

## INTRODUCTION

Malaria is the major cause of mortality and morbidity in the tropical and subtropical regions in the world. An estimated 300-500 million persons suffer from malaria every year and more than 1 million die each year. Majority of these cases and deaths particularly those in children occur in Sub Saharan Africa. Unlike some of the other acute diseases such as encephalitis, meningitis, and most of the chronic diseases, patients of severe malaria can recover completely without any long term effects if treated promptly and correctly. Therefore rationalization and standardization of treatment of cases of severe or uncomplicated malaria at different levels of health care is important. It has several advantages deaths can be reduced by effective use of standard treatment procedures. Patients who require hospitalization and those who need intensive care can be identified promptly and treated before they die or develop complications. The adoption of this approach of standard management can reduce the mortality and morbidity from malaria (WHO, 2004).

Approximately 800 – 1,000 malaria cases are admitted to Bangkok Hospital for Tropical Diseases annually. These include *P. falciparum* (51%), *P. vivax* (46%), mixed infections of *P. falciparum* and *P. vivax* (2%), few cases of *P. malariae* and occasional cases of *P. ovale*. Admitted patients are all treated with antimalarial regimens and most of them are enrolled for clinical trial (Faculty of Tropical Medicine, 2004).

Resistance to antimalarial drugs is increasing nearly everywhere in the tropical world, confounding global attempts to “Roll Back Malaria” (Nosten and Brasseur, 2002) Southeast Asia has the most resistant malaria parasites in the world, which has limited treatment options in this region (WHO, 2001). In Thailand, treatment of acute uncomplicated falciparum malaria is becoming more difficult because of increasing resistance to all antimalarial drugs, except the artemisinin derivatives (Wilairatana *et al*, 2002). To combat the further spread of resistance, it is generally accepted that combinations of antimalarial drugs that include an artemisinin derivative should be used, and, if possible, that preparations should be formulated in a single tablet (Hien *et al*, 2004).

The artemisinin derivatives (artesunate and a recently developed, dihydroartemisinin which is short acting but powerful drug) has been studied extensively in the treatment of falciparum malaria in Thailand, are well tolerated. Their main drawback is that conventional

courses (3-5 days) are associated with high rate of recrudescence, typically >25 %. In addition, there is the risk that parasite resistance will develop when antimalarial drugs are used alone (Warhurst, 1999). Because artemisinin derivatives are now the first-line treatment for multidrug-resistant falciparum malaria in many tropical countries, the appearance of artemisinin-resistant *Plasmodium falciparum* would have serious implications. The development of suitable combinations of an artemisinin compound with a second drug is therefore a priority (WHO, 2001). At present, artesunate has been registered by Thai FDA for use in the treatment of falciparum malaria.

Mefloquine is another antimalarial drug, which is better tolerated than quinine and can be administered during a day, but resistance to mefloquine has developed when used alone. Furthermore, in Thailand where multidrug resistance is encountered, a high dose (25 mg/kg) of mefloquine is recommended for use as a combination with other short acting antimalarial drugs (Nosten *et al*, 1991). Recently, clinical trials of artesunate combination with mefloquine has proved effective and well tolerated (Looareesuwan *et al*, 1992; Looareesuwan *et al*, 1994; Looareesuwan *et al*, 1996; Price *et al*, 1997), therefore this regimen has been chosen for treatment of multidrug resistant falciparum malaria in Thailand. However, some patients can not tolerate adverse effects of mefloquine.

Piperaquine phosphate (1,3-bis[1-(7-chloro-4'-quinoly)-4'-piperazinyl]) phosphate) replaced chloroquine as the recommended treatment for *Plasmodium falciparum* malaria in China in 1978 and was used extensively for mass prophylaxis and treatment. Reported adverse events are generally similar to those observed with chloroquine, although pruitus is uncommon. (Tropical Medicine Institute, 2003). Piperaquine proved to be effective and well tolerated, and no cross-resistance with chloroquine was observed (Chen *et al*, 1982). More recently, piperaquine has been used as part of short-course artemisinin based combination oral therapies designed to have a high cure rate, to have few side effects, and to reduce malaria transmission (Denis *et al*, 2002; Davis *et al*, 2005).

Artekin<sup>®</sup> (compound dihydroartemisinin) a combination of dihydroartemisinin 40mg, piperaquine 320mg per tablet (Batch No. 20011204 Mfg. 120401 Exp. 120403 supplied free of charge by Holleykin Pharmaceutical Co. Ltd., Guangzhou, China is claimed for high effective. In addition, this combination is well tolerated and convenient for use (3 days treatment). This compound has been on clinical trials and proved safe and well tolerated in

China, Vietnam, Laos, Cambodia and else where (Karunajeewa *et al*, 2003; Hien *et al*, 2004). We propose here a clinical trial of Artekin<sup>®</sup> vs artesunate and mefloquine (a standard regimen for treatment of multidrug resistant falciparum malaria in Thailand) at the Bangkok Hospital for Tropical Diseases to determine the safety, tolerability, and efficacy.

## MATERIALS AND METHODS

### Study site and recruitment procedures

All patients who fulfilled inclusion criteria (acute uncomplicated falciparum malaria, either male or female; if female, pregnancy test has to be negative before enrolment to the study, positive asexual forms of *P. falciparum* in blood smear, weight more than 40 kg and age more than 14 years, ability to take oral medication, agreement to stay in the hospital for at least 28 days). Informed consent to the study was obtained from patients, or their guardians, before enrolment to the study. The patients were admitted to the Bangkok Hospital for Tropical Diseases for 28 days to exclude reinfection and to assess the safety and efficacy of Artekin<sup>®</sup> and artesunate plus mefloquine. We excluded severe malaria according to WHO criteria (WHO, 2000), severe vomiting not allowing oral medication, pregnancy or lactating female, significant concomitant systemic diseases (for example systemic bacterial infections, liver and/or kidney insufficiencies, chronic disease or severe malnutrition), diseases requiring therapy except malaria, ingestion of other antimalarials in the past 14 days or presence of urine sulphonamides or 4-aminoquinolones. Clinical evaluation including neurological examination focused on brain stem, cerebellar function, muscle strength in all limb, extraocular and facial muscle strength, deep tendon reflexes, and finger-to-nose tests, and also parasite count were performed 12 hourly until negative then daily for 28 days. Malaria parasite count per microliter was obtained by calculation against the white blood cell count for a thick film. Geometric mean parasites were used as a standard method. Blood films were considered negative if no parasites were seen in 200 oil-immersion microscopic fields. Fever clearance time was taken as the period from the start of treatment until the oral temperature decreased to 37.5 °C and remained below this temperature for the next 48 hours. Side effects were defined as signs and symptoms that occurred or became more severe after treatment started. Cure rate at day 28 (cured patients/evaluable patients X 100%) was defined as the absence of parasite recrudescence during 28 days of follow-up. If there is RI, RII, or RIII failure (World Health Organization, 1973), standard antimalarial drugs of the hospital will be

given. Adverse events will also be treated as standard procedures at the Bangkok Hospital for Tropical Diseases. This study was approved by the Ethics Committee of the Faculty of Tropical Medicine, Mahidol University, Bangkok, Thailand.

