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Judith E. Epstein, … , B. Kim Lee Sim, Stephen L. Hoffman


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Judith E. Epstein,1 Kristopher M. Paolino,2 Thomas L. Richie,3 Martha Sedegah,1 Alexandra Singer,1 Adam J. Ruben,3 Sumana Chakravarty,3 April Stafford,2 Richard C. Ruck,2 Abraham G. Eappen,3 Tao Li,3 Peter F. Billingsley,2 Anita Manoj,3 Joana C. Silva,4 Kara Moser,4 Robin Nielsen,2 Donna Tosh,2 Susan Cicatelli,2 Harini Ganeshan,1 Jessica Case,4 Debbie Padilla,3 Silas Davidson,2 Lindsey Garver,2 Elizabeth Saverino,3 Tooba Mushedkar,3 Anusha Gunasekera,3 Patrick S. Twomey,2 Sharna Reyes,1 James E. Moon,2 Eric R. James,3 Natasha KC,3,6 Minglin Li,3,6 Esteban Abot,1 Arnel Belmonte,1 Kevin Hauns,2 Maria Belmonte,1 Jun Huang,1 Carlos Vasquez,1 Shon Remich,2 Mary Carrington,1 Yonas Abebe,3 Amy Tillman,3 Bradley Hickey,2 Jason Regules,2 Eileen Villasante,1 B. Kim Lee Sim,3,6 and Stephen L. Hoffman3

1Naval Medical Research Center (NMRC), 2Walter Reed Army Institute of Research (WRAIR), Silver Spring, Maryland, USA. 3Sanaria Inc., Rockville, Maryland, USA. 4Institute for Genome Sciences, University of Maryland School of Medicine, Baltimore, Maryland, USA. 5Statistics Collaborative Inc., Washington, DC, USA. 6Protein Potential, Rockville, Maryland, USA.

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**CONCLUSIONS:** We have demonstrated for the first time to our knowledge that PfSPZ Vaccine can protect against a 3-week heterologous CHMI in a limited group of malaria-naive adult subjects. A 3-dose regimen protected against both 3-week and 24-week homologous CHMI (87% and 57%, respectively) in this population. These results provide a foundation for developing an optimized immunization regimen for preventing malaria.

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**Introduction**

In 2014 approximately $2.5 billion was invested in malaria control, and there were at least 214 million clinical cases and 438,000 deaths caused by malaria, with infants and children in Africa bearing the greatest burden (1). Malaria is also a threat to travelers; the US Department of Defense has ranked malaria as its number one infectious threat. A highly effective malaria vaccine would be an ideal tool to
prevent malaria in travelers and deployed military, reduce morbidity and mortality in infants and children, and eliminate malaria from defined geographic areas through mass vaccine administration campaigns. Plasmodium falciparum (Pf) sporozoites (SPZ) are the only immunogens ever shown to induce >90%, sterile protective immunity against malaria (2–4). Until recently, this had only been accomplished by immunization by mosquito bite. Manufacturing processes have now been developed for the production of aseptic, purified, vialed PfSPZ Vaccine (5). In the first clinical trial Sanaria PfSPZ Vaccine administered subcutaneously or intradermally was well tolerated and safe, but poorly protective (6). Studies in non-human primates indicated the vaccine would be protective if administered by intravenous (i.v.) injection (6). In the second clinical trial, 5 doses of PfSPZ Vaccine administered by i.v. injection protected 6 of 6 (100%) subjects against controlled human malaria infection (CHMI) with homologous (same as in vaccine) Pf parasites 3 weeks after final immunization (4). In a third trial, 55% protection was achieved at 14 months after a 4-dose immunization regimen (7).

The WHO Malaria Vaccine Technology Roadmap (2013 update) set a goal for the year 2030 of a malaria vaccine with protective efficacy of at least 75% against clinical malaria, and development of malaria vaccines that reduce transmission of the parasite (8). We have focused on establishment of a regimen that would protect at least 80% of individuals against all Pf infections and could be administered with a maximum of 3 doses. Such a vaccine would meet all WHO objectives. To address these objectives, and to assess the efficiency and tolerability of administering the vaccine by direct venous inoculation (DVI), we designed a study in which numbers of PfSPZ/dose, numbers of doses, and intervals between doses were altered. Protection was assessed at 3 and 24 weeks after the last immunization against homologous Pf parasite challenge and, for the first time to our knowledge, heterologous (different genetically from parasites in vaccine) Pf parasite challenge. Furthermore, we assessed the efficiency and tolerability of DVI administration of PfSPZ.

