

Usefulness of the recombinant liver stage antigen-3 for an early serodiagnosis of *Plasmodium falciparum* infection

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Abstract: In order to develop tools for an early serodiagnosis of *Plasmodium falciparum* infection, we evaluated the usefulness of *P. falciparum* liver stage antigen-3 (LSA-3) as a serodiagnostic antigen. A portion of LSA-3 gene was cloned, and its recombinant protein (rLSA-3) was expressed in *Escherichia coli* and purified by column chromatography. The purified rLSA-3 and 120 test blood/serum samples collected from inhabitants in malaria-endemic areas of Mandalay, Myanmar were used for this study. In microscopic examinations of blood samples, *P. falciparum* positive rate was 39.1% (47/120) in thin smear trials, and 33.3% (40/120) in thick smear trials. Although the positive rate associated with the rLSA-3 (30.8%) was lower than that of the blood stage antigens (70.8%), rLSA-3 based enzyme-linked immunosorbent assay could detect 12 seropositive cases (10.0%), in which blood stage antigens were not detected. These results indicate that the LSA-3 is a useful antigen for an early serodiagnosis of *P. falciparum* infection.

Key words: *Plasmodium falciparum*, falciparum malaria, liver stage antigen-3, recombinant protein, serodiagnosis

INTRODUCTION

Malaria, caused by the apicomplexan protozoa *Plasmodium* sp., is one of the most important infectious diseases worldwide, and the problem appears to be worsening (Breman et al., 2001; Snow et al., 2001).

Although the morbidity and mortality rates have been declining gradually, malaria continues to constitute the major cause of mortality and morbidity in many parts of the world. Approximately 300 to 500 million new malaria cases and 2 to 5 million deaths are estimated to occur annually. The emergence and rapid spread of resistant strains to commonly used anti-malarial drugs, including chloroquine and antifolates, poses a serious challenge to the malaria control (Wellems and Plowe, 2001; Roper et al., 2003; Baird, 2004; Gregson and Plowe, 2005).

Vivax malaria reemerged in 1993 and continues to

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be a big health threat to Korea (Chai et al., 1994; Cho et al., 1994; Yeom et al., 2005). Recent increases of overseas travel and economic activities with other countries, especially those having malaria-endemic regions, make another concern with falciparum malaria. In deed, 476 imported malaria cases have been reported from 1994 to 2004 in the Republic of Korea (unpublished data, Korea Center for Disease Control and Prevention). Most of these patients were infected with *Plasmodium falciparum* and had been infected principally in Asia and Africa where drug resistance strains are prevalent. Therefore, an early diagnosis, with prompt treatment, is highly important. In this respect, development of protocols and materials for an earlier diagnosis of falciparum malaria is urgently necessitated.

The sporozoites of malaria parasites transmitted from the saliva of infected mosquitoes travel rapidly to the liver and invade hepatocytes, where they develop into the exoerythrocytic stage called a tissue schizont. During this stage, the parasites express liver stage specific antigens. In *P. falciparum*, at least 2 of the relevant antigens, liver stage antigen-1 (LSA-1) and liver stage antigen-3 (LSA-3), have been identified and characterized (Guerin-Marchand et al., 1987; Aidoo et al., 2000; Joshi et al., 2000). These proteins are both surface proteins, and are expressed solely in infected hepatocytes and thought to have a role in liver schizogony and merozoite release. Specific humoral, cellular, and cytokine immune responses to LSA-1 and LSA-3 are well documented, with epitopes identified that correlate with the antibody production, proliferative T-cell responses, or cytokine induction (Aidoo et al., 2000; Joshi et al., 2000; Perlaza et al., 2001). Both preerythrocyte antigens have been considered to be vaccine candidates against *P. falciparum* due to their antigenic and protection-inducing immunogenic properties (Daubersies et al., 2000; Kurtis et al., 2001; Sauzet et al., 2001; Taylor-Robinson, 2003). In the present study, we evaluated the usefulness of *P. falciparum* LSA-3 as a serodiagnostic antigen for a more rapid diagnosis of falciparum malaria than using the currently available diagnostic tools.

MATERIALS AND METHODS

Blood samples

Blood samples were collected from 120 inhabitants who visited the malaria clinic with signs and symptoms suggestive of malaria in endemic areas of Mandalay, Myanmar. Thin and thick blood smears were prepared for routine microscopic examinations. Informed consent was obtained from all patients. The study protocol was approved by the Department of Health (Upper Myanmar), the Union of Myanmar.

