicted to be the lowest necessary to achieve the maximum parasite killing and can therefore be defined as the minimum parasiticidal concentration (MPC).

In vitro and in vivo PKs of E209. E209 exhibited metabolic degradation in human, rat and mouse liver microsomes with rates generally being fastest in rat and slowest in the mouse (Supplementary Table 17; Table 2). Comparing metabolite profiles across all three species, hydroxylation(s) on the adamantane ring to form multiple M+16 metabolites represent important pathways in all three species. In addition, N-oxidation to M+16 appears to be an important metabolic pathway in rat whilst tetraoxane cleavage products were observed in all three species (Supplementary Tables 17, 18, 19; Supplementary Fig. 8).

Compartmental PK modelling of preclinical in vivo data showed a clear two compartmental trend for all three species. Simulated PK profiles based on these compartmental model predictions (Table 2) showed good agreement with observed data as shown in Fig. 5a-c). Allometric scaling (see Methods) rat PK data predicted a terminal half-life in humans of 24-30 h and drug levels that exceed E209 IC₅₀ levels for up to 6 days following a single oral dose of 15 mg kg^{-1} (Fig. 5d).

Preclinical safety studies. To assess potential for adverse activities, we profiled E209 using a diverse panel of radioligand binding assays (CEREP screen), ion channel voltage clamp assays and genotoxicity assays. In the CEREP screen, E209 at 10 µM demonstrated a favourable profile apart from moderate potency at Sigma 1 and Sigma 2 receptors (IC50s ca. 200 nM). Risk of adverse cardiac activity including QT interval prolongation is predicted to be low based on the measurement of a high IC₅₀ $(>3.5\,\mu\text{M})$ in the hERG assay and no inhibitory activity at > 100 μ M in Cav1.2 and Nav1.5 assays. E209 was negative in the Ames test and mouse micronucleus test confirming the absence of mutagenic potential.

We also examined the potential haemolytic effect of E209 in NOD-SCID mice engrafted with the African variant G6PD deficient human red blood cells (huRBC) as previously reported for primaquine²⁷. Mice treated with E209 at doses up to $50 \text{ mg kg}^{-1} \text{ day}^{-1}$ for 3 days did not show a significant drugdependent loss of huRBC at 7 days post-treatment with all test groups demonstrating a response similar to the vehicle control (Supplementary Fig. 9A-C). In contrast primaquine, a drug that causes haemolytic toxicity in humans with G6PD deficiency, induced significant loss of huRBC (Supplementary Fig. 9A-C).

In addition, In vitro studies with cytochrome P450 (CYP) isoforms demonstrated that E209 did not inhibit any of the major human CYP450s with $IC_{50}s > 20 \,\mu\text{M}$ (Supplementary Table 20).

The maximum-tolerated dose (MTD) following a single oral administration of E209 in Sprague-Dawley rats was 300 mg kg^{-1} . A 7-day repeat dose, exploratory toxicity study in rats was performed using dose levels of 50,100 and 100 mg kg⁻¹ and parallel groups for toxicokinetic analysis. At 200 mg kg⁻¹ day⁻¹ E209 was not tolerated in either sex. At 100 mg kg⁻¹ there was body weight loss in females. On the basis of weight loss and histopathology findings, the no observed adverse effect level (NOAEL) for this study was 100 mg kg^{-1} day^{-1} for males and 50 mg kg⁻¹ day⁻¹ for females. Systemic



Figure 2 | Parasite reduction ratios of clinically used anti-malarials and E209. (a) *In vitro* PRR—number of viable parasites after E209 treatment $(10 \times IC_{50})$ is compared with the profile shown by standard anti-malarials (b) Lag phase, log_{10} PRR and parasite clearance time (PCT) values for E209 and a selection of standard anti-malarials. The *in vitro* parasite reduction rate assay was used to determine onset of action and rate of killing as previously described²⁰. *P. falciparum* was exposed to E209 at a concentration corresponding to $10 \times EC_{50}$. The number of viable parasites at each time point was determined as described by Sanz *et al.*²⁰ Four independent serial dilutions were done with each sample to correct for experimental variation; error bars, s.d. Previous results reported on standard anti-malarials tested at $10 \times EC_{50}$ using the same conditions are shown for comparison.



