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# SHORT REPORT: HEPATITIS B INFECTION AND SEVERE *PLASMODIUM FALCIPARUM* MALARIA IN VIETNAMESE ADULTS

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*Abstract.* We investigated the prevalence of infection with hepatitis B virus among adult Vietnamese patients hospitalized for severe *Plasmodium falciparum* malaria. Sera from patients admitted with severe malaria in Ho Chi Minh City, Vietnam, between May 1991 and January 1996 were assayed for hepatitis B surface antigen ($HB_sAg$) by a commercial enzyme-linked immunosorbent assay kit. The overall prevalence of $HB_sAg$ was 23.77% (77 of 324). This was higher than reported estimates of prevalence in the general catchment population for the study hospital (mean, 9.8%; range, 9–16%). No association was found between risk of death caused by severe malaria and $HB_sAg$. Patients admitted with cerebral malaria had a slightly greater risk of registering positive for $HB_sAg$ (relative risk, 1.28; 95% confidence interval, 1.04–1.58) relative to other manifestations of severe malaria. Chronic infection with hepatitis B virus may be a risk factor for severe malaria.

Despite remarkable achievements in reducing malaria mortality, malaria remains the most common cause of morbidity and mortality in Vietnam. Of the 71.6 million population, 41.9 million are at risk for malaria; 15 million live in malaria-endemic areas.[1] The majority of people with malaria are not admitted to the hospital. Instead, they receive oral therapy on an outpatient basis. People who develop severe malaria and enter the hospital represent a minority of patients.

This report examines the association of infection with hepatitis B and severe malaria and the impact of that factor on survival. Hepatitis B, a double-stranded DNA virus of the hepadnoviridae family, infects 15% of people in Southeast Asia.[2] Globally, > 2 billion people are infected; 350 million of which are asymptomatic carriers of the virus.[3] The infection kills > 1 million carriers annually, with mortality generally associated with complications of cirrhosis, hepatocellular carcinoma, and, rarely, fulminant liver failure during acute infection.[4,5]

Coendemic falciparum malaria and acute hepatitis B occur through much of Southeast Asia, Africa, and the tropical Americas. Both diseases represent key threats to public health. To our knowledge, large studies evaluating the effect of acute hepatitis B infection on risk of severe disease and death caused by falciparum malaria have not been reported. We hypothesized that acute hepatitis B may exacerbate the risk of sequelae with infection by *Plasmodium falciparum.* By assaying for hepatitis B surface antigen, a marker for active infection, among patients admitted to hospital in Vietnam and enrolled into a treatment trial for severe malaria, we were able to test this hypothesis.[6]

Between May 1991 and January 1996, a randomized, double-blind controlled trial was conducted by Hien and others[6] to compare artemether to quinine for the treatment of severe malaria in Vietnamese adults. The study took place in a research ward at the Center for Tropical Diseases in Ho Chi Minh City, Vietnam. Severe malaria was defined as having one or more of the following: cerebral malaria (Glasgow Coma Scale < 11), anemia (hematocrit < 20%) with a parasite count > 100,000/μL, jaundice (serum bilirubin > 2.5 mg/dL) with a parasite count > 100,000/μL, renal impairment (urine output < 400 mL/24 hr and serum creatinine > 3 mg/dL), hypoglycemia (blood glucose < 40 mg/dL), hyperparasitemia (> 10%), or shock (systolic blood pressure < 80 mm Hg with cool extremities). This definition of severe malaria includes 2 criteria considered supporting by the World Health Organization (WHO), hyperparasitemia and jaundice, and does not include 5 criteria included by WHO: respiratory distress, repeated convulsions, spontaneous bleeding, acidosis, and hemoglobinuria.

Inclusion criteria included the following: age > 14 years, not in the first trimester of pregnancy, no intravenous drug abuse, and receipt of < 3 g quinine or 2 doses of any artemisinin derivatives in the previous 48 hr. Eligible subjects were enrolled after informed consent was obtained from the patients or their guardians.[6] A total of 560 patients were enrolled and randomly assigned to receive either artemether or quinine dihydrochloride intramuscularly, according to the regimens described by Hien and others.[6] Artemether proved as safe and effective as quinine.

Assessable sera samples were obtained from 326 of these subjects. Samples from the remaining subjects were exhausted in a battery of other tests. Samples were collected by disposable syringe, centrifuged, serum aspirated, and maintained for the first 3 months at −20°C. Samples were then transferred to a −70°C freezer and stored at the Institute for Clinical Research in Tropical Medicine, Hanoi, Vietnam. The U.S. Naval Medical Research Unit 2 in Jakarta, Indonesia, analyzed the samples with the commercially available Auszyme monoclonal diagnostic kit for the detection of hepatitis B surface antigen in serum or plasma (Abbott Laboratories, Singapore) according to the manufacturer's instructions.

