

Biochemical and Immunological Properties of a Viral Hybrid Particle Expressing the *Plasmodium vivax* Merozoite Surface Protein 1 C-terminal Region

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Abstract

Background: Mammalian cells expressing the small hepatitis B virus surface protein (HBs) secrete highly immunogenic 20 nm lipoprotein particles. Previous studies demonstrated that the fusion of foreign sequences into certain regions of HBs leads to chimeric particles carrying epitopes for the foreign peptide, as well as for HBs. The present study investigates immunologic and biochemical properties of the fusion of the C-terminal region of the merozoite surface 1 protein of *P. vivax*, the most widely distributed human malaria parasite, and HBs (PvMSP1₁₉-HBs).

Materials and Methods: COS7 cells were transfected with a plasmid coding for PvMSP1₁₉-HBs. The hybrid products were analyzed by density gradient centrifugation and electron microscopy or de-

tected by metabolic labeling and immunoprecipitation with anti-HBs and patient-derived anti-*P. vivax* serum. Mice were immunized with the vector and the antibody response was checked by ELISA.

Results: The fusion PvMSP1₁₉-HBs formed particles of 20–45 nm size, which were secreted from COS7 cells. The particles were immunoprecipitable with anti-HBs and serum of different *P. vivax*-positive individuals. Immunization of mice with the construct as a genetic vaccine showed that antibodies were raised mostly against the PvMSP1₁₉ domain and recognized the native protein.

Conclusion: Due to its biochemical and antigenic properties, the hybrid particle will be useful in future vaccine trials against the asexual blood stages of *P. vivax* as a genetic and/or a proteic subunit candidate.

Introduction

The C-terminal region of the merozoite surface protein 1 of *Plasmodium* (MSP1₁₉) is one of the most solid subunit vaccine candidates against the asexual blood stages of malaria, although its function remains elusive (1). Experiments in rodents, *Aotus* and *Saimiri* monkeys, either with the affinity-purified, yeast-, bacteria- or cell culture-derived MSP1₁₉, showed protection against blood stage challenge with *P. falciparum* (2), *P. vivax* (3), *P. cynomolgi* (4), and *P. yoelii* (5). Recently, a phase I clinical trial was reported (6). Since protection is mostly antibody-dependent (1),

the correct folding of MSP1₁₉, which contains two interspecies-conserved epidermal growth factor (EGF)-like domains stabilized by 10 cysteine residues, was shown to be of crucial importance. Accordingly, a protective monoclonal antibody binds to a reduction-sensitive epitope in the first EGF-like domain of the *P. yoelii* MSP1₁₉ (7). In monkey trials, the *E. coli*-produced MSP1₁₉-glutathion-S-adenosyl transferase (GST) fusion failed to confer protection (8), indicating that the natural conformation of *P. falciparum* MSP1₁₉ may not be achievable in the bacterial expression system. To date, only baculovirus-derived peptides are repeatedly protective in monkey trials (2,4). However, the baculovirus expression system cannot be easily scaled up to produce relevant quantities and yeast-expressed MSP1₁₉ are not as effective (3,9).

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Attempts to increase the immunogenicity of MSP1₁₉ included the fusion to promiscuous T-helper cell epitopes (3,9) or the use of novel adjuvants, nevertheless, none of them are licensed for human use (10,11).

Another way to increase the antigenicity of peptides is the construction of chimeric fusions with the small hepatitis B virus surface protein (HBs). The HBs moiety in this fusion provides, on one hand, widely recognized T-helper cell epitopes (12) and, on the other hand, leads to the formation of subviral particles with increased intrinsic immunogenicity, ideally carrying the fused epitope in high density on its surface. In earlier studies, chimeric fusions of several peptides with HBs were shown to increase the immunogenicity of the fused domains (13–15). Significantly, the most successful (pre-erythrocytic-stage) human malaria vaccine trial ever reported (16) contained large parts of the circumsporozoite protein fused to HBs, forming chimeric subviral particles purifiable from yeast cells (17).

In order to render MSP1₁₉ of the human malaria parasite *P. vivax* (PvMSP1₁₉) more antigenic as DNA, we previously reported on the construction of a fusion construct of the PvMSP1₁₉ gene from the Belem strain and the HBs gene (18). In this study, we report on biochemical and immunological properties of the hybrid particle composed of PvMSP1₁₉ and HBs.