Figure 3 | Ring-stage survival assays show equal potency of E209 against K13 wild-type and C580Y mutant parasites whereas R539T confers a marginal increase of survival. Results show the percentage of early ring-stage parasites (0-3 h post-invasion of human erythrocytes) that survived a 6-h pulse of 700 nM DHA or E209, as measured by flow cytometry 90 hours later. Data show mean ± s.e.m. percentage survival compared to DMSO-treated parasites processed in parallel. Assays were performed on at least 2 separate occasions in duplicate. K13-propeller mutations R539T and C580Y confer resistance to DHA in the clinical isolate Cam3.II and the reference line V1/S (**a**), (**b**). Exposure to E209 reveals generally higher survival with slight crossresistance observed in parasites carrying the R539T mutation (**c**), (**d**). A two-sample t-test with unequal variances was used to calculate *P* values. These can be found in Supplementary Tables 7-14. **P* value <0.05, ***P* value <0.001.

exposure values (C_{max} and AUC_{0-24 h}) at the NOAEL on day 7 were 1.2 µg ml⁻¹ and 19.6 µg h ml⁻¹, respectively, in males and 1.07 µg ml⁻¹ and 14.8 µg h ml⁻¹, respectively, in females. Additional single and multiple dose safety pharmacology studies are in progress in rodent and non-rodent models to establish toxicity profile and predicted exposure multiple at NOAEL over target therapeutic exposure (therapeutic index).

Discussion

A systematic evaluation of the antimalarial tetraoxane scaffold (Supplementary Note 1), revealed the aryloxy template (Fig. 1) to be the most promising in terms of multi-parameter lead-optimization. Through a series of rational chemical modifications we have been able to improve both the inherent pharmacody-mamics (*in vitro* and *in vivo* potency) and PKs (metabolic



Figure 4 | Efficacy of E209 in the *in vivo*-humanized SCID mouse model of *P. falciparum*. (a) Parasitemia in peripheral blood of NSG mice infected with the *P. falciparum* strain $3D7^{0087/N9}$ (n = 2 mice treated with vehicle and n = 6 mice treated with E209). (b) Blood levels of E209 in the efficacy experiment shown in **a** over the first 24 h post-dosing. (c) dose/exposure-response relationship.

Table 2 | *In vivo* and *in vitro* pharmacokinetic (PK) parameters for E2O9 as predicted using compartmental PK analysis in male Beagle dogs, female Swiss outbred mice and the male Sprague-Dawley rat following IV and oral administration.

	Rat (mean)	Mouse (mean)	Dog* (mean)	Human (mean)
In vivo PK parameters				
IV dose $(mg kg^{-1})$	2	2	1	N/A
PO dose $(mg kg^{-1})$	9	10	5	15mg [†]
Central clearance, CL ($1h^{-1}kg^{-1}$)	1.6	4.3	1.1	0.41
Central volume of distribution $V_{\rm c}$ (l kg ⁻¹)	0.54	0.99	0.76	0.54†
Inter-compartmental clearance 1, Q_1 (I $^{-1}h^{-1}kg^{-1}$)	1.76	10.3	2.8	0.45†
Inter-compartmental clearance 2, Q_2 (I h $^{-1}$ kg $^{-1}$)	0.08	0.97	0.13	0.020
Peripheral Volume of distribution 1, V_{p1} (1 kg ⁻¹)	2.1	1.49	0.37	2.1 [†]
Peripheral volume of distribution 2, V_{p2} (1 kg^{-1})	2.0	10.8	2.95	2.0†
Absorption rate constant K_a (h ⁻¹)	0.31	0.79	0.33	0.31†
F (%)	62	82	40	62 [†]
In vitro microsomal data				
Degradation half-life (min)	48	132	173	68
In vitro CL _{int} (μ l min ⁻¹ mg ⁻¹ protein)	36	13	10	25
Microsome-predicted E _H	0.48	0.22	0.38	0.50
*Supplementary Methods and Supplementary Table 9 in Supporting Informatio †On the basis of allometric scaling from rat parameters.	n.			

stability profiles in a range of species and rodent and human blood stability) of the class.

In addition to overall high potency, our studies also demonstrate that the fast PRR for E209 observed in *in vitro* studies translates to a fast action *in vivo* in the humanized SCID mouse model where a maximum parasite clearance was achieved after a single dose of 30 mg kg^{-1} . To achieve this level of parasite clearance with the clinical gold standard endoperoxide artesunate requires four doses of artesunate ($50 \text{ mg kg}^{-1} \text{ day}^{-1}$) in the same model²⁵. Initial safety and pharmacology profiles for E209 meet the requirements of MMV's target candidate profile 1 (TCP1), a drug with a fast killing profile (PRR equivalent to or better than dihydroartemisinin) and PK qualities that would support a single-dose cure either singly or in combination.