### **Study drug administration**

An open randomized clinical trial of Artekin<sup>®</sup> vs Artesunate-Mefloquine was conducted at the Bangkok Hospital for Tropical Diseases, Mahidol University. Upon admission to the ward, patients were randomly treated at ratio of 1:2 into groups A:B as follows:

Group A: AM: Artesunate (4mg/kg/day) was given by oral route once a day for 3 days together with Mefloquine 8mg/kg/day for 3 days

Group B: Artekin: Artekin<sup>®</sup> (2 mg/kg/day of dihydroartemisinin and 15 mg/kg/day piperazine) was given by oral route once a day for 3 days (Artekin<sup>®</sup> 1 tab contains DHA 40mg + Piperazine 320mg)

All patients were treated symptomatically as indicated (e.g. intravenous fluid and antipyretics.) according to the standard practice in the hospital. In cases of RI, RII, or RIII responses (WHO, 1973), other antimalarial drugs (e.g. quinine plus tetracycline for falciparum malaria and chloroquine followed by primaquine for vivax malaria) were used as indicated. Patients who vomited within one hour after drug administration were redosed.

### **Monitoring for safety**

Patients were physically examined and adverse reactions during the study were recorded with the date and time at which they occurred and disappeared. Adverse effects were assessed on the basis of non-suggestive questioning by the study investigators. These include gastrointestinal, central nervous, cardiovascular, dermatological effects, as well as other changes possibly attributable to the study drugs. Routine blood investigations (hematology and biochemistry), and urinalysis were performed prior (Day 0) and weekly for 4 weeks of the study period.

## Statistical analysis

Statistical analysis was performed by using the Analyze It Add Ins Excel for Windows. All the *P-values* reported are from 2-tailed test and the statistically significance level was set at 0.05. The distribution of data was assessed for normality using the Schapiro-Wilks test. Data were expressed, as means and SD. Two statistical tests were performed. We used chi-square analyses to test differences between 2 groups of the qualitative variables and independent t-test to test the difference between 2 groups of the quantitative variables on demographics and baseline laboratory data. (Tabachnick *et al*, 2001)

## RESULTS

A total of one hundred and eighty patients were enrolled in this trial. All pregnancy tests in female patients were negative. Around 90% of patients completed the study as planned. Demographic clinical data and pretreatment laboratory characteristics are shown in Table 1. There were 135 male and 45 female patients aged 14 to 65 years old participated in this trial. There were no significant differences in the distribution of demographic, clinical and laboratory data between the two treatment groups.

At enrolment, patient in both treatment groups showed common malaria symptoms such as headache, asthenia, fatigue, fever, nausea, vomiting, myalgia and anorexia. Most clinical manifestations present on admission gradually disappeared during the first few days of treatment and coincided with high fever. Some baseline laboratory parameters were affected by disease status. However, they all returned to normal within 1-2 weeks.

Nineteen (6 and 13 patients in each group) patients did not complete the 28-days follow up due to social reasons not related to adverse effects. Thus, 161 patients out of 180 patients (89.5%) completed the 28-day study. No patients were deteriorated in clinical or biochemical changes after treatment in both groups. Parasitologic and clinical responses are shown in Table II. All patients in this study showed a prompted response to both antimalarial regimens (Figure 1). The cure rates at 28 days of follow-up were 100% and 99% respectively for the both treatment groups. There were no significant differences in fever clearance time and parasite clearance time between both treatment groups. No patients had RII or RIII failures. Only one patient in Artekin<sup>®</sup> treatment group had recrudescence on day 21 of study

period. The patient who drug failed to clear parasitemia was given the rescue antimalarial chemotherapy according to the hospital's standard regimen. Therefore, all patients had parasitologically negative at the time discharged from our hospital.

Means time for parasite clearances in each treatment group was fast, however there were no statistically significant different differences between the two treatment-groups [ $39.6 \pm 13.7$  hours and  $35.0 \pm 16.2$  hours in group 1 and 2 respectively, ( $p=0.72$ )]. The parasites were all cleared from peripheral blood smears within 84 hours. Similarly, there was no statistically significant difference between the fever clearance times of both treatment groups ( $p=0.67$ ).

No death occurred. No patients had vomiting related to the drugs. There were no major adverse effects and no neurologic or neuropsychiatric manifestations during treatment and during the 28-day follow-up period. Some minor symptoms such as nausea, headache, and dizziness occurred in group A (4, 3, 2 patients) and in group B (5, 4, 4 patients) respectively. However, these signs and symptoms could not be differentiated from malaria symptoms as they were disappeared between 1-4 days after treatment and while fever subsided. In addition, there was no serious adverse event reported during the study.

## DISCUSSION

In Thailand, *plasmodium falciparum* is resistant to chloroquine and there has been a decline in sensitivity to mefloquine (Brockman *et al*, 2000). The use of the artemisinin derivatives has been central to successful malaria control efforts in Thailand, Vietnam and Cambodian (Denis *et al*, 2002; Hien *et al*, 2004; Looareesuwan *et al*, 1997). Artemisinin derivatives are potent, rapidly acting antimalarials that can reduce parasitemias by more than 90% within 24 hours in uncomplicated malaria cases. However, the rate of recrudescence within 28 days when used alone can be as high as 10-25 % depending upon dosage, duration of treatment, and severity of disease (Hien *et al*, 1991; Li *et al*, 1994). These drugs are often combined with other long acting antimalarials such as mefloquine, (in this study combined with piperazine, to improve efficacy and compliance). The rationale using of the combination is as standard treatment of multidrug treatment for tuberculosis, patients with HIV and most cases of cancers. The rapid killings of parasitemias of artemisinin derivatives is accelerate the therapeutic response, prevent dangerous early treatment failures in case of high grade resistance,

reduce the parasite biomass and reduce gametocyte transmission (Looareesuwan *et al*, 1999).

