

Maria M. Mota · Ana Rodriguez *Editors*

Malaria

Immune Response to Infection and
Vaccination

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and Vaccination

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Whole-Sporozoite Malaria Vaccines

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1 Introduction

More than two centuries have passed since Edward Jenner's breakthrough postulates on whole-organism vaccination and more than one since Pasteur's formulation of the idea of immunization through microorganism attenuation [1]. Nowadays, health-care systems around the world rely on numerous successful vaccines to prevent morbidity and mortality in a cost-effective manner [2]. However, despite significant advances in molecular biology, approximately half of all currently approved vaccines continue to rely on the use of live attenuated microorganisms to elicit immunity against the diseases caused by their non-attenuated counterparts [1].

Malaria is one of the most important infectious diseases for which an effective vaccine is still lacking [3]. The gold standard strategies for malaria immunization rely on the administration of live *Plasmodium* sporozoites, the malaria parasite form transmitted to man by mosquitoes. Sporozoite-based vaccination strategies aim at preventing the parasite's life cycle progression from hepatic stages to the symptomatic blood stages of infection while eliciting potent preerythrocytic immune responses. Such whole-sporozoite malaria vaccination strategies are unique in their potential to induce sterile protection against a new infection and have led to the development of various vaccine candidates, currently ongoing preclinical and clinical development. This chapter will explore in detail our current knowledge of the protective immunity elicited by *Plasmodium* sporozoite-based malaria vaccination.

Although it was not until the 1960s that protective immunity against human malaria was first demonstrated [4, 5], the long list of classical studies exploring the use of sporozoites for immunization against malaria spans over a century, with the first studies using heat-inactivated avian parasites dating back to 1910 [6]. However, it was the discovery of the preerythrocytic stages of *Plasmodium* [7], followed by the establishment of a mouse model of malaria [8], that enabled the laboratory production of all stages of the parasite's life cycle [9], which eventually led to the landmark demonstration that live sporozoites attenuated by X-irradiation (RAS) could be used to elicit sterile protection against a new infection [10, 11]. This discovery was soon expanded to humans with the demonstration that volunteers could be protected against homologous and heterologous strains of *P. falciparum* parasites [12–14] (Fig. 1).

The excitement generated by these successes in both animal models and humans ensured that a large number of studies would continue into the following decades, aiming at optimizing the potential of sporozoite-based immunization, as well as at characterizing the immune responses elicited by these strategies [36, 60–64]. Progress also included exploring alternative methods that did not rely on the use of radiation, such as the administration of sporozoites under the cover of antimalarial drugs [37, 38]. Such strategies are based on sporozoite administration with the concomitant administration of a drug that does not interfere with the parasite's ability to complete liver-stage development but prevents blood-stage replication, enabling the induction of a strong immune response in the liver while avoiding the onset of pathology.