It is generally accepted that the mechanism of action of endoperoxide anti-malarials involves intra-parasitic ferrous mediated reductive scission of the endoperoxide bond. This process leads to the generation of reactive oxygen species such as carbon-centered radicals, which react with multiple parasite proteins and result in parasite killing^{28–30}. Because of this

proposed mechanism of action it was originally argued that malaria parasites would find it difficult to acquire resistance to this drug class. However, compelling clinical data from Southeast Asia has confirmed that resistance has emerged and established itself in the region. The resistance phenotype has been difficult to study, as it does not show up as a shift in potency (IC50 value) in traditional drug sensitivity assays²⁰. Extensive molecular investigations have implicated mutations in the K13 gene as key to this unusual resistance phenotype. These mutations allow the parasite to survive exposure to drug at the early ring stage of red blood cell infection even at supra-pharmacological drug concentrations^{20,21}. A concern is that this type of resistance mechanism would blight the long-term clinical utility of the entire endoperoxide class of antimalarial drugs. Our interpretation of this resistance mechanism, which is supported by recent modelling studies²⁸, is that the slow parasite clearance phenotype seen clinically is a result of the loss of drug susceptibility in ring-stage parasites coupled with the extremely short elimination half-life of the endoperoxide. A potential solution to this problem is to extend the current 3-day artemisinin treatment course to four days as suggested by



Figure 5 | Measured rodent and predicted human exposure profiles. PK profiles of E209 in male Sprague–Dawley rats (**a** oral dose = 9 mg kg^{-1} , IV dose = 2 mg kg^{-1}), outbred female Swiss mice (**b** oral dose = 10 mg kg^{-1} , IV dose = 2 mg kg^{-1}) and the male beagle dog (**c** oral dose = 5 mg kg^{-1} , IV dose = 1 mg kg^{-1}) and the male beagle dog (**c** oral dose = 5 mg kg^{-1} , IV dose = 1 mg kg^{-1}) and allometrically scaled predictions for human PK (**d**) based on all of the above assuming an oral dose of 15 mg kg^{-1} . Fits constructed are based on predicted median PK parameter values.

Dogovski et al.³¹ We believe that E209 provides a more elegant solution since it circumvents or minimizes the ring-stage resistance seen with currently deployed endoperoxides with a predicted elimination half-life of 24-30 h compared to current artemisining that have half-lives of circa 1-2h (ref. 32). Hence, not only will E209 plasma levels remain above the IC₅₀ level for 4 days or more in malaria patients as required by MMV's TCP1 criteria (based on PK predictions, Supplementary Methods), but it will also retain killing potential throughout each individual stage of parasite's 48 h intra-erythrocytic cycle. Thus, E209 has the potential for deployment in a superior combination treatment with a partner drug devoid of existing in vivo resistance liabilities (such as the ATP4 inhibitors^{33,34} or DHODH inhibitors³⁵). E209 has the potential to offer a substantial improvement on currently deployed artemisininbased drug combinations and provide an urgently required alternative TCP1 drug for malaria treatment and elimination programmes.

Methods

Synthesis of E209. Details of the synthesis and analytical data on E209 are included in Supplementary Note 2.

Parasite culture and drug sensitivity testing. *P. falciparum* blood stage cultures were maintained by the method of Trager and Jensen³⁶. Drug sensitivity during E209 development and QSAR was determined by the method of Winter *et al.*³⁷ Drug sensitivity of E209 against panel of *P. falciparum* strains was determined based on hypoxanthine incorporation^{38,39}. Mammalian cell cytotoxicity assays were carried out as previously described⁴⁰.

A zinc-finger nuclease (ZFN)-based gene editing approach was employed to specifically introduce or remove K13 mutations from *P. falciparum* lines. Parasites were electroporated with a plasmid containing ZFNs specifically engineered against *K13*. The nucleases were expressed from an episomally maintained plasmid by selecting with 2.5 nM (Cam3.II) or 10 nM (V1/S) WR99210 for 6 days. A 1.5 kb K13 donor fragment harbouring the desired mutation was placed on the same

plasmid and served as template for the double strand break repair. Parasite clones were generated by limiting dilution. A detailed protocol can be found in Straimer $et \ al.^{21}$

Ring-stage survival assays ($RSA_{0-3 h}$) were carried out as previously described with minor modifications^{21,23}. In summary, 10–15 ml parasite cultures were synchronized 1-2 times using 5% sorbitol (Sigma-Aldrich). Synchronous multinucleated schizonts were incubated in RPMI-1640 containing 15 units/ml sodium heparin for 15 min at 37 °C to disrupt agglutinated erythrocytes (purchased from Interstate Blood Bank located in Memphis, TN, USA), concentrated over a gradient of 75% Percoll (Sigma-Aldrich), washed once in RPMI-1640 and incubated for 3 h with fresh erythrocytes to allow time for merozoite invasion. Cultures were then subjected again to sorbitol treatment to eliminate remaining schizonts. The 0-3 h post-invasion rings were adjusted to 1% parasitemia and 2% haematocrit in 1 ml volumes (in 48-well plates), and exposed to 700 nM DHA, TDD-E209, JC3-39 or 0.1% DMSO (solvent control) for 6 h. Duplicate wells were established for each parasite line ± drug. The supernatant was then removed, the culture resuspended in 1 ml culture media and transferred to 15 ml conical tubes, centrifuged at 800g for 3 min to pellet the cells and the supernatants carefully removed. As a washing step to remove drug, 10 ml culture medium were added to each tube and the cells resuspended, centrifuged and the medium aspirated. This washing procedure was repeated three times and included a transfer into a new 15 ml conical tube after the second washing step. Fresh medium lacking drug was then added to cultures, which were returned to standard culture conditions for 90 h. Parasite viability was assessed by microscopic examination of Giemsa-stained thin blood smears. Second-generation ring stages are clearly visible under the microscope but only give a faint signal when measured by flow cytometry. Therefore, we performed a media change 66 h after the drug incubation and allowed the culture to develop for another 24 h into second-generation trophozoite stages. Parasite survival was determined by staining parasites with $2 \times$ SYBR Green I and 165 nM MitoTracker Deep Red (Invitrogen) and a minimum of 8,000 cells were counted by flow cytometry on an Accuri C6 cytometer. Percentage survival was calculated as the parasitemia in the drug-treated sample divided by the parasitemia in the untreated sample \times 100.