The benefit of adding appropriate and suitable long action drug is prevent recrudescence by killing residual parasites, reduce the chance of a resistant mutant surviving and in addition the long acting antimalarial might protect the artemisinin derivative in low transmission areas. Combined administration of artemisinin derivatives and mefloquine in different dosages and duration had been studied in uncomplicated malaria in many countries. This combination is now a standard treatment for multidrug resistant falciparum malaria in Thailand (Wilairatana *et al*, 2002). However, some disadvantages of using artesunate-mefloquine might be seen (e.g. some patients could not tolerate mefloquine).

The combination of DHA and Piperaquine in the form of Artekin<sup>®</sup> was developed as an alternative to established combinations, such as artesunate-mefloquine, primarily to reduce treatment costs and toxicity. Our hospital based study has shown that a combination of dihydroartemisinin and piperaquine, is an effective and well tolerated by Thai adults with acute uncomplicated *plasmodium falciparum* malaria. Like a previous study (Denis *et al*, 2002; Hien *et al*, 2004) most of the patients whose treated by Artekin<sup>®</sup> in this study improved clinically and were parasite negative on the blood smear by the third day of treatment. The present study, all patients responded satisfactorily to the both treatment regimens. As well as, the present study shows a high total cure rate (99-100%) in the both groups. Artekin<sup>®</sup> showed similar cure rate to the standard treatment (artesunate-mefloquine). However at present, it has remained unclear whether the improve cure rate due to synergistic effect of the synergy dihydroartemisinin and piperaquine. There was no fatal patient in the study. Comparing with 3-day combination of artesunate-mefloquine treatment, Artekin<sup>®</sup> is given only in 3 days with 3 doses. The shorter period of Artekin<sup>®</sup> is now on clinical trial and might be better and have high chance of complete treatment course and improved compliance. This combination may serve as alternative regimens for treatment of uncomplicated falciparum malaria. The Artekin<sup>®</sup> has more advantage on these issues and more importantly since the drug was produced as a fix combination, less duration of treatment and possibly lower cost than artesunate-mefloquine. Other long acting drugs combined with artemisinin derivatives are under invention.

In conclusion, the results of this study indicate that Artekin<sup>®</sup> is effective and well-tolerated. Artekin<sup>®</sup> may be an alternative treatment to the standard combination of artesunate- mefloquine in treatment of multidrug resistant uncomplicated falciparum malaria

such as in Thailand. However, additional and more studies in special groups (in children, pregnant women, and field trials) and pharmacokinetic studies to guide rational dosing regimens are needed in order to get more informations of Artekin® in general practice.

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**Table I** Clinical and laboratory characteristics of study groups before treatment.

	Group A	Group B
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	(n = 60)	(n = 120)
Male/Female	49/11	86/34
Age (yr)		
Mean (SD)	26.5 (10.6)	24.3 (8.5)
Range	14-65	14-58
Mean (SD) height in cm	161.5 (9.3)	160.5 (8.1)
Mean (SD) weight in kg	55.6 (10.9)	51.8 (9.7)
Fever[Mean(SD)]		
Duration before admission (days)	5.6 (5.3)	5.3 (4.4)
Highest fever before treatment (°C)	38.2 (0.9)	38.3 (1.0)
No. of patients with:		
Splenomegaly	3	5
Hepatomegaly	12	19
Urine positive for drugs *	0	0
First malaria attack	29	32
Geometric mean parasites count (per µl)	4,645	3,759
Range high	102,500	190,860
Low	13	17
Laboratory data (mean [SD])		
Packed cell volume (%)	35.8 (5.1)	36.0 (6.0)
WBC count (per µl)	6,239 (4,593)	5,579 (1,776)
Blood urea (mmol/L)	14.2 (7.0)	15.3 (7.2)
Serum creatinine (umol/L)	0.8 (0.2)	0.9 (0.2)
Total bilirubin (umol/L)	1.4 (0.9)	1.5 (1.1)
Serum AST	38.1 (22.6)	42.7 (52.7)
Serum AAT	39.8 (34.4)	38.6 (31.8)
Albumin (mg/L)	3.7 (0.5)	3.6 (0.5)
Alk PO <sub>4</sub>	134.9 (56.4)	140.5 (100.4)

WBC = white blood count

AST, AAT = aspartate and alamine aminotransferases (U/L)

Alk PO<sub>4</sub> = alkaline phosphatase (U/L)

\*Sulphonamides and 4-aminoquinolones

**Table II** Therapeutic responses

	<b>Group A</b> (n = 60)	<b>Group B</b> (n = 120)
No. of patients dropout	6	13
No. of patients with 28 days follow up	54	107
No. (%) cured at 28 days	54 (100%)	106 (99%)
Recrudescence on days	-	21
Fever clearance time (hours)		
Mean (SD)	25.2 (28.4)	24.8 (24.7)
Range	4-100	4-124
Parasite clearance time (hours)		
Mean (SD)	39.6 (13.7)	35.0 (16.2)
Range	11-84	4-74