Cloning and sequence analysis of liver stage antigen-3 (LSA-3) of *P. falciparum*

To amplify a portion of *P. falciparum* LSA-3 gene, we designed specific primer sets based on the DNA sequences provided by the Genebank database. LSA-3F contained a *Bam*HI site at its 5' end (5'-GATC-CGAAAATGTAGAAGAAAATGACGACGGAAGT-3'), and LSA-3R contained a *Hind*III site at its 5' end (5'-AAGCTTTGATAAATTGTTGCAACGTTTTTCAGCAAC-3'). Genomic DNA of *P. falciparum* was extracted from the whole blood of a patient using a QIAamp DNA Blood Kit (Qiagen, Valencia, California, U.S.A.) following the manufacturer's instruction. Polymerase chain reaction (PCR) was performed with AccuPower™ PCR Premix (Bioneer, Daejeon, Korea), 50 ng of purified genomic DNA, and 40 pmoles of the above-described reverse and forward primer sets, in a total volume adjusted to 30 μ l with distilled water. The reaction condition was as follows: denaturation at 94°C for 10 min, 35 cycles of 1 min at 94°C, 1 min at 60°C, and 1.5 min at 72°C, and a final extension for 10 min at 72°C. The PCR product was gel-purified using a Gel extraction kit (Qiagen), ligated into pCR2.1-TOPO™ cloning vector (Invitrogen, Carlsbad, California, U.S.A.) and then transformed into *E. coli* TOP10. Sequencing reactions were conducted with a BigDye Terminator Cycle Sequencing Ready Reaction Kit in an ABI 377 automatic DNA sequencer (Applied Biosystems, Foster City, California, U.S.A.). Nucleotide and deduced amino acid sequences were analyzed with the SeqEd.V1.0.3 program, and the CLUSTAL program provided in the Megalign soft-

ware, a multiple-alignment program of the DNASTAR package (DNASTAR, Madison, Wisconsin, U.S.A.).

Expression and purification of recombinant LSA-3 (rLSA-3)

For the expression of *P. falciparum* LSA-3 in *Escherichia coli*, the amplified PCR product was digested with *Bam*HI and *Hind*III and purified with a Gel extraction kit (Qiagen), ligated into pQE30 expression vector (Qiagen), and transformed into *E. coli* M15 (pREP4) cells (Qiagen). The selected clones were grown and induced with 1 mM isopropyl-1-thio- β -D-galactopyranoside (IPTG). The bacteria were then suspended in native lysis buffer (50 mM NaH₂PO₄, 300 mM NaCl, 10 mM imidazole, pH 8.0), sonicated on ice and centrifuged at 4°C for 20 min at 12,000 g. The supernatant was collected and analyzed by SDS-PAGE. The rLSA-3 was purified by nickel-nitrilotriacetic acid (Ni-NTA) chromatography (Qiagen).

Western blot analysis

The purified rLSA-3 was run on 10% SDS-PAGE gel, and the protein was transferred onto a nitrocellulose membrane. After the transfer, the membrane was cut into strips and blocked for nonspecific binding with 3% skim milk for 12 hr at 4°C. The strips were then washed 3 times with phosphate buffered saline (PBS) with 0.15% Tween 20 (PBST) for 10 min each. The strips were allowed to react with the serum samples obtained from either normal or *P. falciparum*-infected persons (diluted 1:100, v/v) for 4 hr, and then washed 3 times with PBST as described above. Diluted peroxidase-conjugated anti-human IgG (1:1,000, v/v; Sigma, St. Louis, Missouri, U.S.A.) was added to each strip followed by 3 hr incubation at room temperature. For color development, 0.2% diaminobenzidine and 0.002% H₂O₂/PBS were used.