P. cynomolgi M strain (a non-human primate malaria closely related to the human malaria *P. vivax* and forming dormant liver stages) was used for *in vitro* liver stage drug assays (The M strain BPRC was obtained from Centers for Disease Control and Prevention (CDC, USA). Liver stage drug assays were performed in primary rhesus monkey hepatocytes as described⁴¹. Read out of the assay was performed using a high-content imaging system (Operetta) and analysed with Harmony software. As described previously⁴² two populations of liver stage parasite can be discriminated in 6-day-old cultures: large forms define liver

schizonts and persistent small forms are dormant liver stages (hypnozoites). Small forms are defined as having a maximum parasite area of $30 \,\mu\text{m}^2$.

Ex vivo P. falciparum and P. vivax drug sensitivity experiments were carried out in Papua Indonesia, an area with documented multidrug-resistant P. falciparum and CQ-resistant P. vivax., using a modified schizont maturation assay as described previously $^{13-17}.$ In brief, 200 μl of a 2% haematocrit blood media mixture (BMM), consisting of RPMI-1640 medium plus 10% AB⁺ human serum (P. falciparum) or McCoy's 5A medium plus 20% AB⁺ human serum (P. vivax) was added to each well of pre-dosed drug plates containing 11 serial concentrations (2-fold dilutions) of the anti-malarials being tested. A candle jar was used to mature the parasites at 37°C for 35-56 h. Incubation was stopped when >40% of ring-stage parasites had reached mature schizont stage in the drug-free control wells. Thick blood films made from each well were stained with 5% Giemsa solution for 30 min and examined microscopically. The number of schizonts per 200 asexual stage parasites was determined for each drug concentration and normalized to the control well. The dose-response data were analysed using nonlinear regression analysis (WinNonLn 4.1, Pharsight) and the IC50 value derived using an inhibitory sigmoid Emax model.

Ethical approval for the *ex vivo P. falciparum* and *P. vivax* efficacy data was obtained from the Human Research Ethics Committee of the NT Department of Health & Families and Menzies School of Health Research, Darwin, Australia (HREC 2010-1396) and the Eijkman Institute Research Ethics Commission, Jakarta, Indonesia (EIREC 47 and EIREC 67). Written informed consent was obtained from all patients participating in the study.

The SMFA study was conducted as previously described where mature gametocytes were incubated for 24 h with different concentrations of E209 prior to performing mosquito feeds^{43–45}. The RSA_{0-3h} was performed as described in Witkowski *et al.*,²¹ whereby tightly synchronized early ring-stage parasites were exposed to a drug pulse for 6 h at 700 nM. After drug removal (by washing) the parasites were returned to culture. Parasitemias were then assessed 66 h later on thin blood smears by two independent microscopists and the percentage survival was assessed in comparison to parasites not exposed to drug, as described by Straimer *et al.*²³ Assays were performed in duplicate on three separate occasions. Parasite reduction ratios were conducted as described in Supplementary Methods²⁰.

In vivo efficacy. Studies of murine P. falciparum infection were ethically reviewed and carried out in accordance with European Directive 2010/63/EU and the GSK Policy on the Care, Welfare and Treatment of Laboratory Animals. In vivo efficacy against P. falciparum was conducted²⁵ in age-matched female immunodeficient NOD.Cg-Prkdcscid Il2rgtm1Wjl/SzJ mice (8-10 weeks of age; 22-24 g) supplied by Charles River, UK, under licence of The Jackson Laboratory, Bar Harbor. Mice were engrafted with human erythrocytes (Red Cross Transfusion Blood Bank in Madrid, Spain) by daily intraperitoneal injection with 1 ml of a 50% haematocrit erythrocyte suspension (RPMI-1640 (Invitrogen), 25 mM HEPES (Sigma), 25% decomplemented AB+ human serum (Sigma) and 3.1 mM hypoxanthine (Sigma)). Mice with $\sim 40\%$ circulating human erythrocytes were intravenously infected with $2 \times 10^7 P$. falciparum Pf3D7^{0087/N9}-infected erythrocytes (day 0). Efficacy was assessed by administering one oral dose of E209 (2.5, 5, 15, 30, 50 and 100 mg kg⁻¹) at day 3 after infection. Treatment group assignments were allocated randomly. Parasitemia was measured by flow cytometry in samples of peripheral blood stained with the fluorescent nucleic acid dye SYTO-16 (Molecular Probes) and anti-murine erythrocyte TER119 monoclonal antibody (Becton Dickinson) in serial 2 µl blood samples taken every 24 h until assay completion. The ED90 was estimated by fitting a four parameter logistic equation using GraphPad 6.0 Software.