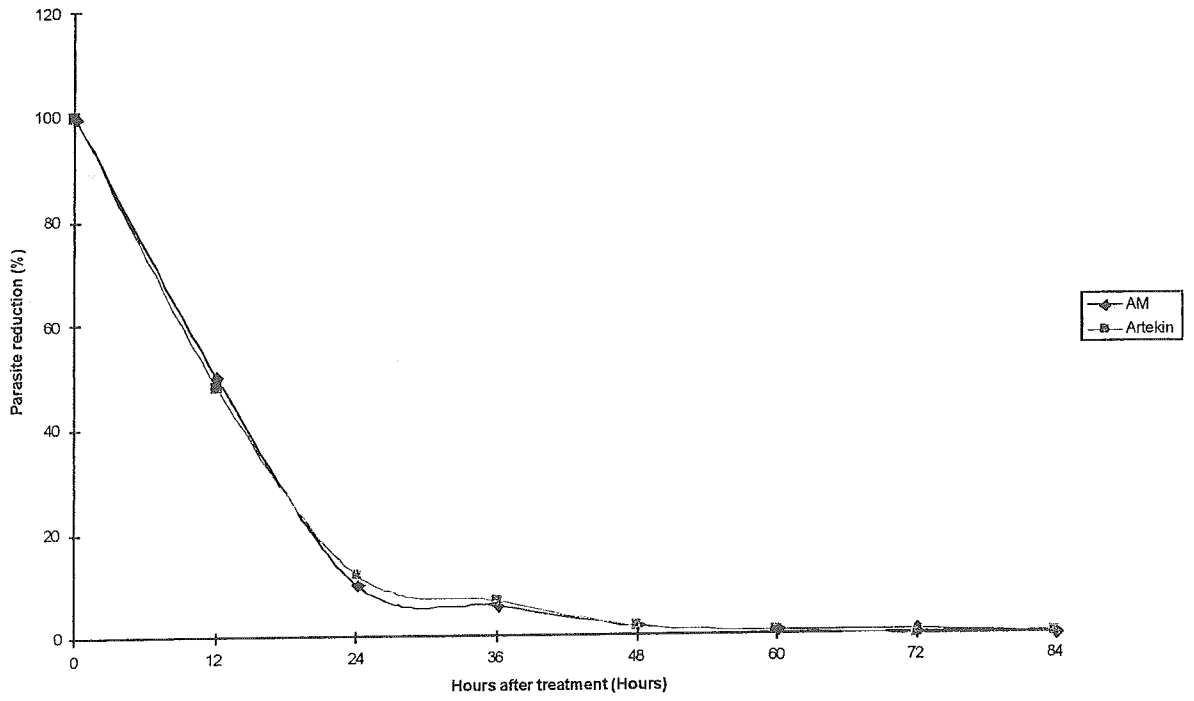


Figure 1 Percentage of malaria parasite reduction after treatment



# *Plasmodium falciparum*: The fungal metabolite gliotoxin inhibits proteasome proteolytic activity and exerts a plasmodicidal effect on *P. falciparum*<sup>☆</sup>

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## Abstract

The in vitro antimalarial activity of the fungal metabolite gliotoxin (GTX) was evaluated, and its mechanism of action was studied. GTX showed plasmodicidal activity against both *Plasmodium falciparum* chloroquine-resistant strain K-1 and chloroquine-susceptible strain FCR-3. GTX cytotoxicity was significantly lower against a normal liver cell line (Chang Liver cells). The intracellular reduced glutathione level of parasitized and of normal red blood cells was not affected by GTX treatment. However, GTX decreased the chymotrypsin-like activity of parasite proteasomes in a time-dependent manner. The results of this study indicate that GTX possesses plasmodicidal activity and that this effect is due to inhibition of parasite proteasome activity, suggesting that GTX may constitute a useful antimalarial therapy.

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**Index Descriptors and Abbreviations:** *Plasmodium falciparum*; Gliotoxin; Proteasome inhibitor; pRBCs, parasitized red blood cells; RBCs, red blood cells; GTX, gliotoxin; GSH, reduced glutathione

## 1. Introduction

Since the report of chloroquine (CQ)-resistant *Plasmodium falciparum* in the 1950s, the spread and persistence of drug-resistant malaria has continued to pose problems for malaria therapy. Multidrug resistant *falciparum* malaria parasites have been reported frequently in the Thai-Myanmar border area (Wongsrichanalai et al., 2002). Considering the increasing prevalence of drug-resistant *falciparum* malaria, the development of new antimalarial drugs is an urgent issue.

Gliotoxin (GTX) is a member of the epipolythiodioxo-piperazine class of fungal metabolites and possesses a diverse range of biologic activities including antimicrobial, antifungal, and antiviral activities, but is lethal to rodents at relatively low concentrations (LD<sub>50</sub> in mice and rats: 25–50 mg/kg) (Taylor, 1971; Waring and Beaver, 1996). GTX has also been reported to induce apoptotic cell death in numerous cell types. An important property of GTX is its ability to go through a redox cycle in the presence of an appropriate reducing agent. Hydrogen peroxide produced by GTX during the redox cycle in a cell-free system directly damages plasmid and cellular DNA (Eichner et al., 1986). GTX thus acts as a redox-active protein and has been shown to form mixed disulfides with accessible thiol residues on proteins or to induce further oxidative modification

<sup>☆</sup> Plasmodicidal activity of fungal metabolite gliotoxin.

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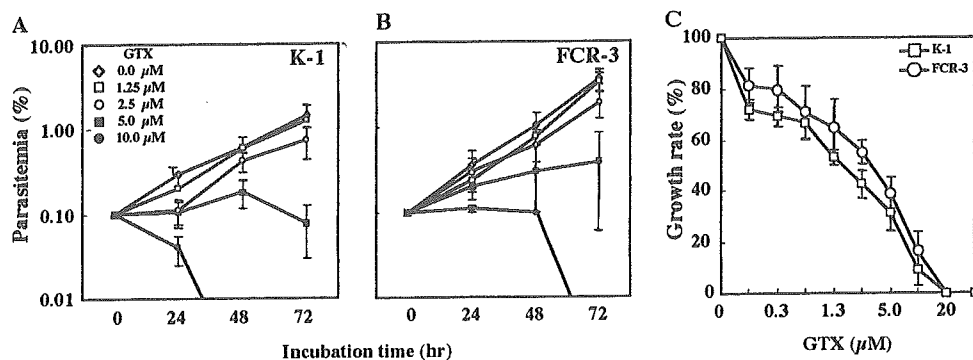


Fig. 1. Plasmodicidal effect of gliotoxin (GTX). GTX was added to cultures of chloroquine (CQ)-resistant strain K-1 (A) or CQ-susceptible strain FCR-3 (B) at a concentration of 10  $\mu\text{M}$  (filled circles), 5  $\mu\text{M}$  (filled squares), 2.5  $\mu\text{M}$  (open circles), 1.25  $\mu\text{M}$  (open squares), or 0.0  $\mu\text{M}$  (open diamonds). Parasitemia (%) was determined every 24 h. (C) Concentration-dependent effect of GTX on inhibition of K-1 (squares) and FCR-3 (circles) maturation in vitro. Results are presented as means  $\pm$  SEM. Parasitemia of the K-1 strain treated with 10  $\mu\text{M}$  GTX at 48 and 72 h are both 0%, and that of the FCR-3 strain treated with 10  $\mu\text{M}$  GTX at 72 h is also 0%.

48 (Bernardo et al., 2003). Recently, it was reported that GTX  
49 efficiently inhibits proteasome catalytic activity (Kroll et al.,  
50 1999).

51 In eukaryotic cells, most proteins in the cytoplasm and  
52 nucleus are degraded not within lysosomes but within pro-  
53 teasomes. The roles of proteasomes in protein turnover are  
54 involved in specific cellular functions, including pro-  
55 grammed inactivation of mitotic cyclins, transcription fac-  
56 tors, and transcriptional regulators; elimination of mutated  
57 or damaged proteins; and antigen presentation. These func-  
58 tions are tightly regulated (Zwickl and Baumeister, 2002).

59 Lactacystin, a proteasome inhibitor, has been reported  
60 to inhibit the growth of *P. falciparum* and to be effective  
61 against CQ-resistant parasites (Certad et al., 1999). Intra-  
62 cellular amastigote-trypomastigote transformation is also  
63 reported to be prevented by lactacystin treatment (de Diego  
64 et al., 2001). GTX has been reported to inhibit the growth  
65 of *Toxoplasma gondii* by inhibiting proteasome activity  
66 (Paugam et al., 2002). In this study, the plasmodicidal effect  
67 of GTX was evaluated in relation to effects on reactive oxy-  
68 gen species and intracellular proteasome activity.

