

# Physiologically-based Pharmacokinetic (PBPK) Modeling for Prediction of the Optimal Dose Regimens of Quinine and Phenobarbital Co-administration in Adult Patients with Cerebral Malaria and Seizures

**Teerachat Saeheng**

Thammasat University

**Juntra Karbwang**

Thammasat University

**Rajith Kumar Reddy Rajoli**

University of Liverpool

**Marco Siccardi**

University of Liverpool

**Kesara Na-Bangchang** (✉ [kesaratmu@yahoo.com](mailto:kesaratmu@yahoo.com))

Chulabhorn International College of Medicine <https://orcid.org/0000-0001-6389-0897>

---

## Research

**Keywords:** Physiologically-based pharmacokinetic (PBPK) modeling, quinine, phenobarbital, cerebral malaria, seizure, cytochrome P450, drug-drug interactions

**Posted Date:** August 7th, 2020

**DOI:** <https://doi.org/10.21203/rs.3.rs-53474/v1>

**License:** © ⓘ This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

---

**Title page****Physiologically-based pharmacokinetic (PBPK) modeling for prediction of the optimal dose regimens of quinine and phenobarbital co-administration in adult patients with cerebral malaria and seizures**

Teerachat Saeheng<sup>1</sup>, Juntra Karbwang<sup>1,2</sup>, Rajith Kumar Reddy Rajoli<sup>3</sup>, Marco Siccardi<sup>3</sup>,  
and Kesara Na-Bangchang<sup>1,2\*</sup>

<sup>1</sup>. Center of Excellence in Pharmacology and Molecular Biology of Malaria and  
Cholangiocarcinoma, Chulabhorn International College, Thammasat University,  
Pathumthani, Thailand

<sup>2</sup>. Drug Discovery and Development Center, Office of Advanced Science and Technology,  
Thammasat University, Thailand

<sup>3</sup>. Department of Molecular and Clinical Pharmacology, University of Liverpool, United  
Kingdom

**Correspondence**

K Na-Bangchang, Chulabhorn International College of Medicine, Thammasat University,  
Rangsit Campus, 99 Moo 18 Phaholyothin Road, Klong Luang District, Pathumthani 12121,  
Thailand

**Email:** kesaratmu@yahoo.com

1  
2  
3  
4  
5  
6  
7  
8  
9  
10  
11  
12  
13  
14  
15  
16  
17  
18  
19  
20  
21  
22  
23  
24  
25

## 26 Abstract

27 **Background:** Cerebral malaria is a fatal disease. Patients with cerebral malaria are at risk of  
28 seizure development, therefore, the co-administration of antimalarial and antiepileptic drugs  
29 are needed. Quinine and phenobarbital are standard drugs for the treatment of cerebral  
30 malaria with seizures. However, there is no information on the optimal dosage regimens of  
31 both drugs when used concomitantly. The study applied physiologically-based  
32 pharmacokinetic (PBPK) modeling for prediction of the optimal dose regimens of quinine  
33 and phenobarbital when co-administered in patients with cerebral malaria and concurrent  
34 seizures who carry wild type and polymorphic cytochrome P450 (CYP450) *2C9/2C19*.

35 **Methods:** The whole-body PBPK models for quinine and phenobarbital were constructed  
36 based on the previously published information using Simbiology<sup>®</sup>. One hundred virtual  
37 population were simulated. Four published articles were used for model verification.  
38 Sensitivity analysis was carried out to determine the effect of the changes in model  
39 parameters on AUC<sub>0-72h</sub>. Simulation of optimal dose regimens was based on standard drug-  
40 drug interactions (DDIs), and actual clinical use study approaches.

41 **Results:** Dose adjustment of the standard regimen of phenobarbital is not required when co-  
42 administered with quinine. The proposed optimal dose regimen for quinine, when co-  
43 administered with phenobarbital for patients with a single or continuous seizure in all  
44 malaria-endemic areas regardless of *CYP2C9/CYP2C19* genotypes, is a loading dose of 1,500  
45 mg IV infusion over 8 hours, followed by 1,200 mg infusion over 8 hours given three times  
46 daily, or multiple doses of 1,400 mg IV infusion over 8 hours, given three times daily. In  
47 areas with quinine resistance, the dose regimen should be increased as a loading dose of  
48 2,000 mg IV infusion over 8 hours, followed by 1,750 mg infusion over 8 hours given three  
49 times daily.

50 **Conclusion:** The developed PBPK models are reliable, and successfully predicted the  
51 optimal doses regimens of quinine-phenobarbital co-administration with no requirement of  
52 *CYP2C9/CYP2C19* genotyping.

53 **Keywords**

54 Physiologically-based pharmacokinetic (PBPK) modeling, quinine, phenobarbital, cerebral  
55 malaria, seizure, cytochrome P450, drug-drug interactions

56

57

58

59

60

61

62

63

64

65

66

67

68

69

70

71

72

73

74

## 75 **Background**

76 Cerebral malaria remains a high burden neurological problem in sub-Saharan African  
77 countries with an increased risk of seizures [1]. Quinine is the standard antimalarial drug for  
78 the treatment of severe malaria, including cerebral malaria [1]. Phenobarbital, a cost-effective  
79 antiepileptic drug, is the standard treatment for cerebral malaria patients with seizures [2, 3].  
80 Phenobarbital-induced severe cutaneous adverse reactions are, however, an issue of concern  
81 for clinical use of this drug [4]. Information on the contribution of host genetics on such  
82 reactions in cerebral malaria patients with seizures has been limited. Since phenobarbital  
83 induces xenobiotic drug-metabolizing enzymes cytochrome P450 (CYP450), the situation is  
84 further complicated when it is co-administered with drugs that are also metabolized by  
85 CYP450 enzymes. Phenobarbital is metabolized mainly in the livers by the polymorphic  
86 isoforms -- CYP2C9 and CYP2C19 [5, 6]. Quinine, on the other hand, is metabolized by  
87 CYP3A4 and UDP-glucuronosyltransferase 1A1 (UGT1A1) enzymes [7]. The activity of  
88 both enzymes is induced by phenobarbital [8, 9]. Cerebral malaria patients with polymorphic  
89 *CYP2C9/CYP2C19* genotypes who receive concomitant treatment with quinine and  
90 phenobarbital are, therefore, at risk of inadequate or toxic therapeutic drug concentrations  
91 due to metabolic drug interactions. To our knowledge, there have been few reports on the  
92 optimal phenobarbital dose for patients with epilepsy [5, 6], but not for cerebral malaria  
93 patients with seizures who carry polymorphic *CYP2C9* and *CYP2C19*. Besides, the optimal  
94 dosage of quinine when co-administered with phenobarbital has not been reported in this  
95 group of patients.

96 Physiologically-based pharmacokinetic (PBPK) modeling and simulation are accepted  
97 by various regulatory authorities as a promising tool to support dose optimization in the  
98 clinical phase of drug development, particularly for the investigation of drug-drug  
99 interactions (DDIs) and non-DDIs [10]. The present study aimed to apply PBPK modeling

100 and simulation for optimization of quinine and phenobarbital co-administration in adult  
101 patients with cerebral malaria and seizures, with consideration of genetic polymorphisms of  
102 *CYP2C9/CYP2C19* and DDIs.

## 103 **Methods**

### 104 **Model construction**

105 The whole-body PBPK models for quinine and phenobarbital (alone and co-administration)  
106 were constructed based on the previously published information [11, 12] using Simbiology®  
107 (version 5.8.2), the product of MATLAB® (version 2019a) (MathWorks, Natick, MA, USA).  
108 The physicochemical and biochemical properties (model parameters) of each drug, including  
109 human physiological parameters, were obtained from the published articles are available in  
110 the supplementary material of this article [9, 11, 13-25] (Additional file 1).

111 Model assumptions included the limitation of blood-flow, immediate dissolution of each  
112 drug in the gastrointestinal tract, absence of drug absorption in the stomach and large  
113 intestine, absence of enterohepatic recirculation, and absence of effect of 3-hydroxyquinine  
114 (quinine metabolite) on quinine disposition.

115

### 116 **Model verification**

117 Four published articles for quinine [26-29] and one article for phenobarbital [30] were used  
118 for model verification. The published data were extracted using Plot digitizer® version 2.6.8  
119 (Free Software Foundation, Inc., Boston, MA, USA). The area under the plasma  
120 concentration-time curve (AUC) was calculated based on the trapezoidal rule using the Excel  
121 spreadsheet. The simulated results from the developed models were compared against the  
122 published data, using the accepted criterion --absolute average-folding errors (AAFEs) of  $\pm$   
123 2-fold [31]. The mathematical equation for AAFEs is given below:

$$124 \text{AAFE} = 10^{(\sum \text{abs}(\log(\text{prediction}/\text{observation}))/n)}$$

125           Where n is the number of samples. The prediction is the simulated results from the  
126 developed model, and the observation is the published clinical data.