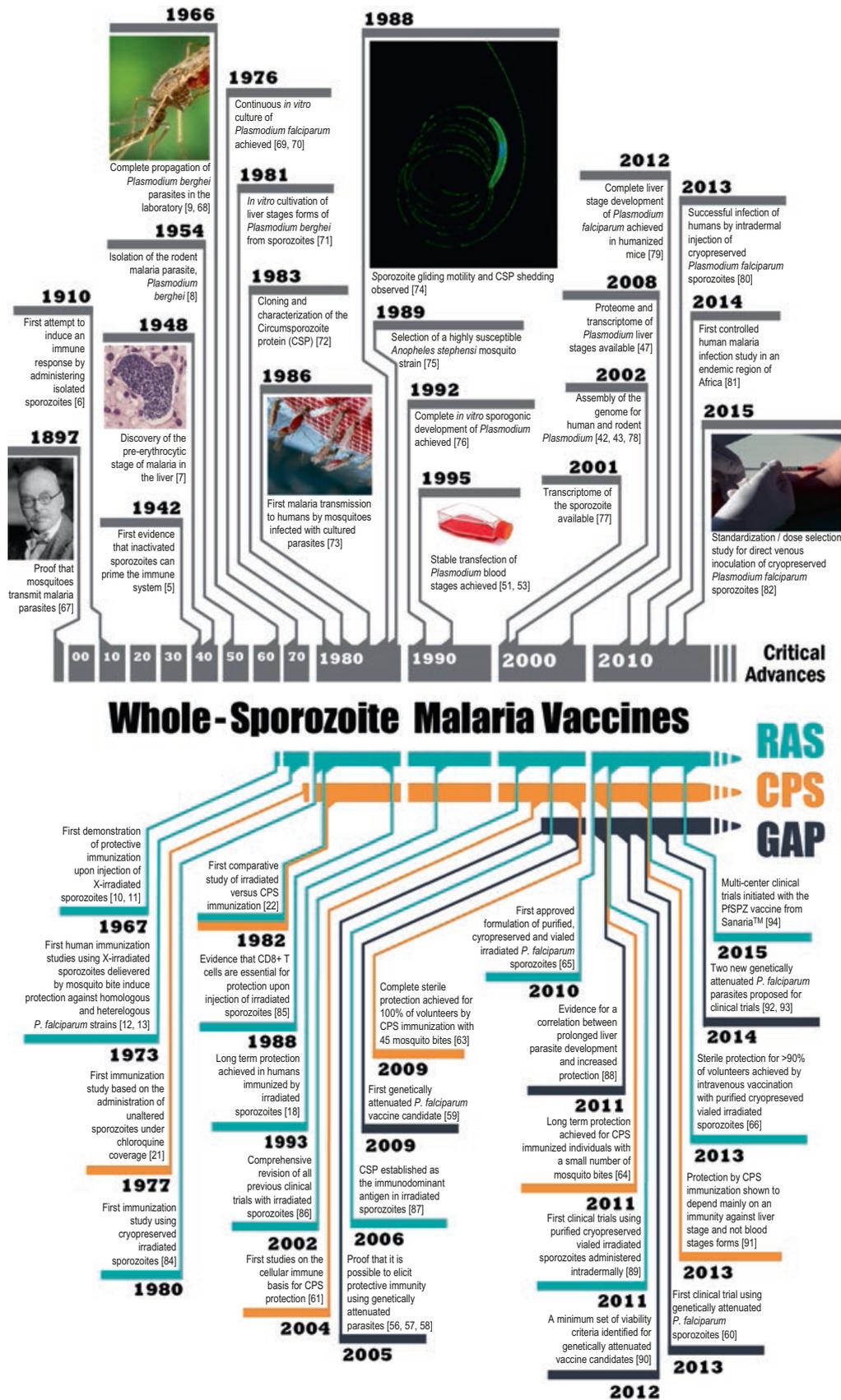


Fig. 1 Timeline of critical events in whole-sporozoite malaria vaccine development. Top [5–9, 15–35], bottom [10, 13, 36–59]

Nonetheless, some of the potential limitations of sporozoite-based vaccination strategies were already evident in the 1970s [65]. Technical procedures to manufacture and store the large amounts of sporozoites required for immunization, as well as to provide safe and efficacious administration to humans, were always a concern and justified the skepticism with which some of these advances were received [66]. These difficulties, along with extraordinary developments in molecular biology methods, led to a substantial shift from sporozoite-based vaccines to subunit vaccination approaches, based on only one or a few parasite antigens with high immunogenic potential [67]. Drawing upon studies aiming to characterize the basis of the protective responses elicited by X-radiation-attenuated sporozoites, the circumsporozoite protein (CSP) [68–70] emerged as the strongest contender for a subunit vaccine and constitutes the basis of the currently most advanced of such vaccine candidates, RTS,S [71–79]. However, the initial optimism generated by RTS,S over 20 years ago has been steadily decreasing over the years [80]. The latest field studies in endemic regions show that vaccine efficacy is substantially inferior to what are the commonly accepted standards for mass distributed vaccines [81–83].

As the reduced efficacy of subunit vaccines became increasingly evident at the beginning of the new century, a renewed call for the development of sporozoite-based immunization strategies took place. At around the same time, malaria research was also entering into the post-genomic era, with the genome sequences of various *Plasmodium* parasite species becoming available, as well as transcriptomic and proteomic data from different parasite developmental stages [15–17, 84–87]. This knowledge fed the search for genes playing essential roles in distinct points of the parasite's life cycle, which could be targeted for deletion using well-established methodologies for stable transfection of *Plasmodium* parasites [18, 19, 88–92]. This led to the design and construction of genetically attenuated parasites (GAPs), whose liver-stage development is arrested by deletion of specific gene(s). Initial studies in mice showed that GAPs were able to elicit a strong immune protection [39, 40, 93]. Shortly afterwards, the first *P. falciparum* GAPs were developed, paving the way for the clinical development of genetically attenuated sporozoites as human vaccine candidates [41, 42]. Concomitantly, strategies that entailed the induction of immunity through sporozoite administration under chemoprophylaxis (CPS) were brought back to the spotlight, with pivotal studies being carried out in mice [43, 94] and in humans [44, 45]. Also in the last decade, substantial progress was made towards overcoming some of the technical hurdles faced by RAS vaccination [46], which eventually culminated in the demonstration of the induction of sterile protection by the intravenous injection of aseptically purified, metabolically active, radiation-attenuated *P. falciparum* sporozoites [47] (Fig. 1).

Recent clinical studies played an important role in consolidating and extending the knowledge accumulated throughout the years regarding the immune responses elicited by sporozoite-based immunization approaches. We now possess a significant understanding of the protective mechanisms at play during whole-sporozoite vaccination. Undoubtedly, our ever-increasing knowledge of these mechanisms will play an instrumental role in guiding future efforts toward the development of a truly effective malaria vaccine.

2 Radiation-Attenuated Sporozoites (RAS)

Radiation-attenuated sporozoites (RAS) have been considered the “gold standard” of whole-sporozoite vaccination [95]. The concept of RAS vaccination was originally described in the late 1960s, when sterile protection from *Plasmodium* challenge was achieved in mice immunized by the mosquito bite administration of *P. berghei* sporozoites attenuated through the X-irradiation of the mosquito vector [10]. These initial studies with rodents paved the way for early studies in humans that established that immunization of volunteers through the bites of irradiated mosquitoes carrying *P. falciparum* or *P. vivax* sporozoites afforded protection against a subsequent *P. falciparum* or *P. vivax* infection [12–14, 96–98]. Between 1989 and 1999, an additional 11 volunteers were immunized by the bites of irradiated *P. falciparum*-infected mosquitoes at the Naval Medical Research Center. Of note, volunteers in these studies averaged nine immunizations with a mean of over 1000 immunizing mosquito bites per volunteer over 9–10 months before the first challenge [51]. Challenges were performed by the bites of mosquitoes infected from cultured *P. falciparum* gametocytes [26, 99], which has since become the method of choice for controlled human malaria infections (CHMIs). A total of 26 challenges were performed in these studies, and protection was observed in 24 of them, with the reported persistence of protective immunity of at least 42 weeks [51].

Despite the protective immunity elicited by these immunization protocols, for a long time, large-scale vaccination with irradiated sporozoites was considered unfeasible, due to the large numbers of parasites required and the need to deliver them alive by infected mosquitoes. As we will see, the twenty-first century brought along significant developments in this regard, to the extent that many researchers no longer consider RAS vaccination impractical. However, recent advances notwithstanding, earlier studies not only established the potential of attenuated sporozoites for malaria vaccination but also began to shed light on the immune responses that govern sterilizing immunity against *Plasmodium* infection.