### Results

Genetic differences between *P. falciparum* isolates

Pf7G8 had been used in the 1990s to establish heterologous protection in volunteers immunized by the bite of irradiated mosquitoes carrying Pf sporozoites (9). To establish that Pf7G8 was representative of a heterologous strain, we utilized whole genome sequencing data to estimate genetic distance between the vaccine strain, PfNF54, and Pf7G8, and between these and 19 clinical isolates from Africa (Supplemental Table 1 and Supplemental Figure 1; supplemental material available online with this article; http://doi.org/10.1172/jci.insight.89154DS1). Comparison of 3D7 (a proxy for its parent PfN54) and 7G8 revealed 22,056 SNPs genome-wide, an average of 0.95 SNPs/kb. 4,925 SNPs fall within a panel of ~106 validated, protein-coding SNPs. This number of SNPs relative to 3D7 is comparable to that from African Pf clinical isolates (Supplemental Figure 1). For 13 selected pre-erythrocytic Pf antigens and vaccine candidates, non-synonomous SNPs in 7G8 relative to NF54 were common in all genes, including 8 in the PICSP gene (Supplemental Table 2), confirming the genetic differentiation between the two strains.
Study population

Forty-five immunized and 24 infectivity control subjects were enrolled. Subjects were recruited from the Baltimore–Washington, DC, area. Baseline demographics were balanced. Study volunteer demographics and baseline characteristics are presented by group for all enrolled subjects (Table 1). Both civilian and military personnel were included. Ten of the 30 subjects in groups 1 and 2 (33%) and 7 of the 15 subjects in group 3 (47%) were military members.

Forty-one of 45 subjects (91%) completed all scheduled vaccinations. In the $4.5 \times 10^5$ PfSPZ group (group 3), all 15 subjects received all immunizations and proceeded to CHMI #1. In the $2.7 \times 10^5$ PfSPZ group (groups 1 and 2), 26 of 30 subjects received all 5 doses were “fully immunized”) and were available for CHMI. Two additional subjects in the $2.7 \times 10^5$ PfSPZ group who did not receive all 5 scheduled immunizations proceeded to CHMI. Of the 28 subjects immunized in the $2.7 \times 10^5$ PfSPZ groups who were available for CHMI, 14 were randomized to group 1 (homologous CHMI) and 14 were randomized to group 2 (heterologous CHMI). Two subjects randomized to heterologous CHMI did not receive CHMI. Of the two remaining subjects who received $2.7 \times 10^5$ PfSPZ immunizations, one subject missed the last immunization and both CHMIs due to deployment overseas, and one subject discontinued from the study after receiving 4 doses (Figure 1).

Randomization was performed using a random generator program (simple randomization). Subjects were informed by a study investigator of their CHMI group approximately 3 weeks prior to first CHMI. Infectivity controls, who did not receive vaccine or placebo, were not recruited and enrolled until the time of each CHMI; thus they were not randomized.

**CHMI #1 at 3 weeks after last vaccine dose.** Fourteen subjects in group 1 (13 of whom who were fully immunized) and 15 subjects in group 3 (all fully immunized) underwent Pf3D7 CHMI. There were 6 Pf3D7 CHMI infectivity controls. A total of 5 fully immunized subjects in group 2 underwent Pf7G8 CHMI. There were 4 Pf7G8 CHMI infectivity controls. One of the 5 vaccinees in the analysis population was bitten by only 3 Pf7G8-infected mosquitoes, and 1 of the 4 infectivity controls was bitten by only 4 Pf7G8-infected mosquitoes. Six of 11 fully immunized subjects in group 2 and 2 of 6 infectivity controls did not undergo Pf7G8 CHMI due to poor mosquito feeding.

**CHMI #2 at 24 weeks after last dose of vaccine.** Eleven subjects in group 1 (10 of whom who were fully immunized), 14 subjects in group 3 (all fully immunized), and 6 infectivity controls underwent Pf3D7 CHMI. Twelve subjects in group 2 (11 of whom who were fully immunized) and 6 infectivity controls underwent Pf7G8 CHMI. One subject in group 2 was dropped from the analysis population because of non-compliance.

**Protective efficacy**

**CHMI #1.** Participants underwent CHMI #1 3 weeks after treatment with the final dose of vaccine (Figure 2, Table 2, Supplemental Table 3, and Supplemental Table 4).

Group 1: Subjects received 5 doses of $2.7 \times 10^5$ PfSPZ (homologous Pf3D7 CHMI). All 6 controls developed parasitemia with a median prepatent period of 11.6 days. Twelve of 13 fully immunized subjects did not develop parasitemia, providing a protective efficacy of 92.3% (95% confidence interval [CI]: 48.0, 99.8, $P = 0.0003$). The prepatent period in the one immunized subject who developed parasitemia was 13.9 days.