127

128 **Sensitivity analysis (the study of the uncertainty of certain model input parameters on**  
129 **the model output)**

130 Sensitivity analysis was performed to determine the effect of the changes in model  
131 parameters on the AUC during the first 72 hours ( $AUC_{0-72h}$ ) following the standard  
132 intravenous (IV) regimen of quinine (quinine model) or phenobarbital (phenobarbital model)  
133 or quinine and phenobarbital co-administration (DDIs model). The model parameters for  
134 sensitivity analysis of the quinine model were a fraction of unbound drug ( $f_u$ ), a fraction of  
135 UGT1A1 on total metabolism ( $f_{m,UGT1A1}$ ), a fraction of CYP3A4 on total metabolism  
136 ( $f_{m,CYP3A4}$ ), inhibitor concentration that produces half-maximal inhibition ( $K_{i3A4}$ ), and  
137 blood-to-plasma partition ratio ( $R_{bp}$ ). The model parameters for phenobarbital were  $f_u$ ,  $R_{bp}$ ,  
138 half-maximal effective concentrations ( $EC_{50,2C9}$ ), maximum effect at the maximum  
139 concentration ( $EC_{max,2C9}$ ), a fraction of CYP2C9 on total metabolism ( $f_{m,CYP2C9}$ ), and a  
140 fraction of CYP2C19 on total drug metabolism ( $f_{m,CYP2C19}$ ). The model parameters for DDIs  
141 included  $E_{max,3A4}$ ,  $EC_{50,3A4}$ ,  $EC_{50,UGT1A1}$ , and  $E_{max,UGT1A1}$ . The effect of the changes in model  
142 parameters on the  $AUC_{0-72h}$  was determined by the change of each model parameter within  
143  $\pm 20\%$ . The mathematical equation for sensitivity analysis is given below:

144 Sensitivity coefficient =  $\%Y/\%X$

145           Where  $\%Y$  is the percent change of the  $AUC_{0-72h}$  and  $\%X$  is the percent change of the  
146 model parameter.

147

148

149

## 150 **Virtual population simulation**

151 One hundred virtual population (50 males and 50 females, aged 18-60 years, weighing 60 kg,  
152 fasting state) were simulated in (i) seizure patients with polymorphic *CYP2C9/CYP2C19*  
153 (phenobarbital model), (ii) patients with cerebral malaria (quinine model), and (iii) patients  
154 with cerebral malaria and seizures with polymorphic *CYP2C9/CYP2C19* (DDIs model). The  
155 intrinsic clearance of phenobarbital in each genotype was obtained from the published  
156 clinical data for wild type [*CYP2C9* extensive metabolizer (EM)/*CYP2C19*EM] [6],  
157 *CYP2C9\*1/\*1CYP2C19\*1/\*3* [*CYP2C9*EM/*CYP2C19* poor metabolizer (PM)] or  
158 *CYP2C9/CYP2C19\*2/\*2* (*CYP2C9*EM/*CYP2C19*PM) or *CYP2C9/CYP2C19\*3/\*3*  
159 (*CYP2C9*EM/*CYP2C19*PM) [5, 6], and *CYP2C9\*1/\*3/CYP2C19\*1/\*1*  
160 (*CYP2C9*PM/*CYP2C19*EM) [32].

161

## 162 **Quinine and phenobarbital dose regimens used in simulations**

163 The standard regimen of quinine for severe malaria is the loading dose of 20 mg kg<sup>-1</sup> (1,000  
164 mg base total dose for the average 60 kg body weight) IV infusion over 4 hours, followed by  
165 the maintenance dose of 10 mg kg<sup>-1</sup> (500 mg base total dose for the average 60 kg body  
166 weight) IV infusion over 4 hours, given three times daily (every 8 hours) [1] for 72 hours.  
167 The simulation time used was based on the average time for patients with cerebral malaria to  
168 regain consciousness (72 hours) [28].

169

## 170 **DDIs model simulation**

171 *Simulation based on standard DDIs study approach:* For standard DDIs study approach,  
172 plasma concentration-time profiles of phenobarbital and quinine following the co-  
173 administration of standard dose regimen of phenobarbital and quinine were simulated.  
174 Phenobarbital is given at 2 mg kg<sup>-1</sup>day<sup>-1</sup> (1-3 mg.kg<sup>-1</sup>.day<sup>-1</sup>) or 120 mg total dose for the

175 average 60 kg body weight) IV infusion over 30 minutes for 17 consecutive days. The  
176 standard dose regimen of quinine for three days (described above) was given on day 14 of  
177 phenobarbital administration when steady-state plasma concentration was achieved.

178 *Simulation based on actual clinical use approach:* For the PBPK simulation based on actual  
179 clinical use study approach, two simulated scenarios were applied with the total simulation  
180 time of 72 hours. The scenario-I applies for patients who have only a single seizure;  
181 phenobarbital (15 mg kg<sup>-1</sup> or 900 mg total dose for the average 60 kg body weight, IV  
182 infusion over 30 minutes) is given as a single dose 6 hours after the first dose of quinine  
183 (average time of occurrence of seizure after admission) [33]. The scenario-II applies for  
184 patients who have continuous seizures; phenobarbital at a loading dose of 15 mg kg<sup>-1</sup> or 900  
185 mg total dose for the average 60 kg body weight IV infusion over 30 minutes, followed by  
186 the maintenance dose of 1.5 mg kg<sup>-1</sup>day<sup>-1</sup> (1-3 mg.kg<sup>-1</sup> day<sup>-1</sup>) or 90 mg total dose for average  
187 60 kg body weight IV infusion over 30 minutes [33] is given every 24 hours, starting 6 hours  
188 after the first dose of quinine until 72 hours. The time of simulation and seizure frequency  
189 was based on the clinical report [34]. The predicted optimal dosage regimens were presented  
190 as amount of quinine base.

191

### 192 **Criteria for optimal dose regimens**

193 The optimal dose regimens of quinine for adult patients with cerebral malaria with seizures  
194 were proposed based on the therapeutic range of quinine, *i.e.*, maximum plasma  
195 concentration ( $C_{\max}$ )  $\leq 20$  mg.L<sup>-1</sup>, and trough plasma concentration ( $C_{\text{trough}}$ )  $\geq 10$  mg.L<sup>-1</sup> [35].

196 The optimal dosage of phenobarbital were proposed based on the therapeutic range of  
197 phenobarbital:  $C_{\max} \leq 40$  mg.L<sup>-1</sup> [36], and  $C_{\text{trough}} \geq 15$  mg.L<sup>-1</sup> [33]. The predicted  
198 pharmacokinetic parameters are presented as mean (95% confidence interval or CI).

199

200

**201 Results****202 Model verification**

203 The AAFEs for quinine and phenobarbital, quinine alone, and phenobarbital alone ranged  
204 from 1.17 (1.03-1.21) [26-30], 1.14 (1.03-1.21) [26-29], and 1.20 [30], respectively. All  
205 were within accepted ranges [31], indicating the reliability of the PBPK models (Table 1).  
206 Visual Predictive Checks (VPCs) between predicted results and published data are shown in  
207 Additional file 2.

208

**209 Sensitivity analysis**

210 Sensitivity coefficient analysis values for  $AUC_{0-72h}$  of quinine, phenobarbital, and quinine  
211 and phenobarbital co-administration ranged from -0.72 to +0.14, -0.02 to +0.10, and -0.48 to  
212 +0.12, respectively. All values were less than one, indicating no significant impact of the  
213 model parameters on model construction.

214

**215 Simulation of standard dose regimen of phenobarbital in patients with seizures with  
216 polymorphic CYP2C9 and CYP2C19**

217 *Simulation based on standard DDIs study approach:* Results of the simulation of optimal  
218 dose regimens of phenobarbital ( $C_{max}$ ,  $C_{trough}$ , and clearance) in patients with wild type  
219 (*CYP2C9EM/CYP2C19EM*) and polymorphic *CYP2C9EM/CYP2C19PM*, and  
220 *CYP2C9PM/CYP2C19EM* are summarized in Table 2. The average values of all parameters  
221 in all genotypes were within the therapeutic range.

222 *Simulation based on actual clinical use study approach:* Results of the simulation of optimal  
223 dose regimens (single and multiple dosing) of phenobarbital ( $C_{max}$ ,  $C_{trough}$ , and clearance) in  
224 patients with wild type (*CYP2C9EM/CYP2C19EM*) and polymorphic

225 *CYP2C9EM/CYP2C19PM*, and *CYP2C9PM/CYP2C19EM* are summarized in Table 3. The  
226 average values of all parameters in all genotypes were within the therapeutic range.