Systemic exposure measurement in infected *Pf* SCID mice. The levels of E209 were evaluated in whole blood to determine standard PK parameters in the individual animals used in the efficacy study. Peripheral blood samples (5 µg ml⁻¹) were taken at different times (0.25, 0.5, 1, 2, 4, 6, 8 and 23 h) after drug administration, mixed with 25 µl of Milli-Q water and immediately frozen on dry ice. The frozen samples were stored at – 80 °C until analysis. Vehicle-treated mice experienced the same blood-sampling regimen. Blood samples were processed by liquid–liquid extraction. Quantitative analysis by Liquid chromatography-tandem mass spectrometry (LC–MS/MS) was performed using a Waters UPLC system and Sciex API4000 mass spectrometer. The lower limit of quantification in this assay was 5 ng ml⁻¹. Blood concentration versus time was analysed by non-compartmental analysis (NCA) using Phoenix ver.6.3 (from Pharsight), from which exposure-related values (C_{max} and AUC₀₋₂₃, AUC_{0-t}) and t_{max} were estimated.

In vitro and *in vivo* PK studies. Details of the *in vitro* PK analysis of E209 are contained in Supplementary Tables 1, 2, 3, 17, 19 and 19). *In vitro* metabolic stability of E209 was carried out as described in Supplementary Methods.

In vivo PK studies in rats and mice were conducted using established procedures in accordance with the Australian Code of Practice for the Care and Use of Animals for Scientific Purposes, and the study protocols were reviewed and

approved by the Monash Institute of Pharmaceutical Sciences Animal Ethics Committee.

Rat PK studies were conducted in fasted 7–9-week-old male Sprague–Dawley rats, as previously described by Charman *et al.*¹² For intravenous administration E209 was dissolved in 5% glucose solution under sonication (for 10 min) prior to addition of Tween 80 and further sonication (5 min) to produce a clear solution. The formulation was filtered through a 0.22 µm syringe filter prior to dosing and aliquots retained to determine the dose administered. E209 was administered intravenously as a 10 min constant rate infusion via an indwelling jugular vein cannula (1 ml per rat, n = 2 rats).

For oral dosing, E209 was dissolved in 0.5% [w/v] hydroxypropyl methylcellulose containing 0.4% [w/v] Tween 80 and 0.5% [v/v] benzyl alcohol and sonicated for 5 min producing a clear solution. The bulk formulation was mixed by inverting the tubes prior to drawing each dosing volume. Aliquots of the dosing solution were retained for analysis to determine the dose administered. After the oral formulation (1 ml per rat, n=2 rats) was administered, an additional volume (1 ml) of Milli-Q water was administered (via a separate syringe).

Blood samples were collected using a Culex automated blood sampler over 48 h post-dosing into tubes containing anticoagulant and stabilization cocktail (Complete, potassium fluoride and EDTA) to minimize the potential for *ex vivo* degradation and immediately centrifuged at 4 °C. Plasma was separated, stored at -20 °C until processing and assayed by LC/MS using either a Waters Micromass Quattro Ultima PT triple quadrupole instrument coupled to a Waters 2795 HPLC or a Waters Micromass Quattro Premier triple quadrupole instrument coupled to a Waters Acquity UPLC.

The PKs of E209 was also studied in non-fasted 6-8 week old female Swiss Outbred mice that had access to food and water *ad libitum* throughout the pre- and post-dose sampling period. The formulations for intravenous and oral administration were the same as used for the rat studies (study protocols were reviewed and approved by the Monash Institute of Pharmaceutical Sciences Animal Ethics Committee).

E209 was administered intravenously by bolus tail vein injection (50 µl per mouse) or orally by gavage (10 ml kg⁻¹) and blood samples collected over 30 h post-dosing (n = 2 mice per time point). A maximum of two samples were obtained from each mouse, with samples being taken either via submandibular bleed (conscious sampling) or terminal cardiac puncture (underinhaled isofluorane anaesthesia). Blood was collected into tubes in the same way as described for the rat studies and immediately centrifuged and stored at -20 °C until analysis by LC–MS.