Group 2: Subjects received 5 doses of $2.7 \times 10^5$ PfSPZ (heterologous Pf7G8 CHMI). All 4 controls exposed to the bites of 5 ($n = 3$) or 4 ($n = 1$) PfSPZ-infected mosquitoes developed parasitemia, with a median prepatent period of 11.9 days. Four of 5 fully immunized subjects exposed to the bite of 5 ($n = 4$) or 3 ($n = 1$) infected mosquitoes did not develop parasitemia. The protective efficacy of this cohort was 80.0% (95% CI: 10.4, 99.5, $P = 0.048$). The prepatent period in the one fully immunized subject who developed parasitemia was 11.9 days. A fully immunized subject who was exposed to only 3 PfSPZ-infected mosquitoes was one of the 4 subjects counted as protected. One fully immunized subject was excluded from the analysis population due to non-compliance.

Group 3: Subjects received 3 doses of $4.5 \times 10^5$ PfSPZ (homologous Pf3D7 CHMI). Controls were the same as those in group 1. Thirteen of 15 fully immunized subjects did not develop parasitemia, providing a protective efficacy of 86.7% (95% CI: 35.9, 98.3, $P = 0.0005$). The prepatent periods in the 2 immunized subjects who developed parasitemia were 13.9 and 16.9 days.

**CHMI #2.** Participants underwent CHMI #2 24 weeks after treatment with the final dose of vaccine (Figure 3, Table 2, Supplemental Table 3, and Supplemental Table 4).
Figure 1. Consort 2010 flow diagram. Thirty subjects were enrolled to receive 5 doses of $2.7 \times 10^5$ PfSPZ/dose at weeks 0, 4, 8, 12, and 20; subjects were then randomized equally to group 1 (homologous CHMIs) or group 2 (heterologous CHMIs). Fifteen subjects were enrolled to receive 3 doses of $4.5 \times 10^5$ PfSPZ/dose at weeks 0, 8, and 16 with homologous CHMIs (group 3, which began immunizations 1 month after groups 1 and 2). Twenty-four subjects were enrolled as infectivity controls for the total of 4 CHMIs (2 homologous CHMIs and 2 heterologous CHMIs); each CHMI included 6 infectivity controls, except heterologous CHMI #1, for which only 4 infectivity controls were challenged. CHMI, controlled human malaria infection.
Group 1: All 6 controls developed parasitemia, with a median prepatent period of 11.6 days. Seven of 10 fully immunized subjects who underwent CHMI #2 did not develop parasitemia, providing a protective efficacy of 70% (95% CI: 17.3, 93.3, $P = 0.011$). The median prepatent period in the 3 fully immunized subjects who developed parasitemia was 15.4 days.

Group 2: All 6 controls developed parasitemia, with a median prepatent period of 10.9 days. One of 10 fully immunized subjects did not develop parasitemia, providing a protective efficacy of 10.0% (95% CI: –35.8, 45.6, $P = 1.00$). The subject who did not develop parasitemia was the subject who did not develop parasitemia in the first CHMI when exposed to the bites of 3 PfSPZ-infect ed mosquitoes. The median prepatent period in the 9 fully immunized subjects who developed parasitemia was 11.9 days. When comparing the prepatent period in the group 2 subjects who developed parasitemia with the controls, there was a statistically significant delay of 1 day (Wilcoxon rank-sum exact test 2-sided $P = 0.005$). One fully immunized subject was excluded from the analysis population due to non-compliance.

Group 3: Controls were the same as those in group 1. Eight of 14 fully immunized subjects did not develop parasitemia, providing a protective efficacy of 57.1% (95% CI: 21.5, 76.6, $P = 0.042$). The median prepatent period in the 6 immunized subjects who developed parasitemia was 14.0 days.

Results for the challenged population, including subjects who missed one or more immunizations, are presented in Supplemental Table 3. All protected individuals were negative for Pf parasites by real-time quantitative PCR (qPCR) on their visits 11, 18, and 28 days after CHMI. All individuals positive by thick blood smear (TBS) were positive by qPCR.

**Antibody responses**

**PfCSP and PfSPZ.** Antibodies against PfCSP by ELISA, PfSPZ by automated immunofluorescence assay (aIFA), and PfSPZ by inhibition of sporozoite invasion (ISI) assay in sera taken 2 weeks after the last dose of vaccine and just prior to the second CHMI (~24 weeks after last dose of vaccine) for the 3 groups are shown in Figure 4, A–C. The median response (bar with number) and those protected (filled symbols) and not protected (open symbols) are shown. The numbers not protected after CHMI #1 were so low that meaningful associations with protection could not be done. However, in group 3 the unprotected subjects had the lowest levels of antibodies to PfCSP, both less than 3,500 (Figure 4A and Supplemental Table 4). Since there were only 2 unprotected subjects, meaningful statistical analysis could not be done. Antibodies diminished between the last dose and the 24-week CHMI. The geometric mean (GM) half-life of PfCSP antibodies in groups 1 and 2 was 17.81 weeks ($n = 21$, range 9.61 to 61.76 weeks), and in group 3 was 19.93 weeks ($n = 14$, range 7.78 to 180.88 weeks). There was no significant association between antibodies by any assay (ELISA, aIFA, or ISI) prior to CHMI #2 and protection.