2.1 Immune Correlates in RAS Immunization

2.1.1 Antibody and T-Cell Responses in Mice and Monkeys

Antibodies have been shown to be effective at limiting the number of parasites that successfully reach the liver [100]. CSP is the most abundant protein on the surface of *Plasmodium* sporozoites and plays a pivotal role in their characteristic gliding motility [101] (Fig. 1). Antibodies against the repeat region of CSP function by blocking both sporozoite motility and invasion of host cells [102]. Passive protection of mice and monkeys by monoclonal or polyclonal antibodies against the central repeat of *P. berghei*, *P. yoelii*, or *P. vivax* CSP has been amply demonstrated (reviewed in [103]). The efficacy of the protective humoral responses targeted to the

extracellular, infective sporozoite depends on the concentration and specificity of the anti-sporozoite antibody response [104, 105]. However, besides humoral responses, cell-mediated immune responses of the RAS-immunized host also play a crucial role in the generation of immunity against sporozoite challenge.

CD8⁺ T cells have been shown to be important for eliminating parasites that successfully invade and replicate within hepatocytes, making the liver stage the primary target for the vaccine-inducible T-cell responses and the presumed target of the RAS vaccination model [100]. It has been suggested that antigens present in the skin following sporozoite injection are presented in the skin-draining lymph nodes, where they prime specific CD8⁺ T cells which then migrate to the liver [106], where CD8⁺ T-cell-mediated elimination of *Plasmodium* hepatic stages occurs [107]. The transporter associated with antigen processing (TAP)-associated processing of exoerythrocytic antigens was also shown to be critical for the presentation of liver-stage antigens by infected hepatocytes [108], and, very recently, it was proposed to play a crucial role in CD8⁺ T-cell-mediated, *P. berghei* RAS-induced sterile protection [109].

The role of T cells in the protection conferred by RAS immunization was originally established by results showing that whereas immunization of T-cell-deficient mice did not result in the generation of protective immunity, B-cell-deficient mice could still be protected following immunization with irradiated sporozoites [110]. Later, passive and adoptive transfer studies in mice further demonstrated that irradiated sporozoites induce potent cell-mediated immunity in the absence of antibodies [111, 112]. Selective depletion of T-cell subsets in RAS-immunized mice by treatment with monoclonal antibodies against CD4⁺ or CD8⁺ T cells subsequently showed that, whereas in vivo depletion of CD4⁺ T cells did not reduce immunity, depletion of CD8⁺ T cells completely reversed protection [50, 113]. CD8⁺ T cells were also shown to play an essential role in sterile immunity of *rhesus* monkeys following attenuated sporozoite vaccination [114]. The exact effector mechanisms utilized by the protective memory CD8⁺ T cells to eliminate liver-stage parasites are yet to be fully elucidated. Studies in mice immunized with *P. berghei* RAS have suggested that IFN γ produced in response to parasite-specific stimulation of CD8⁺ T cells is a critical effector molecule for protection [113, 115, 116]. However, whether protective mechanisms are indeed mediated by IFN γ produced by CD8⁺ T cells or by cytolytic perforin and granzyme through direct contact in the absence of IFN γ is not yet absolutely clear (reviewed in [100, 117, 118]). CD3⁺ $\gamma\delta$ T cells have also been implicated in RAS-elicited protective immunity against *P. yoelii* in the absence of $\alpha\beta$ T cells [119]. Besides, a role for interleukin (IL)-12, nitric oxide (NO), and NK cells for RAS-induced protective immunity initiated by CD8⁺ T cells has also been proposed [120, 121].

The pivotal role of CD8⁺ cytolytic T-cell-mediated protection following RAS vaccination of mice is now fully established, although the quantity or functionality of the protective CD8⁺ T-cell response can differ based on the rodent model [95]. In fact, although most studies suggest that effective RAS immunization in mice appears to depend on CD8⁺ T cells, in some strains of mice, protection mediated by CD8⁺ T cells is not absolute, in which case CD4⁺ T cells are also crucial [103]. Depletion of CD8⁺ T cells in some strains of mice does not affect protective immunity against

P. yoelii [122]. Moreover, *P. yoelii* and *P. berghei* RAS-induced sterile immunity has been observed in CD8⁺ T-cell-deficient mice, where protection appears to be mediated by antibodies and CD4⁺ T-cell-derived IFN γ [123].

Sporozoites and intrahepatic parasite stages contain over 1000 potential antigens [31]. This has led to alternative suggestions that the effectiveness of RAS immunization might result from broad immune responses against multiple antigen targets [124] or, instead, from a few immunodominant antigens that mediate the protection [52]. Several studies have pointed to CSP as a crucial determinant of T-cell responses following RAS immunization. The first evidence for a role for CSP in CD8⁺ T-cell-mediated immunity elicited by irradiated sporozoites came from passive transfer and in vitro studies that identified epitopes in the carboxyl terminus of the CSP of *P. berghei* [125] and *P. yoelii* [126], which are recognized by T cells from immunized mice. An epitope of *P. falciparum* CSP was later identified in studies with mice immunized with sporozoites or with recombinant vaccinia virus expressing *P. falciparum* CSP [127]. More recently, it was shown that in the absence of T-cell-dependent immune responses to CSP, protection induced by immunization with two doses of irradiated *P. yoelii* sporozoites was greatly reduced [52]. Another study has shown that protective immunity induced by *P. yoelii* RAS in transgenic mice expressing K^d molecules under the MHC-I promoter is mediated by CSP-specific, Kd-restricted CD8⁺ T cells [128], further supporting the notion that CSP is indeed an immunodominant protective antigen in mice. However, it has also been shown that, after RAS hyperimmunization, complete CD8⁺ T-cell-mediated protection can occur in transgenic mice that are T-cell tolerant to CSP and cannot produce antibodies [52]. In an effort to identify these non-CSP-protective T-cell antigens, the synthetic peptides corresponding to 34 *P. yoelii* sporozoite antigens that were predicted to contain strong CD8⁺ T-cell epitopes were screened for the presence of peptide-specific CD8⁺ T cells secreting IFN γ in splenocytes of the same hyperimmunized transgenic mice. This study revealed that the numbers of IFN γ -secreting splenocytes specific for the non-CSP antigen-derived peptides were 20–100 times lower than those specific for the CSP-specific peptide. Besides, when mice were immunized with recombinant adenoviruses expressing selected non-CSP antigens, the animals were not protected against challenge with *P. yoelii* sporozoites, despite the large numbers of CD8⁺-specific T cells generated [129]. While the immunodominance of CSP has been largely accepted, its role in immunity has been questioned by studies employing the expression of heterologous CSP proteins by *P. berghei* and *P. yoelii* sporozoites [130, 131]. Interestingly, adoptive transfer of CD4⁺ CTLs from mice immunized with *P. berghei* RAS that recognized an unidentified non-CSP sporozoite and blood-stage antigen conferred protection to naïve mice [132]. Nevertheless, immunization of mice with a peptide representing a CD4⁺ T-cell epitope in the N-terminus of *P. berghei* [133] or *P. yoelii* [134] CSP also conferred protection. It has also been argued that protective subdominant antigens may be expressed both by the sporozoite and by liver-stage parasites to allow recognition by CD8⁺ T cells primed in the lymph nodes by sporozoites, an expression that may be critical to induce protective CD8⁺ T cells [100]. In any event, the overwhelming evidence accumulated over the last few decades strongly suggests that CSP plays a pivotal role in the protection conferred by RAS immunization.