## 2. Materials and methods 69

### 2.1. Parasite and cell culture 70

71 CQ-resistant *P. falciparum* strain K-1 and CQ-susceptible  
72 strain FCR-3 were grown asynchronously, according to  
73 the modified method of Trager and Jensen (1976), in RPMI  
74 1640 (pH 7.4) medium supplemented with 10% human A  
75 serum, 25 mM HEPES, 25  $\mu\text{g}/\text{ml}$  gentamicin (Sigma-Aldrich,  
76 St. Louis, MO), 24 mM sodium bicarbonate, and human O  
77 type red blood cells (RBCs) in disposable sterile culture  
78 dishes under a controlled atmosphere of 5%  $\text{CO}_2$ , 5%  $\text{O}_2$ ,  
79 and 90%  $\text{N}_2$  at 37  $^\circ\text{C}$ .

80 The human normal liver cell line, Chang liver cells, was  
81 kindly provided by Dr. Takeaki Nagamine (Gunma Uni-  
82 versity School of Health Sciences, Gunma, Japan). Cells  
83 were grown continuously in complete Dulbecco's modified  
84 Eagle's medium (DMEM; Sigma-Aldrich) supplemented  
85 with 10% fetal bovine serum (FBS) and penicillin (100 U/  
86 ml)-streptomycin (10 ml/L; Gibco-BRL, Grand Island,  
87 NY) under 5%  $\text{CO}_2$  at 37  $^\circ\text{C}$ .

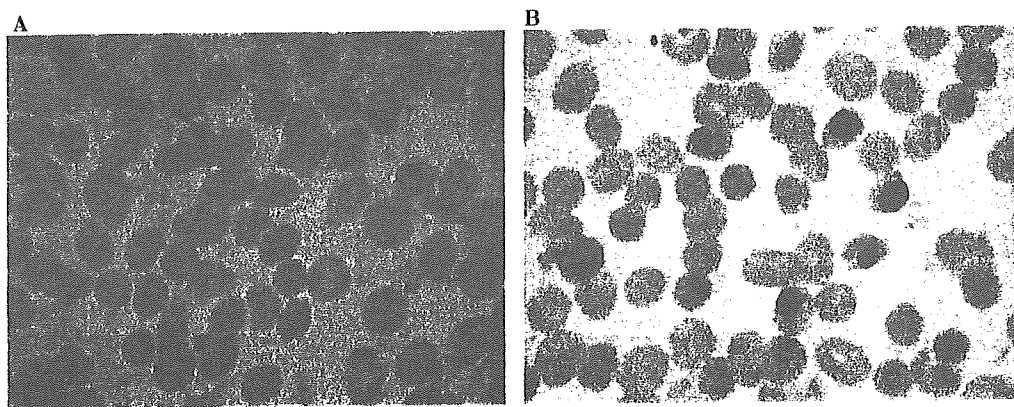


Fig. 2. Morphology of the Giemsa-stained K-1 strain after 24 h of incubation with GTX. The parasites were first synchronized at the ring stage and then cultivated with 10  $\mu\text{M}$  of GTX (A) and drug-free condition (B) for 24 h. Note the degenerative changes of the parasites in the GTX-treated culture (arrows in A). Magnification, 1000 $\times$ .

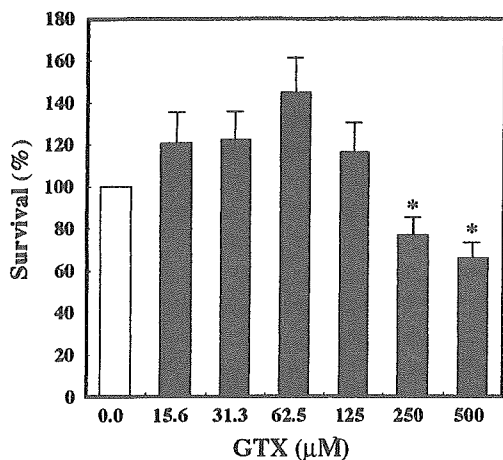


Fig. 3. Cell mortality after treatment with GTX for 72 h. The cytotoxic effect of GTX on Chang liver cells was evaluated by LDH release colorimetric assay (CytoTox-ONE). Results are presented as means  $\pm$  SEM. \*  $P < 0.05$ .

88 2.2. Evaluation of in vitro plasmodicidal effect of GTX

89 RPMI 1640 medium was supplemented with GTX in  
 90 solution at final concentrations ranging from 0.156 to  
 91 10.0  $\mu$ M. GTX-supplemented medium was changed every  
 92 24 h. Five hundred microliters of parasitized-RBC (pRBC)  
 93 suspension was placed into each well of a 24-well culture  
 94 plate (Corning Incorporated, Corning, NY) at a hematocrit  
 95 of 5% and initial parasitemia of 0.1%. Thin-smear speci-  
 96 mens were made every 24 h and stained with Giemsa solu-  
 97 tion, and parasitemia was evaluated by counting the  
 98 number of pRBCs in 3000 RBCs. Morphologic changes of  
 99 the parasites in response to drug treatment were monitored

by observing Giemsa-stained thin-smear specimens under  
 light microscopy.

2.3. Determination of GTX  $IC_{50}$

pRBCs were synchronized with 5% D-sorbitol for 15 min  
 at room temperature as previously described (Taguchi et al.,  
 2004). After three washes in RPMI 1640 medium, pRBCs  
 were resuspended in RPMI 1640 medium (pH 7.4) supple-  
 mented with 10% human O serum, Hepes (25 mM), gentami-  
 cin (25  $\mu$ g/ml), and sodium bicarbonate. Synchronous  
 pRBCs showing approximately 2% parasitemia were seeded  
 onto 24-well cell culture plates as described above. Twenty  
 microliters of GTX solution was added to each well at final  
 concentrations ranging from 0.156 to 40.0  $\mu$ M. Six wells per  
 plate served as GTX-untreated controls to monitor parasite  
 growth. After 24 h of incubation, the control wells were  
 checked for parasite growth. When parasites in control wells  
 became schizonts, the culture plates were removed from the  
 incubator, and thin-smear specimens were made from the  
 contents of each well and stained with Giemsa solution. We  
 counted RBCs in control smears for a total of 50 schizonts.  
 The effect of GTX on parasite growth was evaluated by  
 comparing the number of schizonts in the same number of  
 RBCs as was counted in the control cultures. Growth inhibi-  
 tion (%) was calculated as follows: (test well schizont count/  
 control well schizont count)  $\times$  100. The GTX concentration  
 that inhibited parasite growth by 50% ( $IC_{50}$ ) was calculated  
 by the probit method (Inaba et al., 2001).