Results were expressed as means or frequencies of positive and negative results and were analyzed by the unpaired Student's *t*-test or by the Mantel-Haenszel test. The chi-square test was used to assess differences in proportions. $P \leq 0.05$ was considered statistically significant. Logistic regression was used to evaluate the relationship between several potential confounders and death from severe malaria. Analysis was carried out by SPSS version 8.0 (SPSS, Chicago, IL) and Epi Info version 6 (Centers for Disease Control and Prevention, Atlanta, GA).

The mean age of severe malaria subjects was 33 years (standard deviation, 13), with a range of 15–78 years. Se-

TABLE 1
Characteristics of patients of admission according to hepatitis B status

| Characteristic | HB$_s$Ag positive | HB$_s$Ag negative | P value |
|---|---|---|---|
| Age (years) | | | |
|   Median | 29 | 30 | 0.55 |
|   Range | 15–71 | 15–78 | |
| Sex (% male) | 57/77 (74.0) | 182/247 (73.7) | 0.95 |
| Pregnant (%) | 3/77 (3.9) | 11/247 (4.6) | 0.83 |
| Intravenous drug use (%) | 0/76 (0.0) | 2/236 (0.8) | 0.42 |
| Treatment regimen (% receiving quinine) | 45/77 (58.4) | 121/247 (49.0) | 0.15 |
| Cerebral malaria* (%) | 49/77 (63.6) | 122/246 (49.6) | 0.03 |
| Shock† (%) | 12/77 (15.6) | 28/247 (11.3) | 0.32 |
| Jaundice‡ (%) | 31/73 (42.5) | 126/240 (52.5) | 0.13 |
| Renal failure§ (%) | 17/77 (22.1) | 76/243 (31.3) | 0.12 |
| Severe anemia¶ (%) | 2/77 (2.6) | 18/245 (7.3) | 0.13 |
| Fever# (%) | 67/77 (87.0) | 194/247 (78.5) | 0.10 |
| Transfusion (%) | 13/77 (16.9) | 63/246 (25.6) | 0.12 |
| Pulmonary syndrome** (%) | 26/77 (33.8) | 96/247 (33.1) | 0.42 |
| Convulsions | 8/77 (10.4) | 23/247 (9.3) | 0.78 |

* Glasgow Coma Scale <11.
† Systolic blood pressure <80 mm Hg with cool extremities.
‡ Serum bilirubin >3.0 mg/dL.
§ Serum creatinine >3.0 mg/dL.
¶ Hematocrit <15%.
# Temperature ≥37.5°C.
** Any one of the following: acute respiratory failure, pulmonary distress, or pulmonary edema.
HB$_s$Ag = hepatitis B surface antigen.

roprevalence of hepatitis B surface antigen (HB$_s$Ag) was 77/324 (23.8%). The death rate among patients with severe malaria was 47/323 (14.6%). Table 1 lists the demographic and clinical characteristics of subjects, comparing these between those positive and negative for HB$_s$Ag. The only statistically significant difference between HB$_s$Ag-positive and -negative subjects was in frequency of cerebral malaria: 64 versus 50% (P = 0.03). The relative risk of cerebral malaria with HB$_s$Ag was 1.28 (95% confidence interval [CI], 1.04–1.58, P = 0.03). The frequency of jaundice, renal failure, severe anemia, and the necessity of transfusion were all lower among HB$_s$Ag-positive subjects, but these differences were not statistically significant.

Table 2 lists the death rate stratified by HB$_s$Ag status. HB$_s$Ag positivity carried no elevated risk of death compared with patients without evidence of hepatitis B infection. Multivariate logistic regression analysis controlling for age, sex, pregnancy, and treatment regimen revealed no association between a higher risk of death and HB$_s$Ag positivity. Age was modeled as a categorical variable with the following groups: < 25 years, 26–35 years, 36–50 years, and > 50 years. The adjusted odds ratio (95% CI) was 0.95 (0.44–2.00), similar to the crude odds ratio of 0.99 (0.45–2.16). The sample of analyzable sera in this study would have detected a ≥ 13% difference in incidence of death with 80% power. The analysis thus reveals that HB$_s$Ag in serum at the time of admission for severe malaria was not a significant risk factor for death as an outcome for the treatment regimens applied.

HB$_s$Ag, the outer coat protein of the virus, is a marker for acute infection or a chronic carrier state. HB$_s$Ag is detectable for 2–15 weeks after exposure and then disappears in ~90% of people infected during convalescence. In the remaining 10%, HB$_s$Ag does not clear. These people become chronic carriers. The patients positive for HB$_s$Ag in this study were not suffering acute hepatitis, but were presumably nearly all chronic carriers, although we cannot exclude with certainty that some patients may have had a resolving acute infection. The cross-sectional prevalence of HB$_s$Ag (e.g., at admission to hospital with severe malaria) includes those recovering from acute infection and the cumulative proportion of chronic carriers in the population.