2.1.2 Early Studies of Antibody and T-Cell Responses in Humans

Ever since their identification, anti-CSP antibodies have been considered candidates for mediating protective immunity against malaria [67]. RAS immunization of human volunteers has been shown to elicit the production of antibodies to both the repeat and the flanking regions of CSP [61, 63], with antibody levels paralleling the serum inhibitory activity of sporozoite invasion of hepatoma cells in vitro [63]. Antibodies against the immunodominant B-cell epitope of *P. falciparum* CSP, identified as a repetitive (NANP)₃ sequence [69, 135, 136], inhibit sporozoite infectivity in vitro [70, 137] and in vivo [138]. However, the efficacy of a (NANP)₃ peptide-based vaccine was limited by the absence of *Plasmodium* T-cell epitopes [139, 140], in agreement with the notion that, in naturally exposed individuals, anti-CSP antibody levels seem to correlate with exposure but not necessarily with protection (reviewed in [141]). Antibodies against *P. falciparum* sporozoite surface protein 2 (SSP2) have also been identified in RAS-immunized volunteers protected against malaria [142].

CSP-specific cellular immune responses were described in RAS-immunized volunteers as determined by the proliferation of peripheral blood mononuclear cells (PBMCs) upon in vitro stimulation with recombinant *P. falciparum* CSP [61, 143]. An epitope mapping to the 5' repeat region of *P. falciparum* CSP was identified in T-cell lines and clones obtained from a sporozoite-immunized human volunteer [144]. Another epitope (peptide 326–345) of the *P. falciparum* CSP was later shown to be recognized by human cytolytic class II-restricted CD4⁺ T cells [145]. This epitope contains both a polymorphic [146] and a conserved [69] region of *P. falciparum* CSP, consistent with strain cross-reactivity of RAS-induced protection. A subsequent analysis of the genetic restriction of a series of T-cell clones derived from three RAS-immunized volunteers showed that peptide 326–345 can be presented by multiple MHC class II HLA-DR alleles [147]. Most of the clones analyzed in this study recognized a series of variant peptides representing the amino acid substitutions found in the polymorphic region of different *P. falciparum* strains, suggesting that polymorphism might not constitute a serious obstacle for recognition of this region of CSP by CD4⁺ T cells [147].

Despite the wide array of direct evidence regarding the contribution of *Plasmodium*-specific CD8⁺ T cells in protection in mouse models of RAS immunization, the inability to conduct CD8⁺ T-cell depletion studies in humans makes direct evidence of CD8⁺ T-cell involvement in immunity harder to obtain. However, as we will see below, liver-stage-specific cytokine-producing CD8⁺ T-cell responses are induced by RAS immunization in humans and may indeed contribute to protection.

2.1.3 RAS Immunization-Based Discovery of Preerythrocytic *P. falciparum* Antigens

Following the discovery that particular HLA class I and class II alleles were associated with resistance to severe malaria in Gambian children [148], an HLA-based approach termed reverse immunogenetics has been employed to identify epitopes

from preerythrocytic-stage antigens of *P. falciparum*. This approach enabled the identification of CTL epitopes for two common HLA-B alleles in the liver-stage antigen 1 (LSA-1) protein of *P. falciparum* [149, 150] as well as epitopes for several class I antigens that are found at high frequencies in Caucasians and/or Africans, including CSP, SSP2, and sporozoite threonine- and asparagine-rich protein (STARP) [149, 151].