Plasma PK parameters were calculated based on compartmental PK analysis using Pmetrics⁴⁶ as described in Supplementary Methods. A classical three compartment oral absorption model was used for all tested species due to the tri-phasic profiles displayed over time. For human predictions, allometric scaling was performed based on weight. Clearance rates (including inter-compartmental clearance rates Q1 and Q2) were predicted according to the following equations:

$$CL_{human} = CL_{animal} \times \left[\frac{Humanweight}{Animalweight}\right]^{0.75}$$
(1)

$$Q_{human} = Q_{animal} \times \left[\frac{Humanweight}{Animalweight} \right]^{0.75}$$
(2)

Other PK parameters were scaled linearly and simulation of human exposure to TDD-E209 was performed assuming a 15 mg kg $^{-1}$ dose using Pmetrics where median values were plotted over time on a semi-log scale.

hERG profiling and assessment of cytotoxicity. The potencies (IC_{50} values) for E209 to inhibit the cardiac I_{Kr} (hERG), $Cav_{1.2}$ and $Nav_{1.5}$ channels were determined in an electrophysiology-based assays using IonWorks HT (CHO cells)²⁷.

In vivo (rat) safety studies. A single, rising oral dose study to determine the MTD of E209 and aid in the selection of dose levels for the 7-day toxicity study. The rats (Sprague–Dawley, age 6–7 weeks) were divided into four groups, each with three male and three female. Dose levels employed were 30, 100, 300 and 1,000 mg kg⁻¹ and the MTD was shown to be 300 mg kg⁻¹.

Following this single dose MTD study, rats were dosed once a day at 0, 30, 100 and 300 mg kg $^{-1}$ po E209 for 7 days. E209 was formulated in 0.5 % [w/v] HPMC, 0.4% Tween 80 [w/v], 0.5% [v/v] benzyl alcohol in purified water. In each group there were eight Sprague–Dawley rats of each sex. At the end of the dosing period, five rats/sex/group were killed (day 8) and the remaining three rats/sex/group were left for recovery. In addition, six rats of each sex per group were allocated for toxicokinetics.

In vivo PK studies conformed to AAALAC International and NIH guidelines as reported in the Guide for the Care and Use of Laboratory Animals, National Research Council (2011); People's Republic of China, Ministry of Science & Technology, 'Regulations for the Administration of Affairs Concerning Experimental Animals', 1988.

Data availability. All relevant data are available from the authors upon request.