227 **Simulation of the optimal dose of quinine when co-administered with phenobarbital in**  
228 **cerebral malaria patients with concomitant seizures and polymorphic**

229 ***CYP2C9/CYP2C19***

230 Results ( $C_{\text{trough}}$ ,  $C_{\text{max}}$ , AUC ratio or AUCR, and  $C_{\text{max}}$  ratio or  $C_{\text{max}}R$ ) of the simulation of the  
231 standard dose of quinine, when co-administered with phenobarbital in cerebral malaria  
232 patients with concomitant seizures (*scenario I*: single seizure, and *scenario II*: continuous  
233 seizures) and polymorphic *CYP2C9/CYP2C19* based on DDI and actual clinical use study  
234 approaches, are summarized in Table 4.

235 *Simulation based on standard DDIs study approach*: Standard dose regimen of quinine  
236 provided inappropriate plasma drug concentrations when co-administered with phenobarbital  
237 ( $C_{\text{trough}} < 10 \text{ mg.L}^{-1}$ ) (Table 4). The initially proposed quinine (*regimen-1*: a loading dose of  
238 2,200 mg IV infusion over 4 hours, followed by maintenance doses of 1,200 mg IV infusion  
239 over 4 hours given three times daily) in patients with wild type and polymorphic  
240 *CYP2C9/CYP2C19* provided 2.2-fold lower plasma drug concentrations compared with  
241 standard quinine regimen, with AUCR ranging from 0.42 to 0.45. This 2.2-fold increase of  
242 quinine standard dose provided the  $C_{\text{max}}$  exceeding  $20 \text{ mg.L}^{-1}$ , but the  $C_{\text{trough}}$  lower than 10  
243  $\text{mg.L}^{-1}$  both in patients with wild type and polymorphic *CYP2C9/CYP2C19* (Fig. 1, and Fig.  
244 2 for wild type and polymorphic *CYP2C9/CYP2C19*, respectively) (Table 5). The time to  
245 reach therapeutic concentration ranged from 2 to 3 hours. The three subsequent dose  
246 regimens were therefore simulated, *e.g.*, a loading dose of 2,200 mg IV infusion, followed by  
247 2,000 mg IV infusion (*regimen-2*), a loading dose of 2,000 mg IV infusion, followed by  
248 1,500 mg IV infusion (*regimen-3*), and a loading dose of 1,750 mg IV infusion, followed by  
249 1,500 mg IV infusion (*regimen-4*). All regimens were given as IV infusion over 8 hours, and

250 the maintenance doses were given three times daily. The average  $C_{\max}$  and  $C_{\text{trough}}$  (95%CI) for  
251 each regimen (-2, -3, and -4) are shown in Fig. 1 (wild type), and Fig. 2 (polymorphic  
252 *CYP2C9/CYP2C19*) (Table 5). The time to reach therapeutic concentration ranged from 4 to  
253 6 hours.

254 *Simulation based on actual clinical use study approach:* Standard dose regimen of quinine  
255 provided inappropriate plasma drug concentrations when co-administered with phenobarbital  
256 ( $C_{\text{trough}} < 10 \text{ mg.L}^{-1}$ ) (Table 4). Dose adjustment in both clinical scenarios based on standard  
257 dosage regimen of quinine provided the AUCR of quinine when co-administered with  
258 phenobarbital in patients with wild type genotype, and polymorphic *CYP2C9/CYP2C19*  
259 ranging from 0.47 to 0.53 (Table 4). The 2-fold increase of quinine standard dose (*regimen-*  
260 *5*: a loading dose of 2,000 mg IV infusion over 4 hours, followed by maintenance doses of  
261 1,000 mg IV infusion over 4 hours given three times daily) provided the concentrations out  
262 of therapeutic range ( $C_{\max} > 20 \text{ mg.L}^{-1}$ , and  $C_{\text{trough}} < 10 \text{ mg.L}^{-1}$ ) both in patients with wild  
263 type and polymorphic *CYP2C9/CYP2C19* (Fig. 3 (scenario I and II), Fig. 4 (scenario I), and 5  
264 (scenario II) for wild type and polymorphic *CYP2C9/CYP2C19*, respectively) (Table 6 and 7  
265 for scenario I and scenario II, respectively). The three subsequent dose regimens for the  
266 scenario I (single seizure) and II (multiple seizures) were therefore simulated, *i.e.*, a loading  
267 dose of 2,000 mg IV infusion, followed by 1,750 mg IV infusion three times daily (*regimen-*  
268 *6*), a loading dose of 1,500 mg IV infusion, followed by 1,200 mg IV infusion three times  
269 daily (*regimen-7*), and multiple doses of 1,400 mg IV infusion three times daily (*regimen-8*).  
270 The IV infusion duration in all regimens was 8 hours, and the maintenance doses were given  
271 three times daily. The average  $C_{\max}$ , and  $C_{\text{trough}}$  (95%CI) of quinine for wild type genotype in  
272 each simulation are shown in Fig. 3, and those for polymorphic *CYP2C9/CYP2C19* for the  
273 *scenario I* and *scenario II* are presented in Fig. 4 and Fig. 5, respectively. Table 5, 6, and 7  
274 summarize the parameters predicted based on standard DDI and actual clinical use (single

275 and continuous seizures) study approach, respectively. Time to reach therapeutic quinine  
276 levels ranged from 4 to 6 hours. Plasma quinine concentrations following *regimen-7* and *-8*  
277 except *regimen-5 and -6* were within the therapeutic range for both *scenarios I* and *scenario*  
278 *II*. The  $C_{max}$  of quinine following *regimen-6* in both scenarios ranged from 21-23 mg.L<sup>-1</sup>.

279

## 280 **Discussion**

281 Results of the current study based on PBPK modeling and simulation raise a concern about  
282 the potential DDIs between quinine and phenobarbital in patients with cerebral malaria who  
283 have seizures and require concurrent treatment with both drugs. The conventional dose  
284 adjustment based on the AUCR of both drugs in different clinical scenarios may provide  
285 suboptimal dose regimens with inadequate trough plasma levels, which pose the patients at  
286 risk of treatment failure and/or severe complication. The PBPK modeling approach, on the  
287 other hand, has proved a promising tool for dose optimization of quinine and phenobarbital  
288 co-administration.

289

## 290 **Optimal phenobarbital dose regimens in patients with seizures with polymorphic**

### 291 ***CYP2C9* and *CYP2C19***

292 Simulation of the optimal phenobarbital dose regimens in patients with seizures who carry  
293 polymorphic *CYP2C9/CYP2C19* was investigated using dose regimens based on the two  
294 approaches, *i.e.*, the standard DDIs study approach (at steady-state of phenobarbital level),  
295 and the actual clinical use study approach (scenario I for single seizure and scenario II for  
296 continuous seizures). Results of the present study supported the previous report of the  
297 decrease in total clearance of phenobarbital by 48.2% [32] and 20% [5, 6] in patients carrying  
298 *CYP2C9PM/CYP2C19EM* and *CYP2C9EM/CYP2C19PM* genotypes, respectively. The  
299 reported frequencies of the wild type, *CYP2C9PM/CYP2C19EM*, and

300 *CYP2C9*EM/*CYP2C19*PM genotypes in the Thai population are 42% [37], 2.8% [38], and  
301 13% [37], respectively. Based on the results of PBPK modeling using both DDIs and actual  
302 clinical use study approaches, however, dosage adjustment of phenobarbital may not be  
303 required as plasma drug concentrations were maintained within the therapeutic range ( $C_{\max} \leq$   
304  $40 \text{ mg}\cdot\text{L}^{-1}$  [36], and  $C_{\text{trough}} \geq 15 \text{ mg}\cdot\text{L}^{-1}$  [33]) (Table 2 and 3). The proposed phenobarbital  
305 dosage regimens are optimal for the treatment of patients with single seizure (single dose of  
306 900 mg or  $15 \text{ mg}\cdot\text{kg}^{-1}$ ), as well as patients with cerebral malaria who have continuous  
307 seizures (a loading dose of  $15 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{day}^{-1}$ , followed by  $1.5 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{day}^{-1}$  once-daily)  
308 regardless of patients' *CYP2C9/CYP219* genotypes. Genotyping is therefore not necessary,  
309 which is practical both in developed and developing countries. Besides, the advantage of  
310 using phenobarbital over other anticonvulsants is its relatively low cost [2].

311

312 **Optimal quinine dose regimens when co-administered with phenobarbital in cerebral**  
313 **malaria patients with concomitant seizures and polymorphic *CYP2C9/CYP2C19***

314 Similarly to phenobarbital, simulation of the optimal quinine dose regimens in patients with  
315 concurrent cerebral malaria and seizures with polymorphic *CYP2C9/CYP2C19* were  
316 investigated using dose regimens based on standard DDIs study and actual clinical use  
317 (scenario I and II) study approaches. Dose optimization based on AUCR yielded undesirable  
318 plasma quinine concentrations when co-administered with phenobarbital.