Approximately 77% of the antibodies to the rPfCSP in the ELISA were directed against the NANP repeat region of the PfCSP (Supplemental Table 5). By aIFA against PfSPZ, median IgG antibodies were 11.2-fold higher than median IgM antibodies (Supplemental Figure 2). For all 33 subjects in groups 1–3 who underwent CHMI 3 weeks after last dose of vaccine, median IgG antibodies to NF54 PfSPZ were 1.7-fold higher than to 7G8 PfSPZ by aIFA in sera taken 2 weeks after the last dose of vaccine (2,668 arbitrary
Other antigens. Two weeks after the fifth dose in groups 1 and 2 and after the third dose in group 3, there were no antibodies to PfMSP1, PfEBA175, PfCelTOS, PfEXP1, or PfLSA1 (Supplemental Table 6). At the same time points, there were antibodies to, (a) PfMSP5 in 9 of 26 (35%) subjects in groups 1 and 2, and 12 of 15 (80%) in group 3 (P = 0.009, Fisher’s exact test, 2-tailed); (b) PfAMA1 in 12 of 26 (46%) subjects in groups 1 and 2, and 1 of 15 (6.7%) in group 3 (P = 0.014); and (c) PfSSP2/TRAP in 6 of 26 (23%) subjects in groups 1 and 2, and 2 of 15 (13%) in group 3 (P = 0.687). Because of the low numbers of unprotected vaccinees, we could not establish whether any antibody responses were associated with protection.

Cellular immune responses

Cellular immune responses to stimulation with irradiated PfSPZ were assessed in PBMCs from fresh blood by FluoroSpot 2 (IL-2 and IFN-γ) 2 weeks after the first dose of PfSPZ Vaccine, 2 weeks after the last dose of vaccine (1 week before CHMI#1), and 23 weeks after the last dose of vaccine (1 week before CHMI #2). Responses were in general 1.5–3 times higher after the first dose than 2 weeks after the last dose of vaccine, but there were no significant differences between these time points. There were essentially no measurable responses 1 week prior to CHMI #2, 23 weeks after the last dose of vaccine; all of these responses were significantly lower than prior to CHMI #1 (Kruskal-Wallis test, all P < 0.0002). There was no clear association between these cellular immune responses to PfSPZ or Pf-derived synthetic peptides and protection (Figure 4, D–F, Supplemental Figure 3, and Supplemental Table 4).

Ease of administration, tolerability, and safety

DVI was rapid, required a single stick 96% (182 of 190) of the time, and was well tolerated. Subjects reported no pain 73% of the time and mild pain 24% of the time (Supplemental Table 7).

Details of adverse events (AEs) are presented in Tables 3 and 4 and Supplemental Tables 8–10. Briefly, of the 66 solicited AEs reported within 7 days of immunization considered related, 61 (92%) were grade 1 and 5 (8%) grade 2. No subject experienced a grade 3 or 4 solicited AE. All unsolicited AEs reported within 7 days of immunization were grade 1. The incidence of AEs was not increased in subjects who received 4.5 × 10^5 PfSPZ as compared to 2.7 × 10^5 PfSPZ (Supplemental Table 8). The incidence of AEs
did not increase as subjects received additional doses of the vaccine (Supplemental Tables 9 and 10). 31 of the 45 immunized subjects (75%) had at least one abnormal laboratory value within 7 days of an immunization, but there was no association with dosing schedule, size, or sequence (Table 4). Details on laboratory abnormalities are provided in Supplemental Table 11.

Discussion

An important first step in our development program is to optimize the immunization regimen for non-immune individuals who travel to malarious areas. Their risk was underscored by the recent deaths from malaria of two airline flight attendants after exposure of as little as 1 day (10, 11). For this population we are aiming for a vaccine that is easy to administer, extremely well tolerated, and safe and at a minimum prevents all Pf erythrocytic-stage parasitemia in at least 80% of recipients for 3 months (short-term travel) and ideally 6 months or longer.

Our trial designs have been based on the hypothesis (Figure 5) that increasing the number of PfSPZ/dose and altering intervals between doses will lead to moving from no or minimal protection to high-level (>80%) short-term protection against homologous parasites (threshold 1); high-level short-term protection against heterologous parasites (threshold 2); high-level long-term protection against homologous parasites (threshold 3); and eventually high-level long-term protection against heterologous parasites (threshold 4). In the first successful trial we reached threshold 1, but did not assess for thresholds 2 and 3 (4).