Enzyme-linked immunosorbent assay (ELISA)

The 96-well polystyrene plates were coated with 1 μ g/ml of purified rLSA-3 diluted in PBS (pH 7.4) per well. The plates were incubated overnight at 4°C. After incubation, the plates were washed 3 times with

PBST and blocked via 2 hr incubation with 200 μ l of 3% BSA in PBST per well for 2 hr at 37°C. After 3 washes with PBST, 100 μ l of sera (1:100, v/v) was added to each well, and incubated for an additional 2 hr at 37°C. After washing as described above, the plates were incubated for 2 hr at 37°C, with 100 μ l of peroxidase-conjugated anti-human IgG (1:1,000, v/v, Sigma). The plates were then washed 3 times with PBST and 100 μ l of substrate solution was added. The substrate solution was prepared immediately prior to use by dissolving 30 mg of *o*-phenylenediamine (Sigma) per 60 ml of peroxidase solution B (H₂O₂ in 0.1 M phosphate citric acid buffer, pH 5.0). The plates were then incubated for an additional 30 min at dark, and the reactions were terminated by adding 50 μ l of 4 N H₂SO₄. Optical densities were measured at 450 nm with an ELISA reader.

Immunofluorescent antibody test (IFAT)

IFAT was conducted with smeared *P. falciparum*-infected red blood cells (blood stage antigens; BSA) on multispot slides. The infected red blood cells were routinely obtained by in vitro cultivation of the parasite in our laboratory. The parasite-infected red blood cells in slides were allowed to react with the test sera (1:32, v/v) for 30 min, then rinsed twice with PBS, and incubated for 30 min with fluorescein isothiocyanate (FITC)-conjugated anti human IgG (1:100, v/v, Sigma). The slides were then observed under a fluorescence microscope.

RESULTS

Cloning of LSA-3 from *P. falciparum* genomic DNA

The genomic DNA used for PCR amplification of LSA-3 was prepared from a Myanmar patient. Sequencing analysis of the cloned LSA-3 gene was determined to encompass 1,500 bp and encoded for 500 amino acids (Fig. 1). It exhibited a specific amino acid tandem repeats, 27 repeats of VEESVAEN, and 6 repeats of VEEIVAPTVEEI and VAPTVEEIVAPSV-VESVAPSVEESVEEN. These repeats are highly conserved in different strains of *P. falciparum* and the

1 E N U E E N D D G S U A S S U E 16
 1 GAA AAT GTA GAA GAA AAT GAC GAC CGA AGT GTA GCC TCA AGT GTT GAA 48
 17 E S I A S S U D E S I D S S I E 32
 49 GAA AGT ATA GCT TCA AGT GTT GAT GAA AGT ATA GAT TCA AGT ATT GAA 96
 33 E N U A P T U E E I U A P S U U 48
 97 GAA AAT GTA GCT CCA ACT GTT GAA GAA ATT GTA GCT CCA AGT GTT GTA 144
 49 E S U A P S U E E S U E E N U E 64
 145 GAA AGT GTC CCT CCA ACT GTT GAA GAA AGT GTA GAA GAA AAT GTT GAA 192
 65 E S U A E N U E E S U A E N U E 80
 193 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 240
 81 E S U A E N U E E S U A E N U E 96
 241 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 288