319 *Simulations based on standard DDIs approach:* The three proposed quinine dosage  
320 regimens, *i.e.*, *regimen-2*, *-3* and *-4*) provided adequate  $C_{\text{trough}}$  above  $10 \text{ mg}\cdot\text{L}^{-1}$  in both  
321 wildtype genotype (Fig 1B, C, D for *regimen-2*, *3*, and *-4*, respectively) and polymorphic  
322 *CYP2C9/CYP2C19* (Fig 2B, C, D for *regimen-2*, *-3*, and *-4*, respectively). Nevertheless,  
323 *regimen-2* (a loading dose of 2,200 mg IV infusion over 8 hours, followed by 2,000 mg IV  
324 infusion over 8 hours given three times daily) is not appropriate for clinical use due to high

325 drug concentrations above  $20 \text{ mg}\cdot\text{L}^{-1}$  in individuals with both wild type and polymorphic  
326 *CYP2C9/CYP2C19* and thus, the possibility of toxicity. Since the reported minimum  
327 inhibitory concentration (MIC) of quinine in quinine-resistant areas has been rising to over 10  
328  $\text{mg}\cdot\text{L}^{-1}$  [39], the most optimal dose regimen would be *regimen-3* (a loading dose of 2,000 mg  
329 IV infusion over 8 hours, followed by 1,500 mg IV infusion over 8 hours given three times  
330 daily). In low quinine resistant malaria-endemic areas, *regimen-4* (a loading dose of 1,750  
331 mg IV infusion over 8 hours, followed by 1,500 mg IV infusion three times daily) could be  
332 an alternative regimen. It is noted, however, that the infusion duration of 8 hours might result  
333 in the delay of time to reach therapeutic level compared with the recommended standard  
334 regimens (4-6 and 2-3 hours for the recommended and standard regimens, respectively).  
335 Since the critical clinical period for treatment of patients with cerebral malaria is during the  
336 first 24 hours [27], such delay is unlikely to pose the patients at risk of complicated  
337 manifestation or death. These quinine regimens can be co-administered with phenobarbital  
338 without consideration of *CYP2C9/CYP2C19* genotypes.

339 *Simulations based on actual clinical use study approach:* Optimal  $C_{\text{max}}$  and  $C_{\text{trough}}$  of  
340 quinine were achieved with the two proposed quinine dose regimens (*regimen-7* and *-8*)  
341 when co-administered with phenobarbital in both clinical scenarios (scenario I for a single  
342 seizure and scenario II for continuous seizures) using PBPK modeling and simulation, but not  
343 the AUCR, resulted in adequate plasma concentrations (Fig. 3, Fig. 4, and Fig. 5 for wild  
344 type, the scenario I in polymorphic *CYP2C9/CYP2C19*, and scenario II in polymorphic  
345 *CYP2C9/CYP2C19*, respectively). With the administration time of quinine 6 hours prior to  
346 phenobarbital in actual clinical study approach, the optimal quinine dosage regimens can be  
347 reduced as a loading dose of 2,000 mg IV infusion over 8 hours, followed by 1,750 mg IV  
348 infusion three times daily (*regimen-6*), or alternatively, 1,500 mg IV infusion, followed by  
349 1,200 mg IV infusion (*regimen-7*), or 1,400 mg IV infusion over 8 hours given three times

350 daily (*regimen-8*). These quinine regimens can be co-administered with phenobarbital  
351 without consideration of *CYP2C9/CYP2C19* genotypes since plasma quinine concentrations  
352 in patients with wild type, and polymorphic *CYP2C9/CYP2C19* were comparable. There is  
353 no influence of *CYP2C9/CYP2C19* genotypes on the inducing effect of quinine metabolism  
354 since the steady-state drug concentrations are not achieved with a short duration of  
355 phenobarbital dosing. The possibility of dose reduction could be due to the lack of CYP450  
356 enzyme-inducing effect of phenobarbital during this period (6 hours). In the case of quinine  
357 resistant malaria, particularly with the contribution of large interindividual variability in  
358 quinine clearance, quinine *regimen-6* would be the best option. The higher  $C_{max}$  of 1-3 mg.L<sup>-1</sup>  
359 above the therapeutic range is unlikely to cause toxicity due to high plasma protein binding  
360 during the acute phase of malaria infection. It is noted that the recommended optimal dose  
361 regimens of quinine and phenobarbital co-administration apply for cerebral malaria patients  
362 with seizures who have normal hepatic function but not in those with impaired function.  
363 Therapeutic drug monitoring for quinine in those patients is recommended.

364 The limitations of the study include the exclusion of the contribution of P-  
365 glycoprotein transporter on quinine disposition (due to lack of information on *in vitro* studies),  
366 as well as the inhibitory effect of 3-hydroxyquinine metabolite on CYP3A4 activity.  
367 Nevertheless, the significant impacts of these two factors on quinine disposition are unlikely  
368 [11].

369 In summary, PBPK models are a potential tool for dose optimization of quinine in  
370 patients with cerebral malaria in resource-limited countries. The developed PBPK models for  
371 phenobarbital and quinine-phenobarbital co-administration are reliable, and successfully  
372 predicted the optimal doses regimens of phenobarbital in cerebral malaria patients with single  
373 or continuous seizures with no requirement of *CYP2C9/CYP2C19* genotyping. Dose  
374 adjustment based on PBPK modeling but not AUCR provided desirable plasma quinine

375 concentrations. Dose adjustment of the standard regimen of phenobarbital is not required  
376 when co-administered with quinine. The proposed optimal dose regimen for quinine when  
377 co-administered with phenobarbital for patients with a single seizure (scenario I), and  
378 continuous seizures (scenario II) in all malaria-endemic areas regardless of  
379 *CYP2C9/CYP2C19* genotypes is a loading dose of 1,500 mg IV infusion over 8 hours,  
380 followed by 1,200 mg IV infusion over 8 hours given three times daily (*regimen-7*), or  
381 multiple doses of 1,400 mg IV infusion over 8 hours, given three times daily (*regimen-8*). In  
382 areas with quinine resistance, the dose regimen should be increased as a loading dose of  
383 2,000 mg IV infusion over 8 hours, followed by 1,750 mg IV infusion over 8 hours given  
384 three times daily (*regimen-6*).

385

### 386 **Abbreviations**

387 95%CI: 95% confident interval; AAFEs: absolute average-folding errors; AUC: the area  
388 under the plasma concentration-time curve; AUCR: AUC ratio;  $C_{max}$ : peak plasma  
389 concentration;  $C_{max}R$ :  $C_{max}$  ratio;  $C_{trough}$ : trough plasma concentration; CYP450: cytochrome  
390 P450; DDIs: drug-drug interactions;  $EC_{50}$ : half-maximal effective concentrations; EM:  
391 extensive metabolizer;  $E_{max}$ : maximum effect at the maximum concentration;  $f_m$ : fraction of  
392 metabolism;  $f_u$ : fraction of unbound drug; IV: intravenous;  $K_i$ : inhibitor concentration that  
393 produces half-maximal inhibition;  $R_{bp}$ : blood-to-plasma ratio; PM: poor metabolizer;  
394 UGT1A1: UDP-glucuronosyltransferase 1A1; VPC: visual predictive check.

395

### 396 **Acknowledgements**

397 K.N. receives research support from the National Research Council of Thailand and  
398 Thammasat University (Center of Excellence in Pharmacology and Molecular Biology of

399 Malaria and Cholangiocarcinoma). J.K. receives research support from Thammasat  
400 University (Bualuang ASEAN Chair Professor).

401

#### 402 **Authors's contributions**

403 T.S., J.K., M.S., and K.N. designed the study. M.S., T.S., and R.K.R. collected the  
404 information. T.S., R.K.R., and M.S. performed research. T.S., J.K., and K.N. wrote the  
405 manuscript. All authors had accessed and interpreted the data. All authors approved the final  
406 version.

407

#### 408 **Funding**

409 The study was supported by Thammasat University under the Center of Excellence in  
410 Pharmacology and Molecular Biology of Malaria and Cholangiocarcinoma (TU-CoE 005-  
411 2018).

412

#### 413 **Availability of data and materials**

414 All data generated or analysed during this study are included in this published article [and its  
415 supplementary information files].

416

#### 417 **Ethical approval and consent to participate**

418 Not applicable

419

#### 420 **Consent for publication**

421 Not applicable

422

#### 423 **Competing interests**

424 The authors declare that they have no competing interests.

425

426

427

428

429

### 430 **References**

431 1. Idro R, Jenkins NE, Newton CR: Pathogenesis, clinical features, and neurological  
432 outcome of cerebral malaria. *Lancet Neurol* 2005, 4:827-840.

433 2. Chisholm D, Who C: Cost-effectiveness of first-line antiepileptic drug treatments in  
434 the developing world: a population-level analysis. *Epilepsia* 2005, 46:751-759.

435 3. Meremikwu M, Marson AG: Routine anticonvulsants for treating cerebral malaria.  
436 *Cochrane Database Syst Rev* 2002:CD002152.