References

- WHO. Global Report on Antimalarial Drug Efficacy and Drug Resistance: 2000–2010 (WHO, 2010).
- 2. WHO. Guidelines for the Treatment of Malaria 2nd edn (WHO, 2010).
- Wells, T. N., van Huijsduijnen, R. H. & Van Voorhis, W. C. Malaria medicines: a glass half full? *Nat. Rev. Drug. Discov.* 14, 424–442 (2015).
- Noedl, H. et al. Evidence of artemisinin-resistant malaria in western Cambodia. N. Engl. J. Med. 359, 2619–2620 (2008).
- Dondorp, A. M. et al. Artemisinin resistance in Plasmodium falciparum malaria. N. Engl. J. Med. 361, 455–467 (2009).
- 6. Bruce-Chwatt, L. J. et al. Chemotherapy of Malaria 2nd edn (World Health Organization, 1986).
- 7. WHO. *Chemotherapy of Malaria*. Report of a WHO Scientific Group. WHO Technical Report Series, No. 375 (World Health Organization, 1967).
- WHO. Status Report on Artemisinin Resistance. Available at: (http:// www.who.int/malaria/publications/atoz/status-rep-artemisinin-resistancesep2014.pdf)(2014).
- Leang, R. et al. Evidence of Plasmodium falciparum malaria multidrug resistance to artemisinin and piperaquine in western Cambodia: dihydroartemisinin-piperaquine open-label multicenter clinical assessment. Antimicrob. Agents Chemother. 59, 4719–4726 (2015).
- Gomes, G. d. P., Vil, V., Terent'ev, A. & Alabugin, I. V. Stereoelectronic source of the anomalous stability of bis-peroxides. *Chem. Sci.* 6, 6783–6791 (2015).
- O'Neill, P. M. et al. Identification of a 1,2,4,5-tetraoxane antimalarial drugdevelopment candidate (RKA 182) with superior properties to the semisynthetic artemisinins. Angew. Chem. Int. Ed. Engl. 49, 5693–5697 (2010).
- Charman, S. A. *et al.* Synthetic ozonide drug candidate OZ439 offers new hope for a single-dose cure of uncomplicated malaria. *Proc. Natl Acad. Sci. USA* 108, 4400–4405 (2011).
- 13. Karyana, M. et al. Malaria morbidity in Papua Indonesia, an area with multidrug resistant Plasmodium vivax and Plasmodium falciparum. Malar. J. 7, 148 (2008).
- Ratcliff, A. et al. Therapeutic response of multidrug-resistant Plasmodium falciparum and P. vivax to chloroquine and sulfadoxine-pyrimethamine in southern Papua, Indonesia. Trans. R. Soc. Trop. Med. Hyg. 101, 351–359 (2007).
- 15. Russell, B. *et al.* Determinants of *in vitro* drug susceptibility testing of *Plasmodium vivax. Antimicrob. Agents Chemother.* **52**, 1040–1045 (2008).
- Marfurt, J. et al. Ex vivo activity of histone deacetylase inhibitors against multidrug-resistant clinical isolates of *Plasmodium falciparum* and *P. vivax*. Antimicrob. Agents Chemother. 55, 961–966 (2011).
- Marfurt, J. et al. Ex vivo drug susceptibility of ferroquine against chloroquineresistant isolates of *Plasmodium falciparum* and *P. vivax*. Antimicrob. Agents Chemother. 55, 4461–4464 (2011).
- Vos, M. W. *et al.* A semi-automated luminescence based standard membrane feeding assay identifies novel small molecules that inhibit transmission of malaria parasites by mosquitoes. *Sci. Rep.* 5, 18704 (2015).
- Bolscher, J. M. *et al.* A combination of new screening assays for prioritization of transmission-blocking antimalarials reveals distinct dynamics of marketed and experimental drugs. *J. Antimicrob. Chemother.* **70**, 1357–1366 (2015).
- 20. Sanz, L. M. et al. P. falciparum in vitro killing rates allow to discriminate between different antimalarial mode-of-action. PLoS ONE 7, e30949 (2012).
- Witkowski, B. *et al.* Novel phenotypic assays for the detection of artemisininresistant *Plasmodium falciparum* malaria in Cambodia: *in-vitro* and *ex-vivo* drug-response studies. *Lancet Infect. Dis.* 13, 1043–1049 (2013).
- Witkowski, B. *et al.* Reduced artemisinin susceptibility of *Plasmodium falciparum* ring stages in western Cambodia. *Antimicrob. Agents Chemother.* 57, 914–923 (2013).
- Straimer, J. *et al.* Drug resistance. K13-propeller mutations confer artemisinin resistance in *Plasmodium falciparum* clinical isolates. *Science* 347, 428–431 (2015).
- 24. Peters, W. & Robinson, B. L. Malaria (Academic Press, 1999).
- 25. Angulo-Barturen, I. *et al.* A murine model of falciparum-malaria by *in vivo* selection of competent strains in non-myelodepleted mice engrafted with human erythrocytes. *PLoS ONE* **3**, e2252 (2008).
- 26. Jimenez-Diaz, M. B. et al. Improved murine model of malaria using Plasmodium falciparum competent strains and non-myelodepleted NOD-scid IL2R gamma(null) mice engrafted with human erythrocytes. Antimicrob. Agents Chemother. 53, 4533–4536 (2009).
- Rochford, R. *et al.* A humanized (hu)RBC-SCID mouse model of glucose 6-phosphate dehydrogenase deficiency for *in vivo* assessment of toxic hemolytic responses. *Proc. Natl Acad. Sci. USA* 110, 17486–17491 (2013).
- 28. Wang, J. et al. Haem-activated promiscuous targeting of artemisinin in *Plasmodium falciparum. Nat. Commun.* **6**, 10111 (2015).
- Ismail, H. M. *et al.* Probing the molecular targets of artemisinins in the malaria parasite: a chemical proteomic approach. *Proc. Natl Acad Sci. USA* 113, 2080–2085 (2016).