In this trial, short-term protection against homologous and heterologous CHMI was ≥80%. At 3 weeks after the last dose of vaccine, 92% of 13 subjects immunized with a total of 1.35 × 10^6 PfSPZ in a 5-dose regimen, and 87% of 15 subjects immunized with 1.35 × 10^6 PfSPZ in a 3-dose regimen were protected against homologous CHMI, reaching threshold 1. Furthermore, this trial gave us an indication that there was short-term protection (threshold 2) against heterologous CHMI. All 4 controls developed parasitemia, and 4 of the 5 vaccinees did not develop parasitemia (protective efficacy of 80%, 95% CI: 10.4, 99.5).

At 24 weeks after the last dose of vaccine, 70% of 10 subjects immunized with the 5-dose regimen and 57% of 14 subjects immunized with the 3-dose regimen were protected against homologous CHMI, nearly reaching the 80% level required to achieve threshold 3. Furthermore, recent data indicate that if protected at 6 months, vaccinees will be protected at ~14 months (7).

However, only 10% of 10 subjects immunized with the 5-dose regimen were protected against heterologous CHMI. Nonetheless, there was a significant difference in time to onset of parasitemia in immunized versus control subjects (Figure 3), suggesting vaccine-induced immunity had prevented infection of or eliminated most liver-stage parasites (12, 13). Based on our hypothesis (Figure 5), in the next trial we will immunize with higher doses of PfSPZ in 3-dose regimens. If this is not as effective as expected, we may move to a vaccine containing a mixture of strains of Pf.

PfSPZ Vaccine was well tolerated and safe. A study done in Mali with the same dosage regimen with a blinded control group had the same incidence rates of AEs in vaccine and normal saline placebo recipients (Mahamadou S. Sissoko, Malaria Research and Training Center, Mali-NIAID ICER, University of Science, Techniques and Technologies of Bamako, Mali; and Sara A. Healy, Laboratory of Malaria Immunology and Vaccinology, National Institute of Allergy and Infectious Diseases, NIH; personal communication). In our
study injection by DVI took only 10–15 seconds after the injection site was prepared and caused no pain in 73% of immunizations administered, mild pain in 24%, and moderate pain in 3% (Supplemental Table 7). Administration of PfSPZ Vaccine by DVI appears to be much better tolerated than vaccines administered by traditional routes, likely because all vaccine leaves the site of administration rapidly and there are no inflammatory components in the vaccine. In traditional administration, the most discomfort is experienced when the vaccine is injected and for several hours afterward. Most subjects who received PfSPZ Vaccine by DVI did not sense injection. This will be an advantage in mass vaccine administration campaigns, and in acceptability by infants, children, and travelers.

Through the mid-1980s it was thought that antibodies to PfSPZ, particularly to the PfCSP, that blocked sporozoite invasion of hepatocytes were responsible for irradiated SPZ-induced protection (14, 15). However, when it was shown that treatment of fully immune mice (6, 16–18) and non-human primates (19) with anti-CD8 antibodies eliminated the protective immunity, it was hypothesized that cellular immune responses against the developing liver stages were responsible for protection (16, 17, 20–23). We could not use statistics to assess correlations between antibodies and protection for CHMI #1, and there was no significant correlation between antibodies and protection for CHMI #2 (Figure 4). This is consistent with our

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**Table 3. Solicited and unsolicited adverse events**

<table>
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<tr>
<th>Solicited AE at any immunization</th>
<th>Groups 1 and 2 (5 doses of 2.7 × 10⁵ PfSPZ), n = 30</th>
<th>Group 3 (3 doses of 4.5 × 10⁵ PfSPZ), n = 15</th>
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<tr>
<td>n (%) with grade 1 AE 14/30 (47)</td>
<td>n (%) with grade 2 AE 2/30 (7)</td>
<td>n (%) with grade 1 AE 5/15 (33)</td>
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<td>Pain 0 0 0 0</td>
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<tr>
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<td>Fatigue 4 (13)</td>
<td>0 0 2 (13) 0</td>
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<tr>
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<td>Malaise 5 (17)</td>
<td>0 0 0 0</td>
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<td>Myalgia 3 (10)</td>
<td>0 0 1 (7) 0</td>
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<tr>
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<tr>
<td>Pain in extremity 1 (3)</td>
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</table>