437 4. Manuyakorn W, Siripool K, Kamchaisatian W, Pakakasama S, Visudtibhan A,  
438 Vilaiyuk S, Rujirawat T, Benjaponpitak S: Phenobarbital-induced severe cutaneous  
439 adverse drug reactions are associated with CYP2C19\*2 in Thai children. *Pediatr*  
440 *Allergy Immunol* 2013, 24:299-303.

441 5. Mamiya K, Hadama A, Yukawa E, Ieiri I, Otsubo K, Ninomiya H, Tashiro N,  
442 Higuchi S: CYP2C19 polymorphism effect on phenobarbitone. *Pharmacokinetics in*  
443 *Japanese patients with epilepsy: analysis by population pharmacokinetics. Eur J Clin*  
444 *Pharmacol* 2000, 55:821-825.

445 6. Yukawa E, Mamiya K: Effect of CYP2C19 genetic polymorphism on  
446 pharmacokinetics of phenytoin and phenobarbital in Japanese epileptic patients using  
447 Non-linear Mixed Effects Model approach. *J Clin Pharm Ther* 2006, 31:275-282.

- 448 7. Mirghani RA, Hellgren U, Bertilsson L, Gustafsson LL, Ericsson O: Metabolism and  
449 elimination of quinine in healthy volunteers. *Eur J Clin Pharmacol* 2003, 59:423-427.
- 450 8. Ramirez J, Komoroski BJ, Mirkov S, Graber AY, Fackenthal DL, Schuetz EG, Das S,  
451 Ratain MJ, Innocenti F, Strom SC: Study of the genetic determinants of UGT1A1  
452 inducibility by phenobarbital in cultured human hepatocytes. *Pharmacogenet*  
453 *Genomics* 2006, 16:79-86.
- 454 9. Almond LM, Mukadam S, Gardner I, Okialda K, Wong S, Hatley O, Tay S, Rowland-  
455 Yeo K, Jamei M, Rostami-Hodjegan A, Kenny JR: Prediction of Drug-Drug  
456 Interactions Arising from CYP3A induction Using a Physiologically Based Dynamic  
457 Model. *Drug Metab Dispos* 2016, 44:821-832.
- 458 10. Shebley M, Sandhu P, Emami Riedmaier A, Jamei M, Narayanan R, Patel A, Peters  
459 SA, Reddy VP, Zheng M, de Zwart L, et al: Physiologically Based Pharmacokinetic  
460 Model Qualification and Reporting Procedures for Regulatory Submissions: A  
461 Consortium Perspective. *Clin Pharmacol Ther* 2018, 104:88-110.
- 462 11. Saeheng T, Na-Bangchang K, Siccardi M, Rajoli RKR, Karbwang J: Physiologically-  
463 Based Pharmacokinetic Modeling for Optimal Dosage Prediction of Quinine  
464 Coadministered With Ritonavir-Boosted Lopinavir. *Clin Pharmacol Ther* 2019.
- 465 12. Rajoli RK, Back DJ, Rannard S, Freel Meyers CL, Flexner C, Owen A, Siccardi M:  
466 Physiologically Based Pharmacokinetic Modelling to Inform Development of  
467 Intramuscular Long-Acting Nanoformulations for HIV. *Clin Pharmacokinet* 2015,  
468 54:639-650.
- 469 13. Qalaaquin (FDA product Label)  
470 [[https://www.accessdata.fda.gov/drugsatfda\\_docs/label/2010/021799s0111bl.pdf](https://www.accessdata.fda.gov/drugsatfda_docs/label/2010/021799s0111bl.pdf)]

- 471 14. Dua VK, Sarin R, Prakash A: Determination of quinine in serum, plasma, red blood  
472 cells and whole blood in healthy and Plasmodium falciparum malaria cases by high-  
473 performance liquid chromatography. *J Chromatogr* 1993, 614:87-93.
- 474 15. Supanaranond W, Davis TM, Pukrittayakamee S, Silamut K, Karbwang J, Molunto P,  
475 Chanond L, White NJ: Disposition of oral quinine in acute falciparum malaria. *Eur J*  
476 *Clin Pharmacol* 1991, 40:49-52.
- 477 16. Koudriakova T, Iatsimirskaia E, Utkin I, Gangl E, Vouros P, Storozhuk E, Orza D,  
478 Marinina J, Gerber N: Metabolism of the human immunodeficiency virus protease  
479 inhibitors indinavir and ritonavir by human intestinal microsomes and expressed  
480 cytochrome P4503A4/3A5: mechanism-based inactivation of cytochrome P4503A by  
481 ritonavir. *Drug Metab Dispos* 1998, 26:552-561.
- 482 17. Strauch S, Dressman JB, Shah VP, Kopp S, Polli JE, Barends DM: Biowaiver  
483 monographs for immediate-release solid oral dosage forms: Quinine sulfate. *Journal*  
484 *of Pharmaceutical Sciences* 2012, 101:499-508.
- 485 18. Zhao XJ, Ishizaki T: A further interaction study of quinine with clinically important  
486 drugs by human liver microsomes: determinations of inhibition constant ( $K_i$ ) and type  
487 of inhibition. *Eur J Drug Metab Pharmacokinet* 1999, 24:272-278.
- 488 19. Wallin A, Hengren L: Distribution of phenobarbital in whole blood during pregnancy  
489 and perinatally--an in vitro study. *Eur J Clin Pharmacol* 1985, 29:187-191.
- 490 20. Ehrnebo M, Odar-Cederlof I: Binding of amobarbital, pentobarbital and  
491 diphenylhydantoin to blood cells and plasma proteins in healthy volunteers and  
492 uraemic patients. *Eur J Clin Pharmacol* 1975, 8:445-453.
- 493 21. Bender AD, Post A, Meier JP, Higson JE, Reichard G, Jr.: Plasma protein binding of  
494 drugs as a function of age in adult human subjects. *J Pharm Sci* 1975, 64:1711-1713.

- 495 22. Xu Y, Zhou Y, Hayashi M, Shou M, Skiles GL: Simulation of clinical drug-drug  
496 interactions from hepatocyte CYP3A4 induction data and its potential utility in trial  
497 designs. *Drug Metab Dispos* 2011, 39:1139-1148.
- 498 23. Varma MV, Gardner I, Steyn SJ, Nkansah P, Rotter CJ, Whitney-Pickett C, Zhang H,  
499 Di L, Cram M, Fenner KS, El-Kattan AF: pH-Dependent solubility and permeability  
500 criteria for provisional biopharmaceutics classification (BCS and BDDCS) in early  
501 drug discovery. *Mol Pharm* 2012, 9:1199-1212.
- 502 24. Shou M, Hayashi M, Pan Y, Xu Y, Morrissey K, Xu L, Skiles GL: Modeling,  
503 prediction, and in vitro in vivo correlation of CYP3A4 induction. *Drug Metab Dispos*  
504 2008, 36:2355-2370.
- 505 25. Wilensky AJ, Friel PN, Levy RH, Comfort CP, Kaluzny SP: Kinetics of  
506 phenobarbital in normal subjects and epileptic patients. *Eur J Clin Pharmacol* 1982,  
507 23:87-92.
- 508 26. Salako LA, Sowunmi A: Disposition of quinine in plasma, red blood cells and saliva  
509 after oral and intravenous administration to healthy adult Africans. *Eur J Clin*  
510 *Pharmacol* 1992, 42:171-174.
- 511 27. White NJ, Looareesuwan S, Warrell DA, Warrell MJ, Bunnag D, Harinasuta T:  
512 Quinine pharmacokinetics and toxicity in cerebral and uncomplicated *Falciparum*  
513 malaria. *Am J Med* 1982, 73:564-572.
- 514 28. White NJ, Looareesuwan S, Warrell DA, Warrell MJ, Chanthavanich P, Bunnag D,  
515 Harinasuta T: Quinine loading dose in cerebral malaria. *Am J Trop Med Hyg* 1983,  
516 32:1-5.
- 517 29. Davis TM, White NJ, Looareesuwan S, Silamut K, Warrell DA: Quinine  
518 pharmacokinetics in cerebral malaria: predicted plasma concentrations after rapid