 97 E I U A P T U E E I U A P T U E 112
 289 GAA ATC GTA GCT CCA ACT GTT GAA GAA ATC GTA GCT CCA ACT GTT GAA 336
 113 E I U A P S U U E S U A P S U U 128
 337 GAA ATT GTA GCT CCA ACT GTT GTA GAA AGT GTG GCT CCA AGT GTT GAA 384
 129 E S U E E N U E E S U A E N U E 144
 385 GAA AGT GTA GAA GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 432
 145 E S U A E N U E E S U A E N U E 160
 433 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 480
 161 E S U A E N U E E S U A E N U E 176
 481 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 528
 177 E I U A P T U E E I U A P T U E 192
 529 GAA ATC GTA GCT CCA ACT GTT GAA GAA ATC GTA GCT CCA ACT GTT GAA 576
 193 E I U A P S U U E S U A P S U U 208
 577 GAA ATT GTA GCT CCA ACT GTT GTA GAA AGT GTG GCT CCA AGT GTT GAA 624
 209 E S U E E N U E E S U A E N U E 224
 625 GAA AGT GTA GAA GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 672
 225 E S U A E N U E E S U A E N U E 240
 673 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 720
 241 E S U A E N U E E S U A E N U E 256
 721 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 768
 257 E S U A E N U E E S U A E N U E 272
 769 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 816
 273 E I U A P T U E E I U A P T U E 288
 817 GAA ATC GTA GCT CCA ACT GTT GAA GAA AGT GTA GCT CCA ACT GTT GAA 864
 289 E S U A P S U U E S U A P S U U 304
 865 GAA ATT GTA GCT CCA ACT GTT GTA GAA AGT GTG GCT CCA ACT GTT GAA 912
 305 E S U E E N U E E S U A E N U E 320
 913 GAA AGT GTA GAA GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 960
 321 E S U A E N U E E S U A E N U E 336
 961 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 1008
 337 E S U A E N U E E I U A P T U U 352
 1009 GAA AGT GTA GCT GAA AAT GTT GAA GAA ATC GTA GCT CCA ACT GTT GAA 1056
 353 E I U A P T U U E E I U A P S U U 368
 1057 GAA ATC GTA GCT CCA ACT GTT GAA GAA ATT GTA GCT CCA ACT GTT GTA 1104
 369 E S U A P S U U E E S U E E N U E 384
 1105 GAA AGT GTA GCT CCA ACT GTT GAA GAA AGT GTA GAA GAA AAT GTT GAA 1152
 385 E S U A E N U U E S U A E N U U 400
 1153 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 1200
 401 E S U A E N U U E E I U A P T U U 416
 1201 GAA AGT GTA GCT GAA AAT GTT GAA GAA ATC GTA GCT CCA ACT GTT GAA 1248
 417 E I U A P T U U E E I U A P S U U 432
 1249 GAA ATC GTA GCT CCA ACT GTT GAA GAA ATT GTA GCT CCA AGT GTT GTA 1296
 433 E S U A P S U U E S U E E N U U 448
 1297 GAA AGT GTG GCT CCA ACT GTT GAA GAA AGT GTA GAA GAA AAT GTT GAA 1344
 449 E S U A E N U U E S U A E N U U 464
 1345 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 1392
 465 E S U A E N U U E S U A E N U U 480
 1393 GAA AGT GTA GCT GAA AAT GTT GAA GAA AGT GTA GCT GAA AAT GTT GAA 1440
 481 E I U A P T U U E E I U A E N U U 496
 1441 GAA ATC GTA GCT CCA ACT GTT GAA GAA ATC GTT GCT GAA AAC GTT CCA 1488
 497 T N L S 500
 1489 ACA AAT TTA TCA 1500