*Events occurring within 7 days of any immunization, by group, deemed possibly, probably, or definitely related to immunization and laboratory abnormalities occurring within 7 days of any immunization. No solicited or unsolicited AEs of grade 3 or 4 were reported. Column head counts and denominators are the number of volunteers receiving at least one immunization. Volunteers are counted at most once within each row and under the highest grade reported for related events within 7 days of an immunization.*
Figure 4. Antibody and T cell responses in all immunized participants. For all assays, protected subjects are shown as filled (black) circles and unprotected subjects as open circles. For each of the 3 immunization groups, the interquartile ranges and the median values of responses of subjects in each group are shown. Assessment of antibodies (A–C) was performed in sera from subjects before immunization and 2 and 23 weeks after the last dose of PfSPZ Vaccine, time points that were ~1 week before CHMI #1 and CHMI #2, respectively. Number of samples assessed in A–C for CHMI #1: group 1 (n = 13), group 2 (n = 5), group 3 (n = 15); CHMI #2: group 1 (n = 10), group 2 (n = 10), group 3 (n = 14). (A) Antibodies to PfCSP by ELISA are reported as net OD 1.0 (the difference in OD 1.0 between post- and pre-immunization sera). (B) Antibodies to PfSPZ by aIFA are reported as the reciprocal serum dilution at which the fluorescence units were 2 × 10^5 (AFU 2 × 10^5). (C) Percent inhibition of PfSPZ invasion is reported as the percent reduction of the numbers of PfSPZ that invaded a human hepatocyte line (HC-04) in the presence of post-immunization as compared with pre-immunization serum from the same volunteer, both at a dilution of 1:5. At both CHMI #1 and CHMI #2, all 3 assays correlated with each other (CHMI #1: PfCSP ELISA vs. aIFA, P < 0.001, r² = 0.61; PfCSP ELISA vs. ISI, P = 0.0002, r² = 0.30; aIFA vs. ISI, P = 0.004, r² = 0.19. CHMI #2: PfCSP ELISA vs. aIFA, P < 0.0001, r² = 0.54; PfCSP ELISA vs. ISI, P < 0.0001, r² = 0.56; aIFA vs. ISI, P = 0.0002, r² = 0.34). Assessment of T cell responses by FlouroSpot assay was performed using PBMCs from subjects prior to immunization, 2 weeks after the first dose of PfSPZ Vaccine, and just prior to CHMI #1 and CHMI #2 (D–F). Number of samples assessed in D–F: 2 weeks after dose 1: group 1 (n = 9), group 2 (n = 5), group 3 (n = 15), pre-CHMI #1: group 1 (n = 13), group 2 (n = 5), group 3 (n = 15); pre-CHMI #2: group 1 (n = 10), group 2 (n = 10), group 3 (n = 13). Results are reported as spot-forming cells (SFCs) per 10^6 PBMCs secreting (D–F) IFN-γ only, (E) IL-2 only, or (F) IFN-γ and IL-2. Individual data points have the pre-immunization SFCs/10^6 PBMCs subtracted from the post-immunization SFCs/10^6 PBMCs. aIFA, automated immunofluorescence assay; ISI, inhibition of sporozoite invasion.
hypothesis that antibodies can reduce the numbers of PfSPZ that reach or successfully invade hepatocytes, but cellular immune responses directed against perhaps hundreds of antigens expressed in developing liver stages are required for the sterile protective immunity induced by PfSPZ Vaccine (4, 6).

The cellular immune responses delineated by the FluoroSpot assays peaked after the first dose, were lower after the last dose, and essentially absent at the time of CHMI #2. There was no significant association with protection. This may be because, as in non-human primates, the majority of PfSPZ-specific CD8+ cells are in the liver, where they can attack the Pf-infected hepatocyte, not in the blood, where we can measure them (6). Given that there has been association between T cell responses and protection as assessed by flow cytometry (4, 7), we are proceeding with such an analysis. The low levels of antibodies (Supplemental Table 6) and T cell responses (Supplemental Figure 3) against well-known antigens suggest that as-yet-undefined antigens may be critically important targets of this strong protective immunity, and these may be revealed by flow cytometry studies.

We believe CHMI with heterologous Pf strains will be adequate to predict protective efficacy in the field against heterogeneous Pf populations. This is supported by our study in malaria-exposed Malian adults immunized with the same regimen as groups 1 and 2 in this study, which conferred 28.5% sterile protection against highly heterogeneous naturally transmitted Pf for up to 6 months (M.S. Sissoko and S.A. Healy, personal communication) — several fold better protection than the 10% protective efficacy against Pf7G8 at 24 weeks in the present study. Recent evidence has suggested that the RTS,S/AS01 vaccine induces strain-specific protection in the field (24). Only homologous CHMI has been used in the development and testing of RTS,S/AS01. It is likely that heterologous CHMI would have identified this deficiency early on. It is for this reason we have conducted the first heterologous CHMI, and will continue to use heterologous CHMI in the development of PfSPZ Vaccine.

Whole genome sequencing revealed that the genetic distance between the vaccine strain (Pf3D7, a proxy for its parent strain PINF54) and Pf7G8, the two strains used for CHMI, was substantial (22,056 SNPs). Furthermore, a targeted analysis of 13 known pre-erythrocytic antigens revealed that they all contain non-synonymous SNPs between Pf3D7 and Pf7G8. Thus, we conclude that Pf7G8 is an appropriate parasite for heterologous CHMI.