- 519 intravenous loading using a two-compartment model. *Trans R Soc Trop Med Hyg*  
520 1988, 82:542-547.
- 521 30. Nelson E, Powell JR, Conrad K, Likes K, Byers J, Baker S, Perrier D: Phenobarbital  
522 pharmacokinetics and bioavailability in adults. *J Clin Pharmacol* 1982, 22:141-148.
- 523 31. Saeheng T, Na-Bangchang K, Karbwang J: Utility of physiologically based  
524 pharmacokinetic (PBPK) modeling in oncology drug development and its accuracy: a  
525 systematic review. *Eur J Clin Pharmacol* 2018, 74:1365-1376.
- 526 32. Goto S, Seo T, Murata T, Nakada N, Ueda N, Ishitsu T, Nakagawa K: Population  
527 estimation of the effects of cytochrome P450 2C9 and 2C19 polymorphisms on  
528 phenobarbital clearance in Japanese. *Ther Drug Monit* 2007, 29:118-121.
- 529 33. Kokwaro GO, Ogutu BR, Muchohi SN, Otieno GO, Newton CR: Pharmacokinetics  
530 and clinical effect of phenobarbital in children with severe falciparum malaria and  
531 convulsions. *Br J Clin Pharmacol* 2003, 56:453-457.
- 532 34. Mohapatra MKD, L.K.; Mishra, N.R., et al.: Profile of seizures in adult falciparum  
533 malaria and clinical efficacy of phenytoin sodium for control seizures. *Asian Pac J*  
534 *Trop Dis* 2012, 2:5.
- 535 35. Verdier MC, Bentue-Ferrer D, Tribut O, pour le groupe Suivi Therapeutique  
536 Pharmacologique de la Societe Francaise de Pharmacologie et de T: [Therapeutic  
537 drug monitoring of quinine]. *Therapie* 2011, 66:507-516.
- 538 36. Patsalos PN, Berry DJ, Bourgeois BF, Cloyd JC, Glauser TA, Johannessen SI, Leppik  
539 IE, Tomson T, Perucca E: Antiepileptic drugs--best practice guidelines for therapeutic  
540 drug monitoring: a position paper by the subcommission on therapeutic drug  
541 monitoring, ILAE Commission on Therapeutic Strategies. *Epilepsia* 2008, 49:1239-  
542 1276.

- 543 37. Sukasem C, Tunthong R, Chamnanphon M, Santon S, Jantararoungtong T, Koomdee  
544 N, Prommas S, Puangpetch A, Vathesatogkit P: CYP2C19 polymorphisms in the Thai  
545 population and the clinical response to clopidogrel in patients with atherothrombotic-  
546 risk factors. *Pharmgenomics Pers Med* 2013, 6:85-91.
- 547 38. Sangviroon A, Panomvana D, Tassaneeyakul W, Namchaisiri J: Pharmacokinetic and  
548 pharmacodynamic variation associated with VKORC1 and CYP2C9 polymorphisms  
549 in Thai patients taking warfarin. *Drug Metab Pharmacokinet* 2010, 25:531-538.
- 550 39. Krudsood SC, Watcharee; Silachamroon, Udomsak; Phumratanaprapin, Weerapong;  
551 Viriyavejakul, Parnpen; Bussaratid, Valai; Looareesuwan, Sornchai: Clinical malaria  
552 and treatment of multidrug resistance falciparum in Thailand. *J Trop Med Hyg* 1999,  
553 27:7.

554

## 555 **Figures**

556 **Figure 1** Prediction of quinine dose regimens in wild type *CYP2C9/CYP2C19* based on  
557 standard DDIs approach. Prediction of quinine dosage regimens when co-administered with  
558 phenobarbital based on standard DDIs study approach in cerebral malaria patients with  
559 seizures who carry wild type *CYP2C9/CYP2C19*. Data are presented as mean (95%CI). Set  
560 criteria for dose optimization are  $C_{\max} \leq 20 \text{ mg.L}^{-1}$ , and  $C_{\text{trough}} \geq 10 \text{ mg.L}^{-1}$  (therapeutic range  
561 of quinine).

562

563 **Figure 2** Prediction of quinine dose regimens in polymorphic *CYP2C9/CYP2C19* based on  
564 standard DDIs approach. Prediction of quinine dosage regimens when co-administered with  
565 phenobarbital based on standard DDIs study approach in cerebral malaria patients with  
566 seizures who carry polymorphic *CYP2C9/CYP2C19*. Data are presented as mean (95%CI).

567 Set criteria for dose optimization are  $C_{\max} \leq 20 \text{ mg.L}^{-1}$ , and  $C_{\text{trough}} \geq 10 \text{ mg.L}^{-1}$  (therapeutic  
568 range of quinine).

569

570 **Figure 3** Prediction of quinine dose regimens in wild type *CYP2C9/CYP2C19* based on  
571 actual clinical use approach. Prediction of quinine dosage regimens when co-administered  
572 with phenobarbital based on actual clinical use study approach in cerebral malaria patients  
573 with seizures (scenario I: single, and scenario II: continuous) who carry wild type  
574 *CYP2C9/CYP2C19*. Data are presented as mean (95%CI). Set criteria for dose optimization  
575 are  $C_{\max} \leq 20 \text{ mg.L}^{-1}$ , and  $C_{\text{trough}} \geq 10 \text{ mg.L}^{-1}$ (therapeutic range of quinine).

576

577 **Figure 4** Prediction of quinine dose regimens in polymorphic *CYP2C9/CYP2C19* based on  
578 actual clinical use approach (scenario-I). Prediction of quinine dosage regimens when co-  
579 administered with phenobarbital based on actual clinical use study approach in cerebral  
580 malaria patients with a single seizure (scenario I) who carry polymorphic *CYP2C9/CYP2C19*.  
581 Data are presented as mean (95%CI). Set criteria for dose optimization are  $C_{\max} \leq 20 \text{ mg.L}^{-1}$ ,  
582 and  $C_{\text{trough}} \geq 10 \text{ mg.L}^{-1}$ (therapeutic range of quinine).

583

584 **Figure 5** Prediction of quinine dose regimens in polymorphic *CYP2C9/CYP2C19* based on  
585 actual clinical use approach (scenario-II). Prediction of quinine dosage regimens when co-  
586 administered with phenobarbital based on actual clinical use study approach in cerebral  
587 malaria patients with continuous seizures (scenario II) who carry polymorphic  
588 *CYP2C9/CYP2C19*. Data are presented as mean (95%CI). Set criteria for dose optimization  
589 are  $C_{\max} \leq 20 \text{ mg.L}^{-1}$ , and  $C_{\text{trough}} \geq 10 \text{ mg.L}^{-1}$ (therapeutic range of quinine).

590

591 **Tables**

592 **Table 1** Model verification of quinine and phenobarbital in cerebral malaria, severe malaria,  
593 and healthy adults. Comparisons of quinine and phenobarbital between predicted results and  
594 published data in patients with cerebral malaria, severe malaria, and healthy adults.

595 **Table 2** Simulated standard dose regimens of phenobarbital based on standard DDI study  
596 approach. Simulation of standard dose regimens of phenobarbital in patients with seizures  
597 with wild type and polymorphic *CYP2C9/CYP2C19* based on standard DDI study approach.

598 **Table 3** Simulated standard dose regimens of phenobarbital based on actual clinical use  
599 DDI study approach. Simulation of standard dose regimen of phenobarbital in patients with  
600 seizures with wild type and polymorphic *CYP2C9/CYP2C19* based on actual clinical use  
601 study approach

602 **Table 4** Simulated standard dose regimens of quinine when co-administered with  
603 phenobarbital based on actual clinical use approach. Simulation of the standard dose of  
604 quinine when co-administered with phenobarbital in cerebral malaria patients with  
605 concomitant seizures with wild type and polymorphic *CYP2C9/CYP2C19*.

606 **Table 5** Prediction of quinine dose regimens when co-administered with phenobarbital  
607 based on standard DDI study approach. Prediction of quinine dosage regimens when co-  
608 administered with phenobarbital in cerebral malaria patients with concomitant seizures and  
609 polymorphic *CYP2C9/CYP2C19* based on standard DDI study approach.

610 **Table 6** Prediction of quinine dose regimens when co-administered with phenobarbital based  
611 on scenario-I. Prediction of quinine dosage regimen when co-administered with  
612 phenobarbital in cerebral malaria patients with concomitant seizures with wild type and  
613 polymorphic *CYP2C9/CYP2C19* based on actual clinical study approach (*scenario I: single*  
614 *seizure*).

615 **Table 7** Prediction of quinine dose regimens when co-administered with phenobarbital based  
616 on scenario-II. Prediction of quinine dosage regimen when co-administered with

617 phenobarbital in cerebral malaria patients with concomitant seizures with wild type and  
618 polymorphic CYP2C9/CYP2C19 based on actual clinical study approach (*scenario II*:  
619 continuous seizures).

620

#### 621 **Additional files**

622 **Additional file 1: Table S1.** PBPK model input parameters for quinine and phenobarbital.