Antigen discovery studies in naturally infected populations have been significantly complemented and extended by analyses performed in the context of RAS immunization. Proliferative T-cell responses to *P. falciparum* SSP2 have been reported in protected RAS-immunized volunteers suggesting that this antigen is a target of protective immunity [142]. HLA-B8-restricted responses against two epitopes of SSP2, one of which is conserved and the other variant among *P. falciparum* isolates, have also been reported in two *P. falciparum* RAS-immunized individuals [152]. Later, a panel of 27 open reading frames (ORFs) previously identified by a proteomic analysis [31], representing antigens potentially expressed in the sporozoite and intrahepatic stage of *P. falciparum* life cycle, was interrogated, leading to the identification of 16 antigenic proteins recognized by *P. falciparum* RAS-immunized volunteers [153]. Peptides derived from four previously well-characterized *P. falciparum* preerythrocytic-stage antigens [CSP, SSP2, LSA-1, and exported protein 1 (EXP-1)] were analyzed in parallel. Noteworthy, the reactivity against several of the newly identified antigens, most notably antigen 2 (later designated CelTOS—cell-traversal protein for ookinetes and sporozoites [154]), exceeded those observed against the well-characterized antigens [153]. In a subsequent study, protein microarrays representing 23% of the entire *P. falciparum* proteome were probed with plasma from individuals either showing sterile protection or not protected against infectious *P. falciparum* challenge, following experimental *P. falciparum* RAS immunization [155]. Overall, a markedly different pattern in antibody recognition was observed between the protected and non-protected groups of individuals, in agreement with previously obtained data [156]. A panel of 19 preerythrocytic-stage antigens, 16 of which were novel, was identified as strongly associated with RAS-induced protective immunity, which may form a potential signature associated with sporozoite-induced protection [155]. More recently, 27 out of 151 recombinant *P. falciparum* preerythrocytic antigens selected bioinformatically were recognized by plasma from RAS-immunized volunteers. Twenty-one of these antigens induced detectable antibody responses in mice and rabbits. Moreover, PBMCs from RAS-immunized subjects elicited positive ex vivo or cultured ELISpot responses against peptides from 20 of 22 of these antigens [157]. This study also showed that peptides spanning the full length of *P. falciparum* CelTOS recalled significantly higher positive responses from protected than non-protected subjects [157].

Collectively, these results suggested that immune responses to *Plasmodium* are complex and multispecific. These data further suggest that serodominance does not necessarily correlate with sporozoite-induced immunity and that some of the novel antigens identified by recently employed strategies may represent good candidates for vaccine development [153, 155].

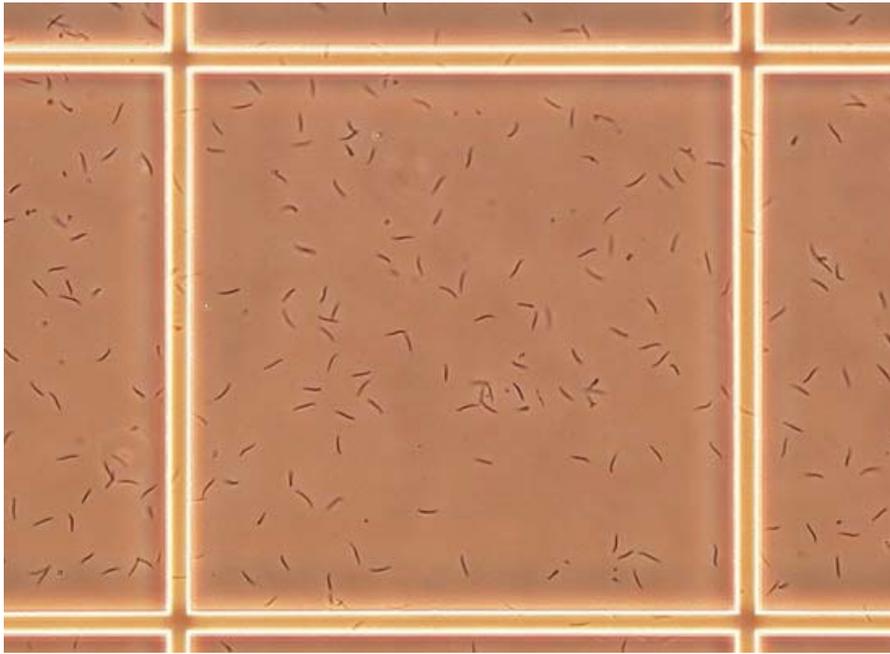


Fig. 2 Purified PfSPZ. The purification process significantly reduces the amount of salivary gland and mosquito material contaminating the PfSPZ (200× magnification). Photo kindly provided by Sanaria™ Inc.

2.2 Recent Progress in RAS Immunization

The twenty-first century witnessed a renewed interest in RAS immunization, which, to a significant extent, was due to the efforts of the company Sanaria™. This company has developed a radiation-attenuated, metabolically active, non-replicating *P. falciparum* sporozoite vaccine, the PfSPZ vaccine (Fig. 2) [46, 66]. The vialled PfSPZ vaccine is aseptic purified and cryopreserved in liquid nitrogen vapor phase (LNVP) [158], by an undisclosed method. Non-attenuated PfSPZ, cryopreserved by the same method, seems to be substantially less infective than their freshly dissected counterparts [159]. Nevertheless, they have been successfully employed in various CHMI studies where they have been administered by intradermal (id; [33, 34, 160]), intramuscular (im; [160–162]), and intravenous (iv; [35, 162]) routes, paving the way to PfSPZ vaccine immunization studies in humans.

2.2.1 The PfSPZ Vaccine Immunization and the Search for Immune Correlates of Protection

The first attempt at vaccination with the PfSPZ vaccine to humans was reported in 2011, when the vaccine was administered to 80 volunteers by id or subcutaneous (sc) injection, prior to challenge with infectious *P. falciparum* sporozoites [54]. Although the vaccine administered by these routes proved safe, it was only poorly

immunogenic and protective, with only two individuals protected out of 44 immunized volunteers challenged by the bites of five *P. falciparum*-infected mosquitoes. Immunized volunteers produced generally low levels of T-cell responses in response to a stimulus with PfSPZ, as measured by IFN γ production, which increased with vaccine dose and number of boosts for the sc but not for the id group of volunteers [54]. Furthermore, PBMCs from immunized volunteers showed a low frequency of PfSPZ-specific CD4⁺ T cells producing any combination of the IFN γ , tumor necrosis factor- α (TNF- α), and interleukin-2 (IL-2) cytokines and no CD8⁺ T-cell responses whatsoever [54]. Also of note, responses to whole sporozoites were more pronounced than to any of the individual *P. falciparum* proteins CSP, SSP2, LSA-1, or CelTOS. The titers of antibodies against whole *P. falciparum* sporozoites and *P. falciparum* CSP were found to increase with vaccine dose and number of doses, but immune sera revealed at best a modest capacity to inhibit *P. falciparum* sporozoite invasion in vitro [54]. These results prompted additional studies in nonhuman primate (NHP) and rabbit models to assess the impact of the route of immunization on protection. The results obtained strongly indicated that immunogenicity and protective efficacy could be greatly improved by iv administration of the PfSPZ vaccine. The suboptimal immunogenicity and protection elicited by vaccine administration by other routes may be explained by the reduced liver load that results from the latter, as was observed in mouse studies employing *P. berghei* [163, 164] and *P. yoelii* [164] sporozoites.