- 30. Ismail, H. M. *et al.* A click chemistry-based proteomic approach reveals that 1,2,4-trioxolane and artemisinin antimalarials share a common protein alkylation profile. *Angew. Chem. Int. Ed. Engl.* **55**, 6401–6405 (2016).
- 31. Dogovski, C. *et al.* Targeting the cell stress response of *Plasmodium falciparum* to overcome artemisinin resistance. *PLoS Biol.* **13**, e1002132 (2015).
- 32. Morris, C. A. *et al.* Review of the clinical pharmacokinetics of artesunate and its active metabolite dihydroartemisinin following intravenous, intramuscular, oral or rectal administration. *Malar. J.* **10**, 263 (2011).
- 33. Jiménez-Díaz, M. B. et al. (+)-SJ733, a clinical candidate for malaria that acts through ATP4 to induce rapid host-mediated clearance of *Plasmodium*. Proc. Natl Acad Sci. USA 111, E5455–E5462 (2014).
- Rottmann, M. et al. Spiroindolones, a new and potent chemotype for the treatment of malaria. Science 329, 1175–1180 (2010).
- Coteron, J. M. *et al.* Structure-guided lead optimization of triazolopyrimidinering substituents identifies potent *Plasmodium falciparum* dihydroorotate dehydrogenase inhibitors with clinical candidate potential. *J. Med. Chem.* 54, 5540–5561 (2011).
- 36. Trager, W. & Jensen, J. B. Human malaria parasites in continuous culture. *Science* **193**, 673–675 (1976).
- Winter, R. W. et al. Evaluation and lead optimization of anti-malarial acridones. Exp. Parasitol. 114, 47–56 (2006).
- Desjardins, R. E. et al. Quantitative assessment of antimalarial activity In vitro by a semiautomated microdilution technique. Antimicrob. Agents Chemother. 16, 710–718 (1979).
- Snyder, C. *et al. In vitro* and *in vivo* interaction of synthetic peroxide RBx11160 (OZ277) with piperaquine in *Plasmodium* models. *Exp. Parasitol.* 115, 296–300 (2007).
- Copple, I. M. *et al.* Examination of the cytotoxic and embryotoxic potential and underlying mechanisms of next-generation synthetic trioxolane and tetraoxane antimalarials. *Mol. Med.* 18, 1045–1055 (2012).
- 41. Zeeman, A.-M. et al. KAI407, a potent non-8-aminoquinoline compound that kills Plasmodium cynomolgi early dormant liver stage parasites in vitro. Antimicrob. Agents Chemother. 58, 1586–1595 (2014).
- Dembele, L. *et al.* Persistence and activation of malaria hypnozoites in longterm primary hepatocyte cultures. A potential role for the epigenetic control of quiescence. *Nat. Med.* 20, 307–312 (2014).
- McNamara, C. W. et al. Targeting Plasmodium PI(4)K to eliminate malaria. Nature 504, 248–253 (2013).
- 44. Baragana, B. et al. A novel multiple-stage antimalarial agent that inhibits protein synthesis. Nature 522, 315–320 (2015).
- 45. Almela, M. J. *et al.* A new set of chemical starting points with *Plasmodium falciparum* transmission-blocking potential for antimalarial drug discovery. *PloS ONE* **10**, e0135139 (2015).
- 46. Neely, M. N., van Guilder, M. G., Yamada, W. M., Schumitzky, A. & Jelliffe, R. W. Accurate detection of outliers and subpopulations with Pmetrics, a nonparametric and parametric pharmacometric modeling and simulation package for R. *Ther. Drug. Monit.* 34, 467–476 (2012).

Acknowledgements

This work was supported by grants from the European Union (Antimal, FP6, S.A.W., P.M.O'N.), the Medicines for Malaria Venture (S.A.W., P.M.O'N., G.A.B., G.L.N., R.K.A.), the NIH (R01 AI109023 to D.A.F.), the Global Health and Innovative Technology (GHIT) Fund (G2015-120, S.A.W., P.M.O'N., G.A.B, G.L.N.) and Medical Research Council, United Kingdom (G0700654, M.H.-L.W., A.E.M., I.M.C.). We would like to thank Dr Sergio Wittlin (Swiss Tropical and Public Health Institute Socinstrasse 57, 4051 Basel, Switzerland) for the data recorded in Table 1 (*Plasmodium berghei* mouse model).

Author contributions

P.M.O'N., S.A.W., G.A.B., G.L.N., D.A.F., S.F.B., I.A.-B., G.A., S.A.C., B.C. wrote the manuscript. P.M.O'N., G.L.N., R.K.A., S.S., E.R.S., N.L.R., M.H.-L.W., C.R., C.P., M.J.P., B.M., W.D.H., M.S., S.C. completed the chemistry, design of compounds, synthetic routes and analysis. J.D. carried out the *In vitro* antimalarial efficacy experiments. G.W., R.N., R.P., J.M. performed the clinical field isolate work; E.R., D.S., G.A., R.S., D.A.S. completed PK/PD modelling and stability studies. F.J.G., L.S. completed the *in vitro* anternative feeding Assay SMFA work. F.J.G., L.S. completed the PRR assays. I.A.-B., M.B.J.-D. completed the *in vivo* efficacy work in SCID mice. S.F.B., M.S.M. undertook the PK experiments in SCID mice. D.F. designed the RSA0-3h assays, which were performed by N.G. and J.S., R.S., M.D. and A.R. performed the transmission-blocking studies and TRA analysis. S.A.C., E.R., D.M.S. performed *in vitro* and *in vivo* PK studies. R.R. designed and performed G6PD deficient mice experiment. B.C. managed the CRO work, project management and toxicology studies. I.M.C., A.E.M., P.S. carried out *in vitro* and *in vivo* toxicology studies. C.K., A.-M.Z. performed liver stage drug assays in primary rhesus monkey hepatocytes

Additional information

Supplementary Information accompanies this paper at http://www.nature.com/ naturecommunications

Competing interests: The authors declare no competing financial interests.

Reprints and permission information is available online at http://npg.nature.com/ reprintsandpermissions/

How to cite this article: O'Neill, P. M. et al. A tetraoxane-based antimalarial drug candidate that overcomes PfK13-C580Y dependent artemisinin resistance. Nat. Commun. 8, 15159 doi: 10.1038/ncomms15159 (2017).

Publisher's note: Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

۲ (∞)

This work is licensed under a Creative Commons Attribution 4.0 International License. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder to reproduce the material. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/

© The Author(s) 2017