623 **Additional file 2: Figure S1.** A comparison of 500 mg quinine IV infusion over 4 hours in  
624 African healthy adults between predicted result and published data. **Figure S2.** A comparison

625 of 10 mg kg<sup>-1</sup> body weight quinine IV infusion over 4 hours in adult Thai patients with  
626 cerebral malaria between predicted result and published data. **Figure S3.** A comparison of 4

627 mg kg<sup>-1</sup> body weight quinine IV infusion over 4 hours in adult Thai patients with severe  
628 malaria between predicted result and published data. **Figure S4.** A comparison of loading

629 dose of 20 mg kg<sup>-1</sup> body weight quinine IV infusion over 4 hours, followed by 10 mg kg<sup>-1</sup>  
630 body weight IV infusion over 4 hours given 3 times daily in adult Thai patients with cerebral

631 malaria between predicted result and published data. **Figure S5.** A comparison of 10 mg kg<sup>-1</sup>  
632 body weight IV infusion over 4 hours given 3 times daily in adult Thai patients with cerebral

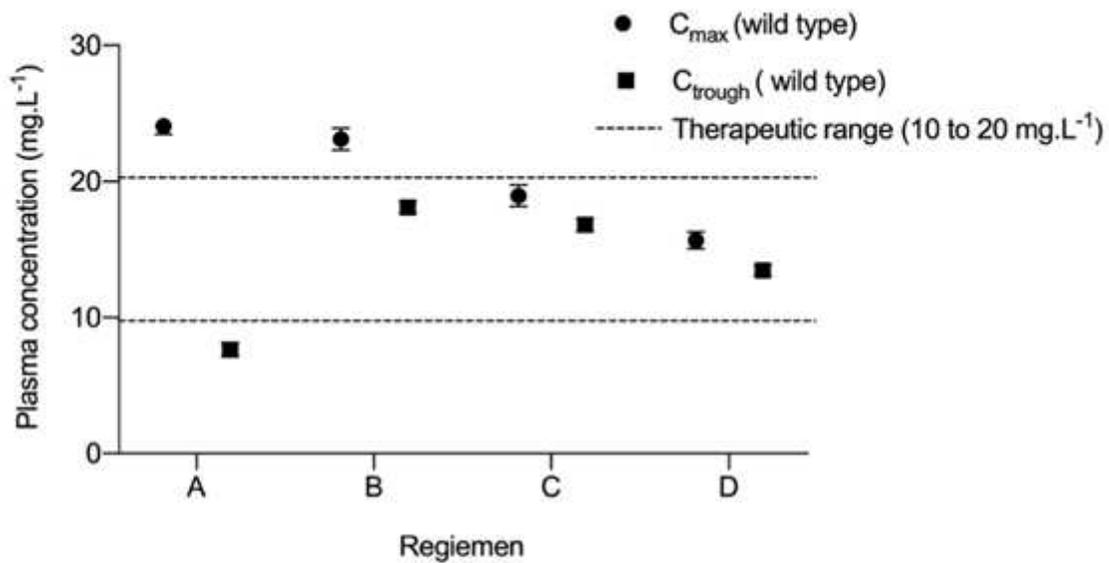
633 malaria between predicted result and published data. **Figure S6.** A comparison of 5 mg kg<sup>-1</sup>  
634 body weight IV infusion over 4 hours given 3 times daily in adult Thai patients with cerebral

635 malaria between predicted result and published data. **Figure S7.** A comparison of 2.6 mg kg<sup>-1</sup>  
636 (218 mg) phenobarbital IV infusion over 6 minutes given in healthy male subjects between

637 predicted result and published data.

638

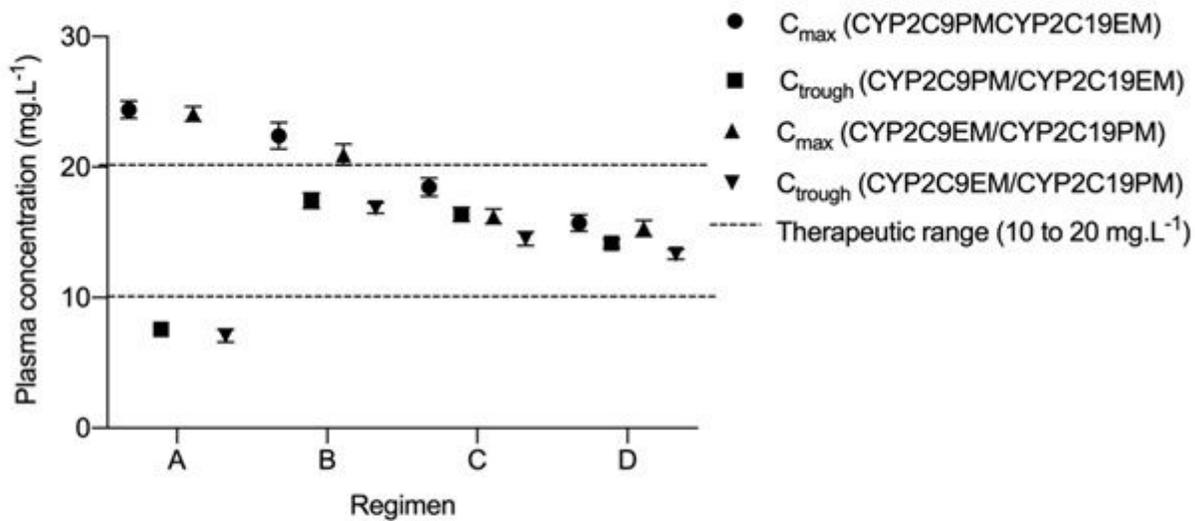
# Figures



- A (Regimen 1): Loading dose of 2200 mg IV infusion over 4 hours, followed by 1200 mg IV infusion over 4 hours (given every 8 hours daily)  
B: (Regimen 2): Loading dose of 2200 mg IV infusion over 8 hours, followed by 2000 mg IV infusion over 8 hours (given every 8 hours daily)  
C (Regimen 3): Loading dose of 2000 mg IV infusion over 8 hours, followed by 1500 mg IV infusion over 8 hours (given every 8 hours daily)  
D (Regimen 4): Loading dose of 1750 mg IV infusion over 8 hours, followed by 1500 mg IV infusion over 8 hours (given every 8 hours daily)

**Figure 1**

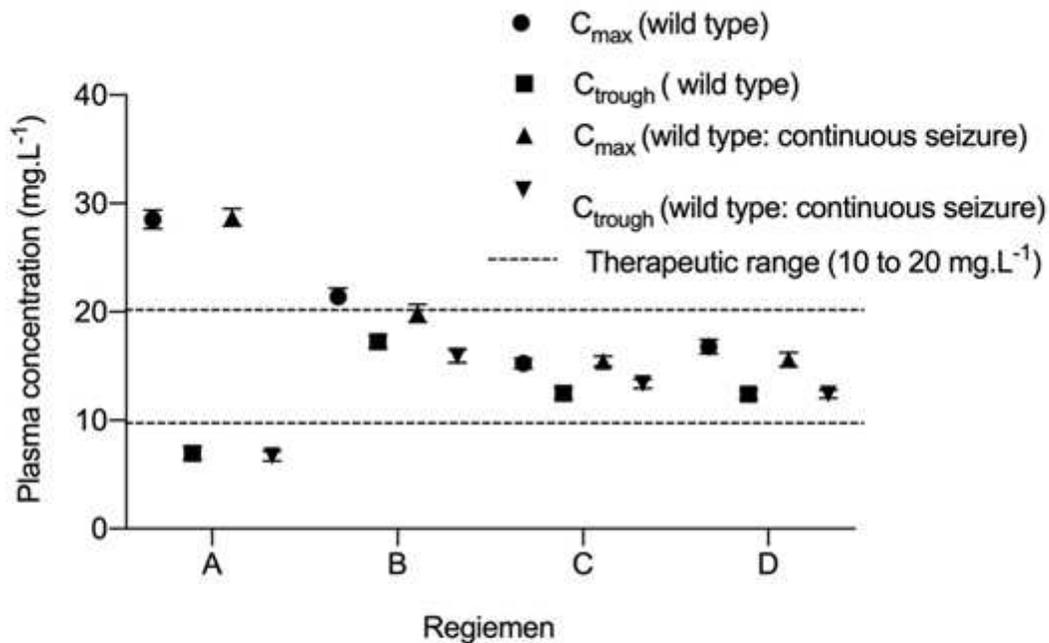
Prediction of quinine dose regimens in wild type CYP2C9/CYP2C19 based on standard DDIs approach. Prediction of quinine dosage regimens when co-administered with phenobarbital based on standard DDIs study approach in cerebral malaria patients with seizures who carry wild type CYP2C9/CYP2C19. Data are presented as mean (95%CI). Set criteria for dose optimization are C<sub>max</sub>  $\geq$  20 mg.L<sup>-1</sup>, and C<sub>trough</sub>  $\geq$  10 mg.L<sup>-1</sup> (therapeutic range of quinine).