Materials and Methods

Plasmid Constructs

The plasmid pSV33M-* coding for HBs [subtype adw2 (19)] is a derivative of pSV33H (20), in which a unique BamHI site was introduced in the downstream position of the first EcoRI site and immediately upstream from the HBs ATG initiation codon. The detailed construction of the recombinant plasmid, VXORF-PvMSP1₁₉-HBs, encoding the fusion PvMSP1₁₉-HBs protein was described elsewhere (18). The plasmid pGEX2T-PvMSP1₁₉ was created by insertion of the PvMSP1₁₉ fragment into pGEX2T (Pharmacia, Uppsala, Sweden) using the 5' and 3' primers (forward 5'-ccgatccATGAGCTCCGAGCACAC and reverse 5'-ccgaattcGAGGACAAGCT-TAGGAAG), respectively. The production of recombinant PvMSP1₁₉ as a fusion protein with GST was done as previously described (21). All insert sequences were con-

firmed by manual, standard dideoxy sequencing. Plasmids used for transfections were purified using the Maxiprep-Qiagen columns according to the manufacturer's instructions (Qiagen, Hilden, Germany). Plasmids were stored at -20°C in 70% ethanol until use and then diluted to 1 mg/ml in phosphate-buffered saline (PBS).

COS7 Cell Transfections

The plasmids pSV33M-* and VXORF-PvMSP1₁₉-HBs were transiently transfected in 50% confluent COS7 cells using the DEAE-Dextran method as described (20). Metabolic labeling with ³⁵S-Translabel (ICN, Costa Mesa, CA), immunoprecipitation and visualization of synthesized proteins essentially were done as described (20), except that immunoprecipitates were washed only once with PBS. Immunoprecipitations were either carried out using commercially available anti-HBs serum (Dako, Hamburg, Germany) or sera from human patients infected with *P. vivax*, which were positive for anti-MSP1 (and blocked before use with the supernatant of mock-transfected COS7 cells). The human sera were obtained from 4 male and 2 female adult patients in the acute phase of vivax malaria before treatment. All donors were infected in the Brazilian Amazon and reported between 2 and 20 previous malaria episodes.

Purification of Samples for Electron Microscopy

COS7 cells were transfected as above. Instead of labeling the cells, however, whole supernatants containing secreted proteins were centrifuged for 10 min at 4000 g and then subjected to CsCl-gradient (10–40% w/w) centrifugation in 10 ml-tubes in a Beckman Ti70 rotor (40 hr, 45,000 rpm, 16°C). Particle-containing fractions were coated to electron microscopy grids for 20 min and negatively stained with 2% Uranyl acetate for a further 20 min and stored under vacuum until analysis.

Immunization of Balb/C Mice

Groups of 10 female Balb/C mice (4–6 weeks old) were immunized intramuscularly in both tibialis anterior muscles with the plasmids VXORF-PvMSP1₁₉-HBs (6 mice) or VXORF (4 mice, control group). Three immunizations were given in three-week intervals with 100 µg (50 µg per leg) per dose per mouse. Pre-im-

mune sera and sera from all animals two weeks after the last boost were collected and stored at 20°C until use. Two independent trials were performed.

ELISA

Sera from all mice were analyzed by enzyme linked immuno-sorbant assay (ELISA) (21) to look for the presence of anti-HBs and/or anti-PvMSP1₁₉ antibodies. Briefly 0.2 µg/well of recombinant PvMSP1₁₉-GST or HBs (kindly supplied by N. Granovski, Instituto Butantan, São Paulo, Brazil) was coated to 96-well microtiter plates, washed, blocked with 4% skimmed milk in PBS, and incubated with antiserum in different serial dilutions in 1% milk/PBS. After further washings, bound antibodies were detected by incubation with anti-mouse immunoglobulin G (IgG)-peroxydase conjugate (Sigma, St. Louis, MO). After addition of tetramethylbenzidine/H₂O₂ substrate (Kierkegaard & Perry Laboratories, Gaithersburg, MD) and 15 min incubation, the colorimetric reaction was stopped with 1 M H₃PO₄ and the optical density (OD)₄₅₀ was determined.