By increasing the dose (Figure 5), we anticipate we will increase the magnitude of protective immune responses against all target antigens, and will expand the immune responses to be direct against subdominant protective and conserved epitopes. They are conserved because there is no immune pressure against these subdominant epitopes, and this should allow for protection against all strains of Pf. We anticipate finalization of
The information from CHMI studies of PfSPZ Vaccine lays a foundation that supports trials designed to finalize a vaccine regimen for mass vaccination program campaigns in Africa.

**Methods**

**Study oversight**

The study was conducted in accordance with the principles of the Declaration of Helsinki and standards of Good Clinical Practice defined by the International Conference on Harmonisation of Technical Requirements for Pharmaceuticals for Human Use.

The NMRC, WRAIR, and Sanaria (sponsor) collaborated on design. There was an independent safety monitoring committee (SMC).

**Study subjects**

Malaria-naive, 18- to 45-year-olds were enrolled during 2014–2015 after providing informed consent. Subjects were excluded if they had a history of malaria infection, travel to a malaria endemic region within 6 months of first immunization, or long-term residence (>5 years) in an area known to have transmission of *P. falciparum*. Subjects were also excluded if they had previously participated in a malaria vaccine trial. All subjects underwent a screening evaluation consisting of a medical history, physical examination, electrocardiogram, complete blood count, clinical biochemistries, urinalysis, sickle cell testing, and serological studies for previous exposure to or infection with HIV, hepatitis B, and hepatitis C. Subjects were excluded if they had hepatic, renal, or autoimmune diseases, allergy to any study component, splenectomy, or evidence of increased cardiovascular disease risk (defined as >10%, 5-year risk) as determined by the method of Gaziano (25). All females had urine pregnancy test at screening, immediately before each immunization and before CHMI; they were to be excluded from further immunization or CHMI if this was positive. All female subjects agreed to use effective means of birth control for the duration of the trial.

**PfSPZ Vaccine**

Sanaria PfSPZ Vaccine is a live attenuated whole parasite vaccine (26) stabilized in liquid nitrogen vapor phase (LNVP), and manufactured and characterized (Supplemental Table 12) as described previously (4–6). Vaccine in 0.5 ml was rapidly (seconds) injected in an arm vein by DVI through a 25-gauge needle.

**Study design**

The study was open label and designed to assess the tolerability, safety, immunogenicity, and protective efficacy of DVI-administered PfSPZ Vaccine in 3 groups of 15 immunized subjects each. Thirty subjects were to receive 5 doses of 2.7 × 10⁵ PfSPZ/dose at weeks 0, 4, 8, 12, and 20 and then were to be randomized equally to group 1 (homologous CHMIs) or group 2 (heterologous CHMIs). Fifteen subjects in group 3 were to receive 3 doses of 4.5 × 10⁵ PfSPZ/dose at weeks 0, 8, and 16, followed by homologous CHMIs.

Tolerability of DVI was assessed immediately after injection by the subjects using a pain scale (none, mild, moderate, or severe). Subjects were followed up in the clinic on days 2, 7, and 14 after each immunization. Solicited and unsolicited AEs were monitored for 7 and 28 days after each immunization, respectively. Safety laboratories were assessed 2 and 7 days after each immunization. A TBS was assessed 14 ± 2 days after the first immunization to rule out breakthrough infection. See supplemental material for details of AE recording and grading.
CHMI was conducted 3 and 24 weeks after the final immunization. Groups 1 and 3 underwent CHMI by bites of 5 *Anopheles stephensi* mosquitoes infected with Pf3D7 parasites (27), which are a clone of the NF54 parasites in the vaccine (presumed African origin) (homologous CHMI). Group 2 underwent CHMI by bites of 5 *A. stephensi* infected with Pf7G8 parasites (South American) (heterologous CHMI) (28). As is standard for malaria vaccine trials in malaria-naive subjects with CHMI, for each CHMI there were non-immunized infectivity controls. From day 7 to day 18 after exposure to the bites of PfSPZ-infected mosquitoes, subjects spent evenings and nights in a hotel. AEs and TBSs were assessed at least daily from day 7 to day 18 after CHMI, and then again on days 20, 22, 25, and 28 after initiation of CHMI or until detection of parasitemia as previously described (13). qPCR was done retrospectively (see supplemental materials for specifics) (29, 30). Clinical laboratory tests were also monitored. Upon detection of parasites, subjects were treated with a 3-day course of atovaquone/proguanil (Malarone).

Assessment of immunological responses

*Antibody assays.* Sera were assessed for antibodies to selected Pf proteins by ELISA, and to Pf sporozoites by aIFA. Functional activity of sera was assessed by the ISI assay. See supplemental materials for specifics.

*Cellular immunology assays.* PBMCs were assessed for activity against PfSPZ and Pf peptides by FluoroSpot assay (Mabtech). See supplemental material for specifics.