- A (Regimen 1): Loading dose of 2200 mg IV infusion over 4 hours, followed by 1200 mg IV infusion over 4 hours (given every 8 hours daily)
- B: (Regimen 2): Loading dose of 2200 mg IV infusion over 8 hours, followed by 2000 mg IV infusion over 8 hours (given every 8 hours daily)
- C (Regimen 3): Loading dose of 2000 mg IV infusion over 8 hours, followed by 1500 mg IV infusion over 8 hours (given every 8 hours daily)
- D (Regimen 4): Loading dose of 1750 mg IV infusion over 8 hours, followed by 1500 mg IV infusion over 8 hours (given every 8 hours daily)

**Figure 2**

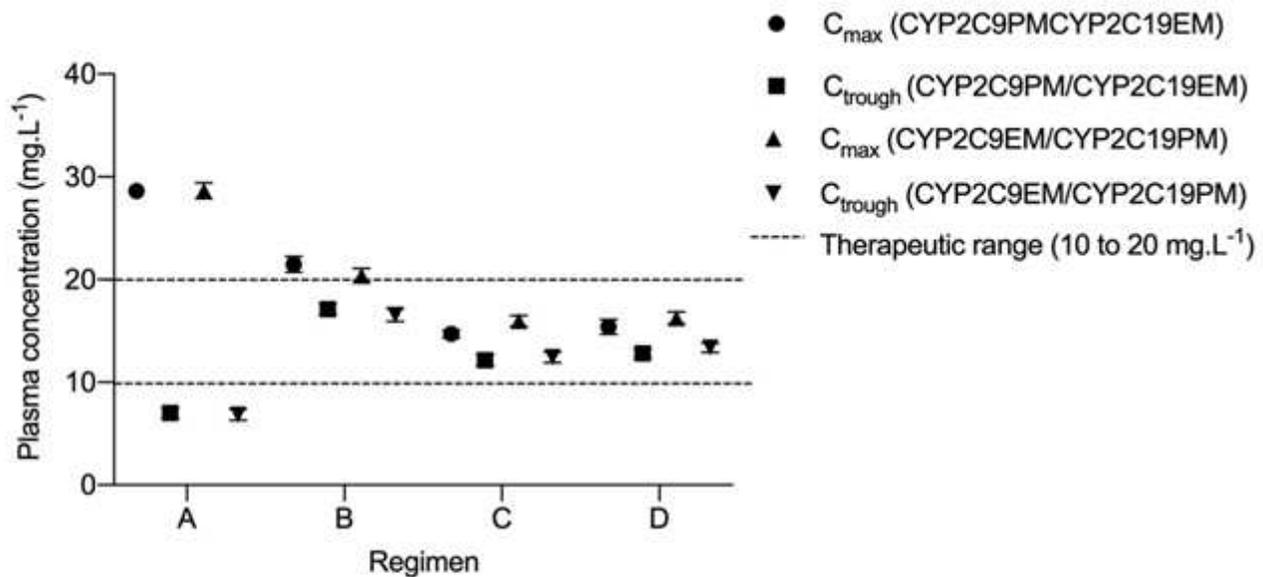
Prediction of quinine dose regimens in polymorphic CYP2C9/CYP2C19 based on standard DDIs approach. Prediction of quinine dosage regimens when co-administered with phenobarbital based on standard DDIs study approach in cerebral malaria patients with seizures who carry polymorphic CYP2C9/CYP2C19. Data are presented as mean (95%CI). Set criteria for dose optimization are  $C_{max} \geq 20$  mg.L<sup>-1</sup>, and  $C_{trough} \geq 10$  mg.L<sup>-1</sup> (therapeutic range of quinine).



A (Regimen 5): Loading dose of 2000 mg IV infusion over 4 hours, followed by 1000 mg IV infusion over 4 hours (given every 8 hours daily)  
 B: (Regimen 6): Loading dose of 2000 mg IV infusion over 8 hours, followed by 1750 mg IV infusion over 8 hours (given every 8 hours daily)  
 C (Regimen 7): Loading dose of 1500 mg IV infusion over 8 hours, followed by 1200 mg IV infusion over 8 hours (given every 8 hours daily)  
 D (Regimen 8): Multiple doses of 1400 mg IV infusion over 8 hours (given every 8 hours daily)

**Figure 3**

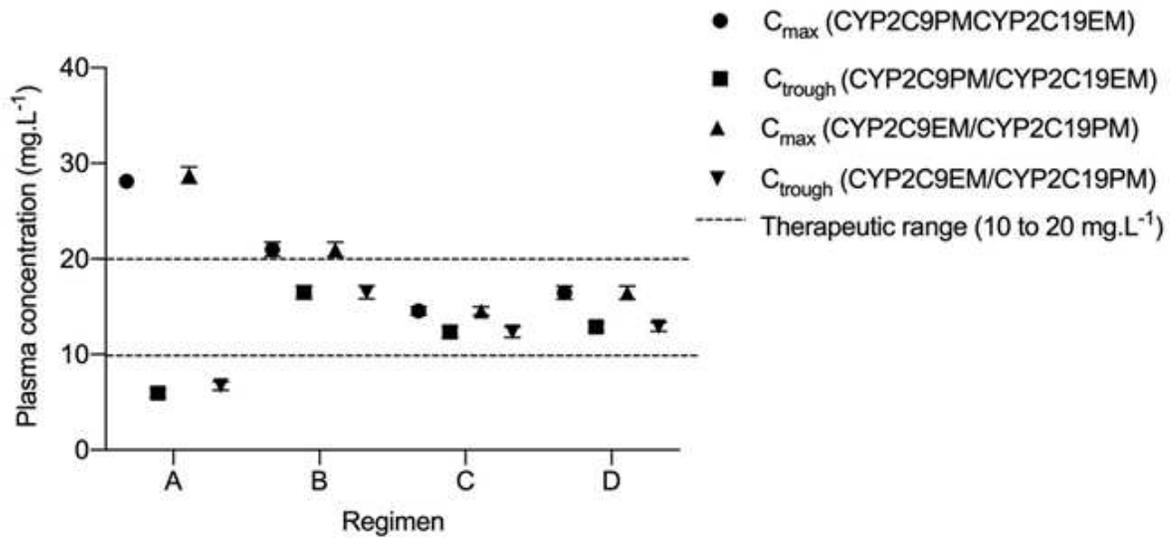
Prediction of quinine dose regimens in wild type CYP2C9/CYP2C19 based on actual clinical use approach. Prediction of quinine dosage regimens when co-administered with phenobarbital based on actual clinical use study approach in cerebral malaria patients with seizures (scenario I: single, and scenario II: continuous) who carry wild type CYP2C9/CYP2C19. Data are presented as mean (95%CI). Set criteria for dose optimization are  $C_{max} \approx 20$  mg.L<sup>-1</sup>, and  $C_{trough} \approx 10$  mg.L<sup>-1</sup> (therapeutic range of quinine).



A (Regimen 5): Loading dose of 2000 mg IV infusion over 4 hours, followed by 1000 mg IV infusion over 4 hours (given every 8 hours daily)  
 B: (Regimen 6): Loading dose of 2000 mg IV infusion over 8 hours, followed by 1750 mg IV infusion over 8 hours (given every 8 hours daily)  
 C (Regimen 7): Loading dose of 1500 mg IV infusion over 8 hours, followed by 1200 mg IV infusion over 8 hours (given every 8 hours daily)  
 D (Regimen 8): Multiple doses of 1400 mg IV infusion over 8 hours (given every 8 hours daily)

**Figure 4**

Prediction of quinine dose regimens in polymorphic CYP2C9/CYP2C19 based on actual clinical use approach (scenario-I). Prediction of quinine dosage regimens when co-administered with phenobarbital based on actual clinical use study approach in cerebral malaria patients with a single seizure (scenario I) who carry polymorphic CYP2C9/CYP2C19. Data are presented as mean (95%CI). Set criteria for dose optimization are  $C_{max} \geq 20$  mg.L<sup>-1</sup>, and  $C_{trough} \geq 10$  mg.L<sup>-1</sup>(therapeutic range of quinine).



A (Regimen 5): Loading dose of 2000 mg IV infusion over 4 hours, followed by 1000 mg IV infusion over 4 hours (given every 8 hours daily)  
 B: (Regimen 6): Loading dose of 2000 mg IV infusion over 8 hours, followed by 1750 mg IV infusion over 8 hours (given every 8 hours daily)  
 C (Regimen 7): Loading dose of 1500 mg IV infusion over 8 hours, followed by 1200 mg IV infusion over 8 hours (given every 8 hours daily)  
 D (Regimen 8): Multiple doses of 1400 mg IV infusion over 8 hours (given every 8 hours daily)

**Figure 5**

Prediction of quinine dose regimens in polymorphic CYP2C9/CYP2C19 based on actual clinical use approach (scenario-II). Prediction of quinine dosage regimens when co-administered with phenobarbital based on actual clinical use study approach in cerebral malaria patients with continuous seizures (scenario II) who carry polymorphic CYP2C9/CYP2C19. Data are presented as mean (95%CI). Set criteria for dose optimization are  $C_{max} \geq 20$  mg.L<sup>-1</sup>, and  $C_{trough} \geq 10$  mg.L<sup>-1</sup>(therapeutic range of quinine).

## Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [BPBKcerebralmalariaTablesMalariaJournal.docx](#)
- [Additionalfile1TableMalariaJournal.docx](#)
- [Additionalfile2FiguresMalariaJournal.docx](#)