Results

COS7 Cells Efficiently Secrete a Hybrid Structure Containing PvMSP1₁₉ and HBs Epitopes

In the first experiment, we determined whether a hybrid PvMSP1₁₉-HBs protein was efficiently translated and secreted by COS7 cells. As shown in Figure 1A, after transfection of COS7 cells with pSV33M^{-*} and metabolic labeling, the HBs proteins of 24 and 27 kD (glycosylated variant) could be immunoprecipitated by anti-HBs from cell lysates and culture supernatants (lanes 1 and 2). When COS7 cells were cotransfected with VXORF-PvMSP1₁₉-HBs and pSV33M^{-*}, both forms (~46 kD and 24/27 kD, respectively) could be observed in the supernatant of cells by immunoprecipitation with anti-HBs (lane 4). The secretion, however, of either form seemed slightly less efficient than that of cells transfected with a single plasmid. After transfection of cells with VXORF-PvMSP1₁₉-HBs, a protein in the supernatant in the expected size of ~46 kD was precipitated by anti-HBs (lane 6). Under these conditions, no signal was seen in mock-transfected cells (lanes 7 and 8).

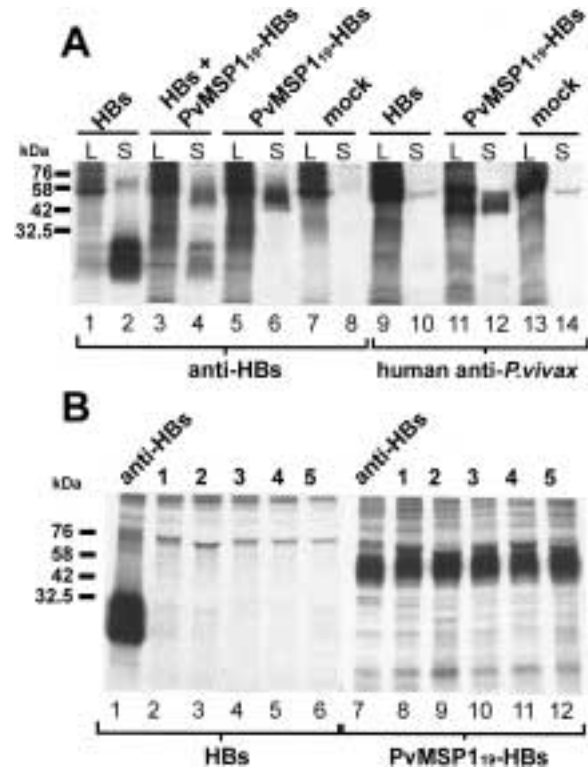


Fig. 1. Expression of HBs and PvMSP1₁₉-HBs in COS7 cells. (A) COS7 cells were transfected with different plasmids and lysates (L) and Supernatants (S) were immunoprecipitated with polyclonal anti-HBs (diluted 1:500) or a presaturated human serum positive for *P. vivax* anti-PvMSP1₁₉ (diluted 1:1000), but negative for anti-HBs. HBs, transfection with pSV33M^{-*}, PvMSP1₁₉-HBs + HBs, Cotransfection with pSV33M^{-*} and VXORF-PvMSP1₁₉-HBs, PvMSP1₁₉-HBs, transfection with VXORF-PvMSP1₁₉-HBs, mock: transfection with plasmid pVXORF1. (B) Anti-PvMSP1 contained in 5 different human sera recognized the PvMSP1₁₉-HBs protein produced in COS7 cells. COS7 cells were transfected with pSV33M^{-*} or VXORF-PvMSP1₁₉-HBs, pulse labeled and the supernatants immunoprecipitated with the sera from 5 different patients (dilution of each serum 1:1000). They were analyzed by SDS-PAGE. As a positive control, supernatants were immunoprecipitated with anti-HBs. HBs, small hepatitis B virus surface protein.

When the material produced by transfected and labeled cells was immunoprecipitated with the serum from a *P. vivax*-infected patient blocked previously with mock-transfected unlabeled COS7 culture supernatant, a band of 46 kD could be observed (Fig. 1A, lanes 11 and 12). This band was absent in material from cells transfected with pSV33M^{-*} or mock-transfected cells (lanes 9, 10, 13, 14). As the antiserum previously tested negative for anti-HBs in ELISA and failed to precipitate wildtype

HBs (lanes 9 and 10), precipitation of the hybrid structure was conferred by the PvMSP1₁₉ domain (lanes 11 and 12). This confirmed that the PvMSP1₁₉ portion of the chimeric protein was localized on the particle surface and accessible for precipitating antibodies specific for the natural PvMSP1₁₉ peptide.