Assessment of genetic differences between *P. falciparum* isolates

SNP calling utilizing previously generated whole-genome short-read sequencing data was used to estimate genetic distance between the vaccine strain PfNF54 and Pf7G8, as well as between these and 19 clinical isolates from Africa (Supplemental Table 1 and Supplemental Figure 1), with details described in the supplemental material. Given the negligible difference (55 detectable SNPs) between the Pf3D7 clone and its parent, PfNF54, 3D7 (27) was used as a proxy for NF54 (31).

Statistics

Sample sizes for the groups were selected based on comparison of the number of immunized subjects per group versus 6 infectivity controls for each CHMI. Accounting for a dropout rate of immunized subjects of 20% (from 15 to 12 subjects), a sample size of 6 in the control group and 12 in the vaccinated group was selected to be able to show with a power of 80% without a continuity correction that if 5 of 6 become infected in the control group and 2 of 12 become infected in the immunized group, the proportion of subjects infected in the control group would be significantly different from the proportion of subjects infected in the immunized group (α < 0.05, 2-sided Fisher’s exact test).

Immunological analyses, differences in responses were analyzed by nonparametric tests.

Protective efficacy was calculated as 1 – (relative risk of parasitemia infection). Exact CIs for protective efficacy were calculated by inverting two 1-sided tests for the score statistic in StatXact 9 (Cytel Studio 9). Statistical comparison tests of protective efficacy were performed using 2-sided Fisher’s exact tests comparing vaccinated subjects with controls within similar CHMI strains. The Hochberg procedure was applied to the comparisons of the 3D7 strain CHMIs (group 1 versus control and group 3 versus control for both CHMIs), for a total of 4 comparisons to preserve the overall type I error rate at 0.05 (32). A hierarchical testing procedure was applied to the comparison for the 7G8 strain CHMIs (group 2 versus controls). Because a priori we believed that protection was more likely at 3 weeks than at 24 (see Discussion and Figure 5), we planned to perform the 3-week comparison first at a type I error rate of 0.05 and, only if that was significant, use 0.05 for the 24-week CHMI. Analyses were performed on two populations: (a) the analysis population and (b) the challenged population. The analysis population consisted of fully immunized subjects who participated in the respective CHMI. The text of this article reports results for the analysis population. The challenged population consisted of all subjects who received the respective CHMI, regardless of the number of vaccinations received or compliance status. Results for the challenged population are shown in Supplemental Table 3.

Study approval

The IRB of WRAIR approved the protocol in compliance with applicable federal regulations. Written informed consent was obtained from all subjects prior to inclusion in the study.
Author contributions

JEE, KMP, BKLS, S Reyes, EV, TLR, and SLH contributed to the study design. Protocol development and conduct of the clinical trial were performed by JEE, KMP, MS, A Singer, A Stafford, RCR, RN, DT, S Cicatelli, SD, LG, ES, PT, S Reyes, JEM, CV, S Remich, BH, and JR. AJR, S Chakravarty, AG, TL, PFB, AM, ERJ, NKC, ML, YA, BKLS, and SLH were involved with vaccine manufacture. Regulatory support was provided by TM and AG. MS, S Chakravarty, JCS, HG, KM, NKC, ML, EA, AB, KH, MB, JH, and BKLS performed the assays. EA, AB, and MS carried out the thick smear microscopy. JEE, TLR, AJR, JC, AT, DP, and SLH conducted the data analysis. JEE, TLR, AJR, and SLH wrote the manuscript, and MS, S Chakravarty, JC, S Reyes, MC, YA, and AT assisted with final manuscript preparation.

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Address correspondence to: Stephen L. Hoffman, Sanaria Inc., 9800 Medical Center Drive, Suite A209, Rockville, Maryland 20805, USA. Phone: 240.403.2701; E-mail: slhoffman@sanaria.com.

KMP’s present address is: Center for Global Health and Translational Sciences, Upstate Medical University, New York, USA. AS’s present address is: Virginia Cancer Specialists, Arlington, Virginia, USA. RCR’s present address is: U.S. Army’s Health Clinic–Schofield Barracks, Schofield Barracks, Hawaii, USA. RN’s present address is: Clinical Research Management, Frederick, Maryland, USA. DP’s present address is: University of Virginia, Charlottesville, Virginia, USA. SD’s present address is: Armed Forces Research Institute of Medical Science, Bangkok, Thailand. KH’s present address is: U.S. Army Medical Directorate–Armored Forces Research Institute of Medical Services, Bangkok, Thailand. CV’s present address is: Bio Clinica Inc., Newark, California, USA. MC’s present address is: USS Ashland LSD 48. BH’s present address is: Navy Experimental Diving Unit Medical Department, Panama City, Florida, USA. JR’s present address is: Department of Clinical Research, U.S. Army Medical Research Institute of Infectious Diseases, Frederick, Maryland, USA.