Fig. 1. Nucleotide and deduced amino acid sequences of the liver stage antigen-3 (LSA-3) of *Plasmodium falciparum* Myanmar isolates. Specific amino acid tandem repeats are observed in the sequence.

LSA-3 gene from the Myanmar isolate evidenced its highest degree of homology (99.6%) with a Kenya isolate (AJ007010) (data not shown).

Expression and purification of recombinant LSA-3 (rLSA-3)

The expression of the rLSA3 was induced by adding IPTG and analyzed on SDS-PAGE. The size of the expressed protein, rLSA3, was approximately 52 kDa, which is consistent with the predicted molecular weight of the protein (Fig. 2A). In the western blot analysis of the purified rLSA-3, 5 serum samples from 14 malaria patients infected with *P. falciparum* were shown to be reactive against rLSA-3 (Fig. 2B).

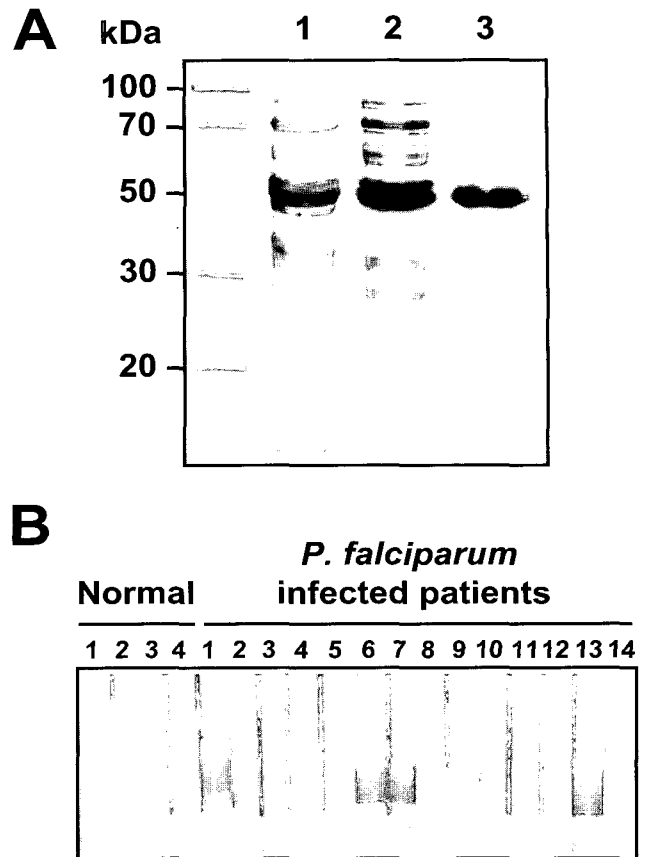


Fig. 2. Expression and purification of recombinant LSA-3. (A) SDS-PAGE analysis. Lane 1, uninduced *E. coli* lysate; lane 2, IPTG induced *E. coli* lysate; lane 3, Ni-NTA affinity purified rLSA-3. (B) Western blot analysis. Each strip was incubated with sera from *P. falciparum*-infected patients. Some of the sera (5 of 14 samples) exhibited positive reactions.

Comparison of parasite detection rates

In order to evaluate the utility of rLSA-3 as a serodiagnostic antigen, we conducted several diagnostic tests and the parasite detection rate of each test was compared. In the case of microscopic examinations of the 120 blood samples collected in malaria-endemic areas of Mandalay, Myanmar, the positive rate was 39.2% (47/120) in thin smear trials and 33.3% (40/120) in thick smear trials (Table 1). Cases of mixed infection with *P. vivax* and *P. falciparum* were also detected in thin blood smear tests. IFAT employing the whole BSA revealed that 85 (70.8%) among 120 sera tested were determined to have antibodies against BSA. Meanwhile, ELISA employing the rLSA-3, positive rate was lower than that for blood stage antigens, at

Table 1. Microscopic examinations of 120 blood samples using thick and thin smears

	No. of positive samples (%)	
	Thin smear	Thick smear
<i>Plasmodium vivax</i>	23 (19.2)	14 (11.7)
<i>Plasmodium falciparum</i>	19 (15.8)	26 (21.7)
<i>P. vivax</i> + <i>P. falciparum</i> ^{a)}	5 (4.2)	0 (0.0)
Negative	73 (60.8)	80 (66.6)
Total	120 (100.0)	120 (100.0)

^{a)}Coinfection with *P. vivax* and *P. falciparum*.

Table 2. Comparison of the antibody positive rates of 120 sera against liver stage antigen-3 (LSA-3) and blood stage antigen (BSA)

	No. of positive (%)
LSA-3 ^{a)}	12 (10.0)
LSA-3 + BSA ^{b)}	25 (20.8)
BSA ^{c)}	60 (50.0)
Negative	23 (19.2)
Total	120 (100.0)

^{a)}Positive only for LSA-3.

^{b)}Positive for both LSA-3 and BSA.

^{c)}Positive only for BSA.

30.8% (37/120). However, rLSA-3 based assays could detect 12 seropositive cases (10.0%), in which BSA were not detected (Table 2).

DISCUSSION

In the present study, we attempted to develop materials through which malaria parasites could be detected prior to the clinical onset of the disease. All human malaria including *P. falciparum*, remain for several weeks or months in liver cells after their introduction into the human body by mosquitoes, in the form of sporozoites. If the disease could be diagnosed in patients during these pre-onset periods of incubation, chances of early treatment might be greatly improved.

The infected sporozoites of malaria parasites developed and matured in the hepatocytes. During this incubation period, the parasites express liver stage-specific antigens, including LSA-1 and LSA-3. These

preerythrocyte antigens evoked specific humoral and cellular immune responses in the hosts and have been recognized as candidates for vaccine (Aidoo et al., 2000; Daubersies et al., 2000; Joshi et al., 2000; Kurtis et al., 2001; Perlaza et al., 2001; Sauzet et al., 2001; Taylor-Robinson, 2003). In this consideration, the liver stage-specific antigen could also be a good diagnostic candidate for early diagnosis of malaria.

In order to determine the feasibility of this, we evaluated the rLSA-3 as a promising candidate marker for screening of patients with liver-resident parasites. As we expected, the antibodies against LSA-3 were proved to be detectable in the patient's sera prior to the production of antibodies specific for BSA. Although positive rate of rLSA-3 based ELISA was lower than that of BSA, rLSA-3 based assay could detect 12 seropositive cases (10.0%), which were not detected by the BSA. Therefore, LSA-3 appears supplemental to the BSA, and may be proved useful as a serodiagnostic adjunct. A larger prospective study with a larger number of samples and different infectious status of samples are needed to generalize this method for a diagnostic use.

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