Immune Sera Specifically Recognized PvMSP1₁₉ Epitopes Expressed on the Surface of the Secreted Hybrid Protein

We then asked if different sera of individuals infected with *P. vivax* were able to recognize the COS7 cell-expressed PvMSP1₁₉-HBs construct. Therefore, supernatants of VXORF-PvMSP1₁₉-HBs or pSV33M-* transfected and pulse-labeled COS7 cells were immunoprecipitated with sera from 5 different individual *P. vivax* patients containing anti-PvMSP1₁₉ antibodies, but no detectable anti-HBs antibodies. Significantly, all sera precipitated the PvMSP1₁₉-HBs construct, but not HBs, further confirming that the structure displayed PvMSP1₁₉ epitopes recognized in natural vivax infections (Fig. 1B, lanes 7–12).

Chimeric Structures Form Particles and Have Slightly Different Densities than Wildtype HBs

To test if the COS7-expressed PvMSP1₁₉-HBs hybrid protein formed particles morphologically, compared with wildtype HBs 20-nm particles, COS7 cells were transfected with pSV33M-* and VXORF-PvMSP1₁₉-HBs and metabolically labeled. The resulting supernatants of cells were subjected to CsCl gradient centrifugation. Fractions were dialyzed and immunoprecipitated with anti-HBs and the precipitate analyzed by SDS-PAGE. The peak fraction of particles containing wildtype HBs had a density of 1.17 g/ml; whereas, the PvMSP1₁₉-HBs hybrid structure peaked at 1.22 g/ml (data not shown). In a second experiment, secreted material from transfected unlabeled cells was subjected to CsCl gradient centrifugation as above and the material from the peak fractions (1.17 g/ml and 1.20–1.22 g/ml, respectively) was examined by electron microscopy. As shown in Fig. 2A, HBs particles of ~22 nm were found in the supernatant of cells transfected with a plasmid coding for wildtype HBs. Material from cells transfected with the hybrid construct VXORF-PvMSP1₁₉-HBs showed particles with differing diameters of

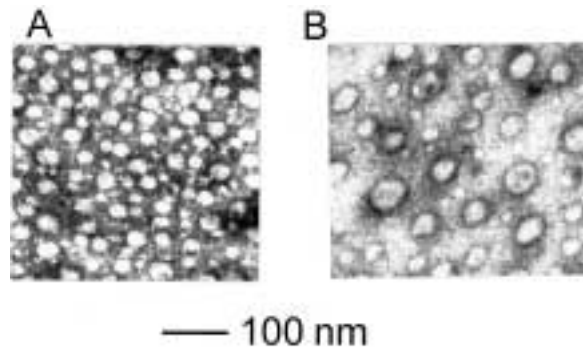


Fig. 2. PvMSP1₁₉-HBs forms particles. (A) Electron micrograph showing particles of the supernatant of pSV33M-transfected cells. (B) Micrograph showing hybrid particle structures of differing size (20–45 nm) derived from supernatant of cells transfected with pVXORF-PvMSP1₁₉-HBs. Magnification in A and B was 100,000 \times .

up to 45 nm (Fig. 2B). The hybrid particles differed in sizes ranging from 18 to 45 nm in diameter and this may be due to the voluminous PvMSP1₁₉ domain. The average size distribution of particles counted from an entire electron micrograph ($n = 163$) were 4% 40–45 nm, 30% 25–40 nm and 66% 18–22 nm. Variations in the morphology of HBs particles also occurred naturally (22). Regardless, these data indicated that, similar to wildtype HBs, PvMSP1₁₉-HBs proteins assembled into particle-like structures, which were secreted from expressing cells.

Immunization of Mice with the Recombinant Plasmid VXORF-PvMSP1₁₉-HBs Generated Predominantly Antibodies against PvMSP1₁₉

We recently demonstrated that immunization of mice with the recombinant plasmid VXORF-PvMSP1₁₉-HBs elicited antibodies capable of interacting with recombinant *E. coli* PvMSP1₁₉ (18). In order to determine whether these antibodies recognized PvMSP1₁₉ and HBs epitopes expressed on the viral chimeric particle, Balb/C mice were immunized with the plasmid VXORF-PvMSP1₁₉-HBs. After three immunizations, antiserum titers against recombinant PvMSP1₁₉ and HBs were determined using ELISA. As shown in Table 1, all mice developed considerable titers for anti-PvMSP1₁₉ antibodies, but only 2 of 6 or 4 of 6 mice contained low titers for anti-HBs antibodies. In a parallel experiment, COS7 cells were transfected with the plasmids pSV33M-* and VXORF-PvMSP1₁₉-HBs, pulse-labeled