Collectively, the data obtained in this study argue in favor of iv vaccine administration to elicit sterile protection by RAS immunization. Indeed, this is the subject of a groundbreaking study in 2013, where various doses of the PfSPZ vaccine were administered iv to 40 volunteers, prior to challenge with fully infectious *P. falciparum* sporozoites [47]. The results of this study showed that the vaccine was protective, with six out of six and three out of nine volunteers immunized with five and four doses of 135,000 PfSPZ vaccine, respectively, fully protected against malaria. Humoral responses in protected and unprotected subjects were assessed 2 weeks after the last immunization by measuring antibody titers against CSP and PfSPZ, as well as by in vitro assays of inhibition of PfSPZ invasion (ISI). A dose-dependent increase in anti-CSP and anti-PfSPZ antibody titers, as well as in ISI, was observed. However, despite a clear trend for higher antibody titers and ISI in protected than in unprotected individuals, neither of these parameters could directly predict protection. Multiparameter flow cytometry was also employed to assess the frequency of PfSPZ-specific IFN γ , IL-2, TNF- α , or perforin-producing CD3⁺CD4⁺, CD3⁺CD8⁺, CD3⁺ $\gamma\delta$ T cells and NK cells from PBMCs 2 weeks after the final immunization. Although no absolute correlates of cell-mediated protection could be derived from this analysis, important information about the immune responses elicited by RAS immunization could be obtained: (1) a dose-dependent increase in frequency of PfSPZ-specific CD3⁺CD4⁺ T cells and CD3⁺CD8⁺ T cells producing any combination of IFN γ , IL-2, or TNF- α was observed; (2) there was no significant difference between protected and unprotected individuals for PfSPZ-specific CD3⁺CD4⁺ IFN γ -producing T cells, whereas protected subjects

showed a trend toward higher and more consistent PfSPZ-specific CD3⁺CD8⁺ IFN γ -producing T-cell responses; (3) no differences were observed between protected and unprotected subjects in terms of the quality of the PfSPZ-specific CD3⁺CD4⁺ T-cell responses, while a somewhat higher percentage of PfSPZ-specific CD3⁺CD8⁺ T cells that produced IFN γ only was observed in protected than in non-protected individuals; and (4) the overall frequency of CD3⁺ $\gamma\delta$ T cells increased in subjects who received the highest dose of vaccine. Finally, ELISpot analyses showed that IFN γ responses to PfSPZ were highest in the group that received the highest vaccine load, but IFN γ responses to CSP, SSP2, LSA-1, and CelTOS were low or undetectable in all subjects that received 135,000 PfSPZ vaccine [47].

More recently, another study assessed long-term protection of human volunteers following PfSPZ vaccination. In this report, 7 out of 9 and 6 out of 11 volunteers immunized with four doses of 270,000 iv-administered PfSPZ were protected following CHMI at 3 and 21 weeks after immunization, respectively. Five of the individuals protected at 21 weeks were re-challenged at 59 weeks after immunization and none developed parasitemia [165]. This study further showed that im injection of PfSPZ was substantially less efficient in inducing protection than iv administration of the vaccine, as only 3 out of 8 subjects vaccinated im with 2.2×10^6 PfSPZ were protected 3 weeks after vaccination, and all three developed parasitemia upon repeat CHMI at 25 weeks [165]. On average, anti-CSP and anti-PfSPZ antibody levels measured 2 weeks after the last immunization were significantly higher in protected than in non-protected individuals who underwent CHMI 3 and 21–25 weeks after the final immunization. Interestingly, the percentage of IFN γ -, IL-2- and/or TNF α -producing *P. falciparum*-specific CD4⁺ and CD8⁺ T cells in the blood did not correlate with outcome at the 3-week or the 21- to 25-week CHMI. However, the absolute frequency of unstimulated V δ 2⁺ T cells as a percentage of total lymphocytes correlated with outcome at the 21- to 25-week CHMI, while the frequency of the V δ 2⁺ T cell subset measured prior to the first immunization also correlated with outcome of CHMI at 3 and 21–25 weeks [165]. Nevertheless, it is important to note that neither the antibody nor the cellular correlates found in this study constitute absolute predictors of protection, as no clear threshold of antibody levels or V δ 2⁺ T cell frequency exists, above which all subjects are protected and below which none are.

Despite the encouraging results obtained with the PfSPZ vaccine, several challenges remain. These include the high numbers of parasites required for immunization, the requirement for iv delivery, the relatively high number of doses required for protection, the possibility that the vaccine might be strain specific, and the duration of protection after immunization [166, 167]. However, efforts to tackle these and other issues are currently being made by SanariaTM [168] and others. With five stage 1 clinical trials already completed, and an additional seven stage 2 trials in the USA, Europe, and Africa planned to address a set of well-defined objectives [59], significant progress in the clinical development of the PfSPZ vaccine can safely be expected in the near future.