2.4. Toxicity of GTX in Chang liver cells

To assess GTX cytotoxicity in a mammalian cell line,  
 Chang liver cells were analyzed with a lactate dehydrogenase

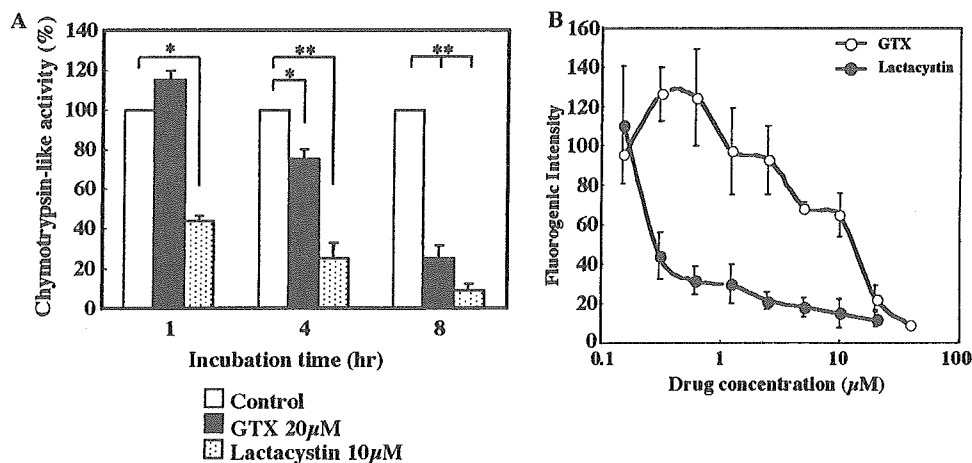


Fig. 4. GTX inhibits chymotrypsin-like activity of *P. falciparum* proteasomes in time- and concentration-dependent manner. (A) Chymotrypsin-like activity was measured on crude parasite extracts after addition of GTX together with the fluorogenic peptide, succinyl-Leu-Leu-Var-Tyr-4-methylcoumarinyl-7 amide. Activities are presented relative to GTX-untreated control (open columns, 100% activity). Data are presented as means  $\pm$  SEM of three independent experiments. Filled and hatched columns indicate GTX-treated (20  $\mu$ M) and lactacystin-treated (10  $\mu$ M) parasites as positive controls. \*  $P < 0.05$ , \*\*  $P < 0.01$ . (B) Proteasomes isolated from the parasites, plus lactacystin (open circle) and GTX (closed circle) at concentrations, were incubated with fluorogenic substrate succinyl-Leu-Leu-Var-Tyr-4-methylcoumarinyl-7 amide to measure chymotrypsin-like activity. Each point is the mean of the fluorescences of triplicate wells  $\pm$  SEM.



130 release colorimetric assay (CytoTox-ONE 96 Assay Kit;  
131 Promega, Madison, WI) as described previously (Hatabu  
132 et al., 2005). In brief, Chang liver cells were seeded at  $1 \times 10^5$   
133 cells/ml in 96-well culture plates (Corning Costar) and incu-  
134 bated at 37°C for 72h in DMEM supplemented with GTX.  
135 After incubation, cells were washed three times with Hanks  
136 solution (Sigma–Aldrich), and 100  $\mu$ l of reaction reagent was  
137 added to each well and incubated at 37°C for 10min. After  
138 the reaction was terminated, fluorescence levels were deter-  
139 mined with a multi-well plate reader (CytoFluor; Perseptive  
140 Biosystems, Framingham, MA) at an excitation wavelength  
141 of 530 nm and an emission wavelength of 580 nm.

#### 142 2.5. Measurement of reduced glutathione levels in pRBCs 143 and normal RBCs

144 Intracellular reduced glutathione (GSH) levels were  
145 measured according to the method of Hissin and Hilf  
146 (Hissin and Hilf, 1976). In brief, cell suspensions were sup-  
147 plemented with various concentrations of GTX (2.5–  
148 20  $\mu$ M). After a 24-h incubation, cells were counted with a  
149 hemocytometer. After a wash in PBS, homogenization  
150 buffer (4:1 mixture of 0.1 M sodium phosphate, 5 mM  
151 EDTA buffer, pH 8.0, and 25% metaphosphoric acid) was  
152 added. Cell suspensions were homogenized on ice with a  
153 sonic wave homogenizer for 40 s. Homogenates were cen-  
154 trifuged (21,000g, 30 min, 4°C), and supernatants were  
155 collected for analysis. After addition of *o*-phthalaldehyde  
156 (0.1% in methanol), GSH levels were determined with a  
157 multi-well plate reader (CytoFluor) at an excitation wave-  
158 length of 360 nm and an emission wavelength of 420 nm.  
159 Standards contained 0–2  $\mu$ g GSH/ml.

#### 160 2.6. Enzymatic assay of *P. falciparum* proteasome activity

161 Chymotrypsin-like activity of *P. falciparum* proteasomes  
162 was assessed by the fluorometric method of Gonzalez et al.  
163 (1999). In brief, GTX-treated and control pRBCs contain-  
164 ing trophozoite/schizont-stage parasites were harvested and  
165 incubated in 0.05% saponin (Sigma–Aldrich) on ice for  
166 30 min. After a PBS wash, homogenization buffer was  
167 added as described above. Cell suspensions were homoge-  
168 nized on ice with a sonic wave homogenizer for 40 s.  
169 Homogenates were centrifuged (21,000g, 30 min, 4°C), and  
170 supernatants were collected for analysis.

171 Chymotrypsin-like activity of 20S proteasomes was  
172 measured with the use of the fluorogenic substrate *N*-succinyl-  
173 Leu-Leu-Val-Tyr-7-amino-4-methyl-coumarin (Suc-  
174 LLVY -amc; Peptide Institute, Osaka, Japan). Lactacystin  
175 was purchased from the same source. A concentration of  
176 10  $\mu$ M of lactacystin was used as a positive control. Mean  
177 values of three experiments are shown.

#### 178 2.7. Statistical analysis

179 Data presented are the means  $\pm$  SEM values of at least  
180 three independent experiments. Student's *t* test was used for

181 statistical analysis.  $P < 0.05$  was considered statistically sig-  
182 nificant.