Table 1. Antibody titers in ELISA after immunization of mice with plasmids VXORF1 or VXORF-PvMSP1₁₉-HBs

| Plasmid | anti-PvMSP1 ₁₉ | Responders* | anti-HBs | Responders** |
|---------------------------------|---------------------------|-------------|--------------|--------------|
| VXORF-PvMSP1 ₁₉ -HBs | 1:12800–1:51600 | 6/6 // 6/6 | 1:200–1:1000 | 2/6 // 4/6 |
| VXORF1 | <1:100 | 0/0 // 4/4 | <1:50 | 0/0 // 4/4 |

*, ** Responders are defined as $\geq 1:500^*$ or $\geq 1:200^{**}$ in endpoint dilution ELISA. HBs, small hepatitis B virus surface protein

with ³⁵S-methionine, and cell lysates and supernatants were immunoprecipitated with pooled sera from mice immunized with VXORF-PvMSP1₁₉-HBs or commercially available anti-HBs serum. Significantly, the genetic immunization of mice elicited antibodies capable of specifically and predominantly interacting with PvMSP1₁₉ epitopes (Fig. 3). Moreover, these results were very similar to those observed with sera from *P. vivax* patients (Fig 1B).

Discussion

In this study, we describe biochemical and immunological properties of the merozoite surface protein 1 C-terminal region of *P. vivax* (PvMSP1₁₉), presented as a hepatitis B virus subviral surface antigen hybrid particle. Transfection of COS7 cells with a recombinant plasmid encoding a fusion PvMSP1₁₉-HBs protein led to the production of subviral chimeric particles of ~18–45 nm diameter with a peak density of 1.22 g/ml. Moreover, the hybrid protein was precipitable with sera from naturally infected *P. vivax* patients and with sera from mice genetically immunized with this recombinant plasmid.

von Brunn et al. (15) previously described the generation of a recombinant HBs particle containing central parts of MSP1 of *P. falciparum* in the vaccinia virus expression system. Thus, a significant part of the HBs loop forming the immunodominant a-epitope of HBsAg was exchanged for sequences of MSP1 and the different constructs inserted into the vaccinia virus genome by homologous recombination. After infection of CV1 cells, subviral particles that were mostly identical in their biochemical behavior to wildtype HBs particles were observed. In a different report, Gordon and colleagues (17) inserted the C-terminal part of the *P. falciparum* circumsporozoite (CS) protein,

which contains several cysteines, into the N-terminal position of HBs. The authors were able to express the construct in *Saccharomyces cerevisiae* as a hybrid particle only if a certain quantity of wildtype HBs was coexpressed. Apparently, the 189 amino acid containing the CS domain caused sterical problems during the budding process of the nascent particle at the endoplasmic reticulum (ER) membrane and, therefore, needed the “help” of wildtype HBs to pass this compartment.

In our studies, we cloned PvMSP1₁₉ (containing 10 cysteines) also N-terminal of HBs and under the control of the cytomegalo virus (CMV) promoter (18). Significantly, even as an episomal vector and in the absence of co-expressing wildtype HBs, transiently transfected COS7 cells were able to produce subviral particles (Fig. 2). This result is probably due to the strong nature of the CMV promoter and to the secretion signal from the human tissue plasminogen activator N-terminal to PvMSP1₁₉. Both might facilitate the transfer of this domain into the ER lumen, thus, dispensing the necessity for additional wildtype HBs for secretion of the recombinant particle. Regardless, the availability of hybrid particles prompted us to determine whether it exposed PvMSP1₁₉ epitopes mimicking natural PvMSP1₁₉.