3 Genetically Attenuated Parasites (GAPs)

3.1 Creation of GAPs

As discussed above, human test subjects immunized with *P. falciparum* RAS can be effectively protected from a subsequent challenge of fully infectious *P. falciparum* wild-type sporozoites. In 2005, employing rodent models of malaria, it was reported that similar sporozoite attenuation can be achieved by targeted gene deletion, by removing genes essential for liver-stage development, thereby creating so-called genetically attenuated parasites (GAPs) [40, 93]. Importantly, mice immunized with rodent *Plasmodium* GAPs can induce sterile immunity that is comparable to RAS immunization [169]. An advantage of GAPs over RAS immunization is that GAPs constitute a homogeneous population of parasites with a defined attenuation phenotype. This removes issues related to the delivery of correct doses of irradiation in order to ensure sufficient attenuation without killing the parasites or leaving them competent to produce a blood infection. The conceptual basis of vaccines consisting of GAP, just like RAS, is that after inoculation sporozoites invade but only partially develop in the liver. In the years following the first description of GAPs, a number of gene deletion mutants have been described in the rodent malaria models, *P. berghei* and *P. yoelii*, which are completely or partially attenuated, and manifest an arrest phenotype at different points during liver-stage development which provokes different levels of protective immunity (Fig. 3) [171].

The generation of GAPs, which can serve as a malaria vaccine, requires the creation of genetically engineered sporozoites that can invade liver cells but are unable to complete hepatic development. Furthermore, these parasites must provoke a strong protective immunity against a subsequent human malaria infection. These two features are critical to the development of a malaria vaccine consisting of attenuated parasites: first and foremost it must be safe (i.e., complete attenuation in the liver) and next it needs to be potent (generates strong protective immunity). *Plasmodium* genes that may govern exclusively sporozoite and/or liver-stage-specific functions were first identified by characterizing mRNA transcripts that are only expressed (or highly upregulated) in salivary gland sporozoites [30, 172]. Deletion of some of these gene candidates from the genomes of rodent malaria parasites resulted in the creation of some of the first GAPs [39, 93]. Moreover, as with irradiated sporozoites, immunization with sporozoites lacking expression of these proteins generated sterile protective immunity against malaria in mice. Independent studies, also in rodent models, investigating proteins of the *Plasmodium* 6-Cys protein family revealed that one member of this family, P52, was vital for parasite development in the liver and immunization of mice with sporozoites lacking P52 could also induce protective immunity [40]. Mutants lacking the 6-Cys protein P52 either by itself or in combination with the deletion of the gene encoding the closely related protein, P36, were among the first GAP to be created in *P. falciparum* [41, 55, 173]. Recently, a newly identified member of the 6-Cys protein family, B9, has

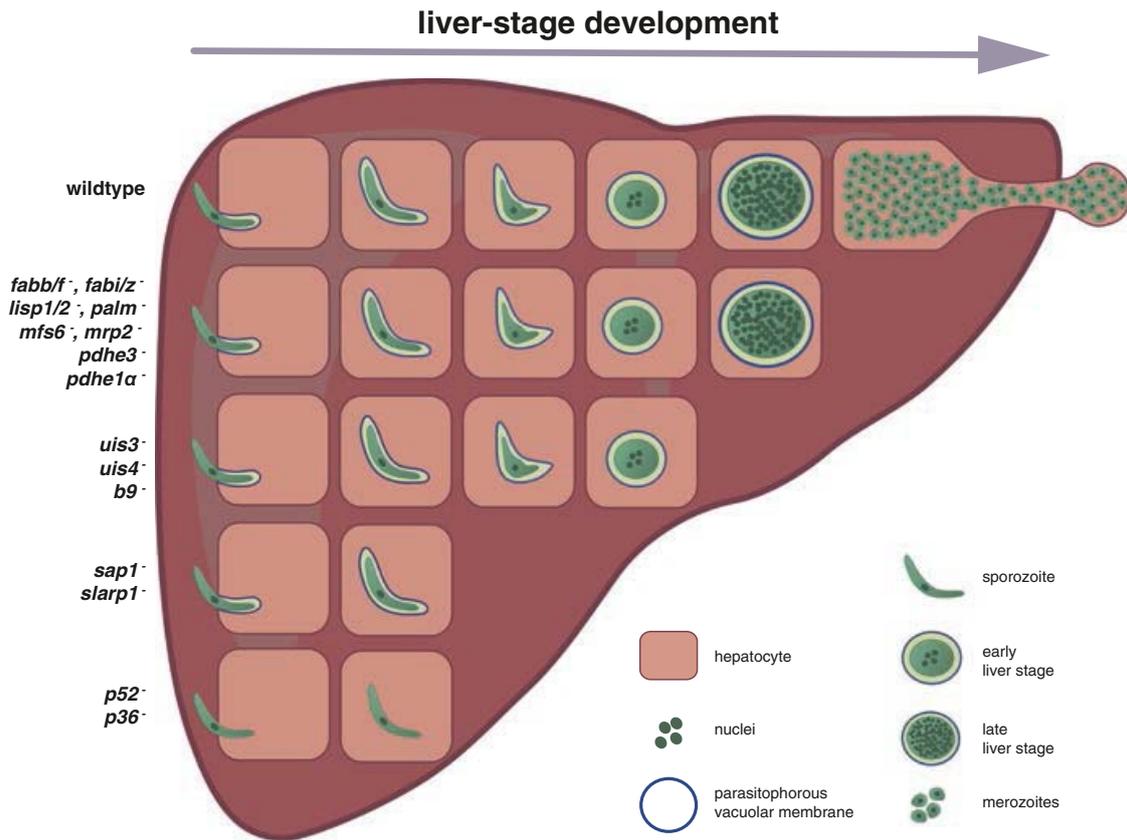


Fig. 3 Liver-stage developmental arrest of GAPs in rodent malaria models. Adapted from [170] and kindly provided by Britta Nyboer