### 183 3. Results

#### 184 3.1. Plasmodicidal activity of GTX in CQ-resistant and 185 CQ-susceptible *P. falciparum*

186 To confirm the plasmodicidal activity of GTX, CQ-resis-  
187 tant *P. falciparum* strain K-1, and CQ-susceptible strain  
188 FCR-3 were included in medium containing GTX for 3 days,  
189 and parasite growth was monitored by determining the level  
190 of parasitemia (Figs. 1A and B). Growth of both strains was  
191 inhibited in a concentration-dependent manner by GTX, the  
192 parasites being degenerative (Fig. 2). In the presence of 5  $\mu$ M  
193 GTX, the growth of both strains was inhibited but not com-  
194 pletely. In the presence of 10  $\mu$ M GTX, the growth of both  
195 parasite strains was completely inhibited. However, CQ-resis-  
196 tant strain K-1 looked a bit more susceptible to GTX than  
197 CQ-susceptible strain FCR-3 by the graph.

198 Dose–response curves for GTX in K-1 and FCR-3 are  
199 shown in Fig. 1C. IC<sub>50</sub> values for GTX were  $3.60 \pm 0.97 \mu$ M  
200 for K-1 and  $4.17 \pm 1.08 \mu$ M for FCR-3. K-1 seemed to be  
201 a little more susceptible to GTX, but there was no  
202 significant difference between IC<sub>50</sub> values of CQ-resistant  
203 and CQ-susceptible parasites ( $P > 0.05$ ).

#### 204 3.2. GTX toxicity in Chang liver cells

205 Toxicity of GTX in Chang liver cells was also evaluated  
206 (Fig. 3). Although the survival of Chang liver cells was  
207  $\approx 70\%$  in response to  $\geq 250 \mu$ M GTX, survival was not  
208 affected by  $\leq 125 \mu$ M GTX.

#### 209 3.3. GTX effect on intracellular proteasome activity

210 To determine whether GTX inhibits proteasome-associ-  
211 ated proteolytic activity, we tested the effect of 20  $\mu$ M GTX  
212 on chymotrypsin-like activity (Fig. 4A). GTX decreased chy-  
213 motrypsin-mediated proteolysis of Suc-LLVY-amc by  $\approx 25\%$   
214 after 4 h treatment compared to untreated controls ( $P < 0.05$ ).  
215 After 8 h GTX treatment, 74% inhibition of Suc-LLVY-amc  
216 activity was observed compared to untreated controls  
217 ( $P < 0.01$ ). When tested on proteasomes isolated from para-  
218 sites, GTX showed inhibitory activity that paralleled its plas-  
219 modicidal activity (Fig. 4B). Thus, GTX decreased  
220 chymotrypsin-like activity of parasite proteasomes in a time-  
221 and concentration-dependent manner.

222 To study another plasmodicidal mechanism of GTX,  
223 intracellular GSH content was determined. Intracellular  
224 GSH levels of both pRBCs and RBCs were not affected by  
225 treatment with GTX (data not shown).

### 226 4. Discussion

227 The results of this study indicate that GTX possesses  
228 plasmodicidal activity and is effective against *P. falciparum*

229 CQ-resistant strain K-1 and CQ-susceptible strain FCR-3.  
 230 GTX has a multitude of effects, many of which are due to  
 231 proteasome inhibition. The results also indicate that GTX  
 232 inhibits chymotrypsin-like activity of parasite proteasomes.  
 233 Many parasites showed degenerative changes after GTX  
 234 treatment, as determined by light microscopy. Gonzalez  
 235 et al. (1999) reported inhibition of normal development to  
 236 the schizont stage by treatment with another proteasome  
 237 inhibitor, lactacystin. Morphologic changes in their report  
 238 were similar to those observed in this study.

239 GTX acts as a redox-active protein, and in cell-free  
 240 experiments, it has been shown to form mixed disulfides  
 241 with accessible thiol residues on proteins or to induce fur-  
 242 ther oxidative modification (Bernardo et al., 2003). Kweon  
 243 et al. (2003) reported that GTX induces apoptosis in  
 244 hepatic stellate cells by forming reactive oxygen species and  
 245 that the formation of reactive oxygen species by GTX is  
 246 concentration-dependent. However, our data indicate that  
 247 intracellular GSH levels are not altered, and there was no  
 248 difference in GSH levels between pRBCs and RBCs in  
 249 response to 10  $\mu$ M GTX, suggesting that the plasmodicidal  
 250 activity of GTX is not induced by oxidative damage. These  
 251 findings suggest that GTX primarily targets parasite pro-  
 252 teasomal chymotrypsin-like proteolytic activity.

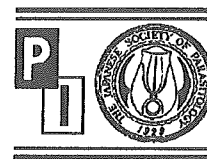
253 Proteasome inhibitors, including lactacystin, are candi-  
 254 dates for antimalarial therapy (Gantt et al., 1998). Taylor  
 255 (1971) reported that GTX was lethal to rodents at relatively  
 256 low concentrations ( $LD_{50}$  in mice and rats: 25–50 mg/kg).  
 257 However, the present study showed that GTX exerted  
 258 approximately 25-fold lower cytotoxicity on normal liver  
 259 cells than on parasites, indicating that GTX may be a  
 260 promising treatment for drug-resistant malaria.

261 In conclusion, GTX showed an inhibitory effect on plas-  
 262 modial development in vitro by inhibiting parasite protea-  
 263 some activity, and selective inhibitors of parasite  
 264 proteasomes may be useful antimalarial agents.

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## Down-regulation of tight junction mRNAs in human endothelial cells co-cultured with *Plasmodium falciparum*-infected erythrocytes

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### Abstract

To understand the mechanism of sequestration in the microvasculature of patients with falciparum malaria, we examined the patterns of expression of mRNAs for adhesion molecules (ICAM-1, VCAM-1, and E-selectin) and tight junction molecules (occludin, vinculin, and ZO-1) in human umbilical vein endothelial cells (HUVECs) co-cultured with *Plasmodium falciparum*-parasitized red blood cells (PRBCs) in vitro. The PRBCs were collected from patients with uncomplicated, severe, or cerebral malaria (CM). Patterns of mRNA expression in HUVECs co-cultured with PRBCs were examined by real-time quantitative reverse transcription-polymerase chain reaction (RT-PCR). Levels of mRNAs for all the three adhesion molecules increased with increased culture time within 3 h, regardless of the source of the PRBCs. In contrast, the patterns of mRNA expression for the tight junction molecules varied between the different co-cultures. When HUVECs were cultured with PRBCs from uncomplicated malaria patients, levels of mRNAs for tight junction molecules increased according to the culture time. HUVECs co-cultured with PRBCs from severe malaria patients showed no change in the mRNAs levels during 3 h of observation. When HUVECs were cultured with PRBCs from CM patients, levels of mRNAs for tight junction proteins decreased according to the culture time. Although the mechanisms underlying these phenomena are not clear, our results suggest that PRBCs can alter expression of tight junction proteins in endothelial cells at the site of sequestration and thereby influence disease severity.