We previously demonstrated that naturally acquired IgG antibodies from *P. vivax* patients specifically and predominantly recognized PvMSP1₁₉, as opposed to other regions of the PvMSP1 protein (23). Accordingly, we used sera from individual vivax patients in immunoprecipitation assays and clearly demonstrated that the hybrid PvMSP1₁₉-HBs particles exposed epitopes recognized in natural infections (Fig. 1B). We next demonstrated that antibodies elicited by genetic immunization of mice with the recombinant plasmid encoding PvMSP1₁₉ could also immunoprecipitate the hybrid viral particles (Fig. 3). Moreover, these

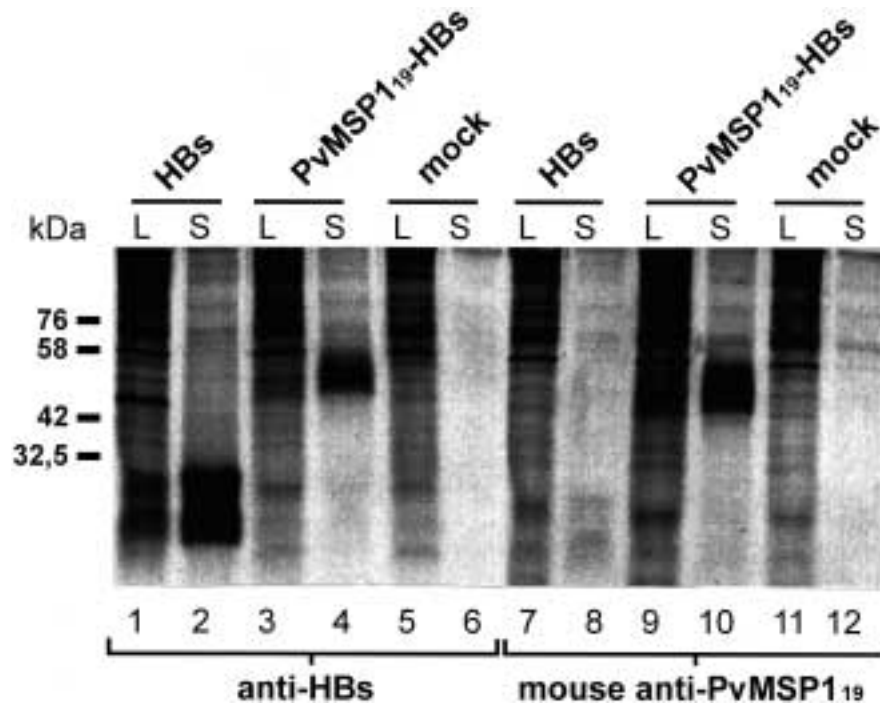


Fig. 3. Sera of VXORF-PvMSP1₁₉-HBs immunized mice recognize COS7 cell expressed PvMSP1₁₉-HBs. COS7 cells were transfected with pSV33M⁻ or VXORF-PvMSP1₁₉-HBs, pulse labeled and cell lysates or supernatants were immunoprecipitated with anti-HBs (lanes 1–6) or pooled sera

derived from VXORF-PvMSP1₁₉-HBs-immunized mice (lanes 7–12; dilution of both sera 1:500). HBs transfection with pSV33M⁻, PvMSP1₁₉-HBs transfection with VXORF-PvMSP1₁₉-HBs, mock transfection with plasmid pVXORF1. HBs, small hepatitis B virus surface protein.

antibodies were able to recognize the native protein in immunofluorescence analysis, further suggesting that genetic immunizations with PvMSP1₁₉-HBs presented a properly folded PvMSP1₁₉, in spite of its fusion with HBs (data not shown). These results further emphasize the importance of this construct, not only as a proteic subunit vaccine candidate, but also as a DNA vaccine candidate. In addition, the antibody titers against HBs were always lower than against the PvMSP1₁₉. This could be due to an intrinsically higher immunogenicity of PvMSP1₁₉ than of HBs; alternatively, PvMSP1₁₉ may render the immunodominant a-epitope in HBs partially inaccessible for immunoglobulins. In studies with a form of CS-HBs (24), in which 16 NANP repeats were fused to HBs, a similar result was obtained, even with a smaller peptide size devoid of any conformational structure mediated by disulphide bonds, as was the case in PvMSP1₁₉.

Although a significant humoral immunity against PvMSP1₁₉ was obtained in the genetic immunizations (Table 1), others correlated pro-

tection of rodents against malaria by immunization with MSP1₁₉ to highest antibody titers (25). Still others attributed it to the presence of antibodies directed against a certain reduction-sensitive epitope contained in the first EGF-like domain (7). Studies employing the MSP1₁₉-HBs produced in yeast and cell culture for the rodent malaria models *P. chabaudi* and *P. yoelii* are currently under way to address these issues. Due the fact that there are very few sequence variations in the *P. vivax* MSP1₁₉ gene which are identical in immunologic terms (26), the herein described hybrid particle may be useful as a component in future vaccine trials against blood stage forms of *P. vivax*.

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