The prevalence of HB$_s$Ag in subjects with severe malaria was 23.77% (77/324), higher than in cross-sectional estimates from the general adult population of the region. The study by Kakumu and others represents an exception to this as the population surveyed included children (mean age, 28 years; range, 2–81 years).[2] Table 3 summarizes measurements of the prevalence of HB$_s$Ag in populations from which the severe malaria patients were drawn and compares these to HB$_s$Ag prevalence in those patients. In all instances, the prevalence among subjects with severe malaria was significantly higher (P < 0.001). Measurements of the prevalence of HB$_s$Ag in the general population ranged 9–16%.[7,8] The lower range, 3.1–5.7%, occurred among residents of rural areas of Ho Chi Minh City. We considered 9.8% HB$_s$Ag positivity reported from the large study of Tran and others[8] a surrogate control for the severe malaria cases we evaluated.

TABLE 2
Assessment of outcome

| Variable | HB$_s$Ag positive (%) | HB$_s$Ag negative (%) | Relative risk (95% confidence interval) | P value |
|---|---|---|---|---|
| Prevalence of death | 11/76 (14.5) | 36/247 (14.6) | 0.99 (0.53–1.85) | 0.98 |

Relative risk is for the HB$_s$Ag-positive group. Groups were compared via the Mantel-Haenszel test. HB$_s$Ag = hepatitis B surface antigen.

Table 3
History of HB$_s$Ag prevalence studies in Ho Chi Minh City, Vietnam

| Date of study | Population of Ho Chi Minh City | Prevalence of HB$_s$Ag (%) | Chi-Square test statistic* (P value) |
|---|---|---|---|
| 1996 | Rural area[2] | 5.7 (51/890) | 81.91 (<0.001) |
| 1993 | Subpopulations[7] | | |
| | Overall | 10.5 (64/610) | 29.09 (<0.001) |
| | Range | 8–14% | |
| 1992 | Blood donors[9] | 3.1 (15/491) | 83.61 (<0.001) |
| 1989–1991 | Subpopulations[8] | | |
| | Overall in normal population† | 9.8 (3,837/39,080) | 69.87 (<0.001) |
| | Range in normal population† | 9–16% | |

* Comparing the prevalence in patients with severe malaria with each population. HB$_s$Ag = hepatitis B surface antigen.
† Excluding intravenous drug abusers, patients with hepatitis, and patients with liver cancer.

Excluding intravenous drug abusers, hepatitis patients, and liver cancer patients yielded a population of 39,080 subjects, closely approximating the population from which our sample was drawn. Using that estimate of normal risk of HB$_s$Ag positivity, the odds ratio for the likelihood of HB$_s$Ag positivity in patients with severe malaria versus the normal population was 2.9 (95% CI, 2.19–3.73, $P < 0.0001$). Obviously, a prospective case control study would have provided a more accurate estimate of this ratio. Finally, malaria tends to be a rural disease in Vietnam, where prevalence of hepatitis B is relatively low. If the severe malaria patients at Ho Chi Minh Hospital came predominately from rural areas with relatively low rates of hepatitis B, the relative risk of concurrent events would be considerably higher. Therefore, we considered the 2.9 odds ratio of HB$_s$Ag in severe malaria patients versus the normal population to be a conservative indirect estimate of elevated risk of severe malaria with hepatitis B infection. This finding warrants studies that directly measure the risk of severe malaria with hepatitis B disease.

Our analysis, which was based on samples collected in the setting of a therapeutic trial, suggests that acute hepatitis B infection may exacerbate falciparum malaria, causing an elevated risk of admission to the hospital for severe malaria. The analysis lacked carefully selected controls for the severe malaria cases and applied a relatively crude classification of hepatitis B disease. In addition, it is possible that some sampling bias existed because sera samples were not available from all patients. Last, because this study included only adult subjects, the conclusions cannot be automatically extrapolated to children. The findings nonetheless point to the possibility of an interaction between these diseases that may increase the risk of morbidity and mortality for the many millions of people exposed to endemic risk of both infections. The mechanism of this apparent increase in susceptibility is not known. Perhaps chronic hepatitis B carriers are less efficient at limiting parasite multiplication. There was no suggestion that hepatitis B carriers were older or more likely to come from malaria endemic areas, which might imply a failure to develop specific protective immunity. Understanding this relationship may help refine strategies intended to diminish disease and death caused by malaria.

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