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**Keywords:** *Plasmodium falciparum*; HUVEC; PRBC; Adhesion molecule; Tight junction molecule

### 1. Introduction

Approximately 40% of the world's population lives in malaria endemic areas, which are distributed across 100 countries in tropical and subtropical regions. Malaria infection results in 300–500 million clinical cases and 1.5–2.7 million deaths annually, with approximately 1 million deaths occurring in children less than 5 years of age. Approximately 90% of the

cases and deaths occur in tropical Africa. *Plasmodium falciparum* is responsible for the majority of severe clinical cases of malaria, which occur primarily in young children, non-immune adults, and pregnant women [1].

The most serious complication of malaria is cerebral malaria (CM). It is a diffuse reversible encephalopathy characterized by an altered level of consciousness, focal neurologic findings, and seizures. Sequestration of *P. falciparum*-parasitized red blood cells (PRBCs) to the microvasculature, predominantly that in the brain, heart, lungs, and submucosa of the small intestine, may be directly related to the pathogenesis of the disease. In vitro studies have shown that sequestration is

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Table 1  
Patient characteristics at time of admission per disease severity and isolate

Patient's severity	Uncomplicated malaria		Severe malaria		Cerebral malaria	
	MP 058	AA 863	NAC 42	NAC 44	AQ 1133	AQ 1142
Isolate no.						
Age (years)	38	20	18	21	50	45
Sex	M	F	F	F	F	M
RBC ( $\times 10^6/\mu\text{l}$ )	5.55	4.48	3.09	3.9	3.55	4.87
Hb (g/dl)	13.1	11.9	7.2	8.2	10.4	15.5
Hematocrit (%)	41	35	22	26	31	45
WBC ( $\times 10^3/\mu\text{l}$ )	3.5	3.8	12.6	7.5	14.3	7.1
Plt ( $\times 10^4/\mu\text{l}$ )	15.2	25	28	38	11	40
BUN (mg/dl)	7	8.2	171	17.1	65	55.1
Cr (mg/dl)	0.8	0.77	4.71	0.65	1.9	2.4
Albumin (g/dl)	4.3	3.9	2.7	3.1	2.9	2.6
Total bilirubin (mg/dl)	0.7	0.6	2.59	7.1	7.1	11.76
AST (U/L)	14	53	284	73	175	156
ALT (U/L)	13	59	106	66	208	102
Parasite count (/ $\mu\text{l}$ )	44,400	98,560	954,810	561,600	188,150	735,370
Schizont presented	No	No	Yes	Yes	Yes	Yes
Acute renal failure	No	No	Yes	No	No	No
Jaundice	No	No	Yes	Yes	Yes	Yes
Blood transfusion (no. of unit transfusion)	No	No	Yes (2)	Yes (2)	No	No
Respiratory distress with endotracheal tube insertion	No	No	Yes	No	Yes	Yes
Glasgow coma score	15	15	13	15	8	7
Fever clearance time (h)	48	56	144	24	96	116
Parasite clearance time (h)	40	51	73	50	88	164

RBC, red blood cell; Hb, hemoglobin; WBC, white blood cell; Plt, platelet; BUN, blood urea nitrogen; Cr, creatinine; AST, aspartate aminotransferase; ALT, alanine aminotransferase.

mediated by the interaction of a parasite protein, *P. falciparum* membrane protein-1 (PfEMP-1), which is located on knob-like protrusions on the surface of PRBCs, with receptors on the host endothelium [2]. Many host adhesion molecules, including intercellular adhesion molecule-1 (ICAM-1 or CD54), thrombospondin (TSP) [3,4], CD36 [5], chondroitin sulfate A (CSA) [6,7], vascular cell adhesion molecule 1 (VCAM-1), and endothelial leukocyte adhesion molecule 1 (ELAM-1) or E-selectin [8], bind PRBCs.

However, there is evidence that endothelial junctions are disrupted in patients with CM [9]. Immunohistochemistry of post-mortem tissue from Vietnamese adults and Malawian

children with CM showed loss of endothelial-cell junctional proteins ZO-1, occludin, and vinculin, most notably in vessels containing sequestered PRBCs [9–11]. Therefore, the pathogenesis caused by sequestration in CM may be related to expression of adhesion molecules and tight junction molecules in endothelial cells.

To examine whether *P. falciparum* isolates from patients can alter expression of mRNAs for adhesion molecules (ICAM-1, VCAM-1, and E-selectin) and tight junction molecules (occludin, vinculin, and ZO-1) in endothelial cells, we co-cultured human umbilical vein endothelial cells (HUVECs) with PRBCs from patients with malaria of different severities.

Table 2  
Sequences of oligonucleotide primers and probes used for TaqMan RT-PCR

Gene	Amplicon length (bp)	Oligonucleotide sequence (5'-3')	GenBank accession no.	
ICAM-1	90	Forward	CATAGAGACCCCGTTGCCTAAA	NM_000201
		Reverse	TGGCTATCTTCTTGCACATTGC	
		Probe	TACACCTTCCGGTTGTTCCAGGCA	
VCAM-1	124	Forward	ATGACCTTCATCCCTACCATTGA	NM_001078
		Reverse	CATTGACATAAAGTGTTTGCCTACTCT	
		Probe	TGATGACATGGAATTCGAACCCAAACAA	
E-selectin	122	Forward	CCACTGGGAAACTTCAGCTACAA	NM_000450
		Reverse	GCTGGAATAGGAGCACTCCATT	
		Probe	CCAAGCAGCATGGAGACCATGCA	
Occludin	77	Forward	CCCATCTGACTATGTGGAAAGA	NM_002538
		Reverse	AAAACCGCTTGTCACTTCTTTG	
		Probe	TTGACAGTCCCATGGCATACTCTCCAATG	
Vinculin	136	Forward	GCATCGTCTGGCTAATGTTATGAT	NM_014000
		Reverse	TGCTCGTGCCTGAGGACTCT	
		Probe	CTGTCACTGGTCCACTCGGTCCACA	
ZO-1	144	Forward	CACCTTTTGATAATCAGCACTCTCA	NM_003257
		Reverse	CTCTAGGTGCCCTGTTCTGTAACGT	
		Probe	TTGAAGAGCCAGCCCTCTGTCTTACG	