also been shown to be critical to parasite development in the liver, and mutants lacking this protein have been shown to elicit strong protective immunity in rodent models [58, 174]. After the initial identification of these rodent GAPs, a number of other gene deletion mutants have been identified that are completely or partially attenuated and shown to induce protective immunity [171, 175] (Fig. 3). However, of all the GAP candidates in rodent models that could be advanced as a potential *P. falciparum* GAP vaccine, only mutants lacking the gene *slarp* (alternatively known as *sap1*), which encodes a protein involved in the regulation of transcripts in sporozoites, have been shown to fully arrest during liver-stage development, even at very high sporozoite doses (up to 5×10^6 sporozoites inoculated intravenously) [176–178]. Recently, GAPs generated in both *P. berghei* and *P. falciparum* lacking the gene *mrp2* that arrest mid-late during liver-stage development are only the second mutant that appears to be fully arrested in the liver cell even at very high sporozoite doses [179]. All other GAPs have been shown to be capable of leading to breakthrough blood infections when the dose of sporozoites is increased [171]. Consequently, the GAPs that are most advanced for human use and currently ready for clinical evaluation consist of mutants where *slarp/sap1* has been removed in addition to the deletion of one or more of the 6-Cys genes from the *P. falciparum* genome. Specifically, the *P. falciparum* GAPs “*p52(-)/p36(-)/sap1(-)*” and “ $\Delta b9\Delta slarp$ ” have both been examined in blood-stage culture, in mosquitoes, in cultured human hepatocytes, and in mice engrafted with human liver tissue and have

been shown to only exhibit a growth defect after liver cell invasion [57, 58]. They completely abort intrahepatic development, demonstrating that they are likely to be safe for use in humans; currently these GAPs await clinical testing to evaluate their protective efficacy in humans. Importantly, these *P. falciparum* GAPs are expected to arrest early in hepatic development, soon after invasion of hepatocytes by sporozoites.

It has been reported that a late-arresting rodent GAP was a superior immunogen to irradiated sporozoites and early-arresting GAPs [53]. GAPs that progress into late liver-stage development will express both a greater quantity and a greater diversity of antigens, compared to early-arresting GAPs and RAS. The hypothesis is that parasites that arrest late into liver-stage development will not only expose the immune system to a greater magnitude of antigens (a mature liver stage contains up to 40,000 times more protein than an invading sporozoite) but also to a greater diversity of antigens, as they will express antigens expressed in irradiated sporozoites as well as antigens expressed later in the liver and in blood stages (Fig. 3).

3.2 Immunity Induced After GAP Immunization

Depending on the type and/or strength of attenuation, parasites arrest at different time points during liver-stage development or shortly thereafter. Heat-killed or over-irradiated sporozoites fail to invade hepatocytes and hence do not confer protection [180]. This emphasizes the importance of the preerythrocytic stage and the requirement for metabolically active, attenuated parasites that progress at least to early hepatic development before they arrest [181]. Sterile protective immunity in whole-sporozoite experimental vaccinations highly depends on the parasite species, mouse strains, and the immunization regimens. As such, sterile protection is more easily achieved in the *P. yoelii*/Balb/c experimental model than in the *P. berghei*/C57/Bl6 model [121, 182].

Protective immunity has been reported to rely on CD8⁺ T cells and IFN γ as the key mediators of protection in all different GAP models [169, 183, 184]. Depletion of CD4⁺ T cells did not abrogate protection, although adoptive transfer of CD4⁺ T cells from, e.g., *uis3(-)*-GAP-immunized mice protected 50% of naïve animals from wild-type infection, indicating a role also for this T-cell subset in GAP-induced immunity. Humoral responses to CSP were measured in the *P. berghei* GAP model but appeared to be dispensable since B-cell-deficient mice were capable of sterile protection [183].

Similarly to RAS, GAP immunization induces a significant increase in intrahepatic CD8⁺ T cells displaying a T effector memory phenotype (Tem) that is clearly associated with long-term protection [185, 186]. Concerning direct effector mechanisms, Trimmell et al. hypothesized that contact-dependent cytotoxicity is mediated by CD8⁺ T cells since sterile protection was abolished in perforin-deficient mice. Moreover, CD8⁺ T cells induced apoptosis in infected hepatocytes in vitro [169, 186].

One dose of *P. yoelii uis4(-)* sporozoites was sufficient to induce protection against challenge infection in a Balb/c mouse background [184] that is mediated by the significant upregulation of specific effector CD8⁺ T-cell activation markers such as CD11c. It was suggested that the upregulated CD11c expression phenotype on effector CD8⁺ T cells induced by *P. yoelii* GAP vaccination is not only responsible for their potent inflammatory function but also marks their highly proliferative but short-lived, antigen-specific effector activity, which will ultimately lead to controlling both infectious sporozoite hepatocyte invasion and intrahepatic differentiation [187].

A second generation of GAP leading to a late arrest in liver-stage development was introduced more recently. Immunization with *Pyfabbf(-)* parasites that are depleted of a gene encoding an essential enzyme of the apicoplast-localized, type II fatty acid biosynthesis (FAS II) pathway induced long-lasting protection in rodent models [53, 188], despite later evidence of breakthrough infections by *Pbfabbf(-)* parasites [55]. The results of the underlying study further suggested that a larger number of effector memory CD8⁺ T cells with a broader antigenic specificity were elicited by immunization with *Pyfabbf(-)* GAPs as compared to early-arresting GAP or even RAS. These characteristics of *Pyfabbf(-)*-induced immune responses correlated with enhanced protection, and thus it was postulated that late-arresting GAP confers superior protective immunity than early-arresting attenuated parasites. Besides, it was shown that this superior protection is stage transcending, i.e., mice immunized with *Pyfabbf(-)* are not only sterile protected against sporozoite-induced challenge infection but also enjoy protective immunity after intravenous challenge with homologous *P. yoelii* blood-stage parasites [53].

As such late-arresting live attenuated parasites are promising for next-generation vaccine design, it is now imperative to gain a more detailed overview about the underlying immunological functions triggered during experimental vaccination, before such attenuated parasites can be optimized for human clinical trials. Protective immunity induced by the late-arresting experimental whole-organism model parasite *Pyfabbf(-)* was comprehensively studied over the last few years. It was recently found that not only CD8⁺ T cells but also CD4⁺ T cells contribute to protective responses and that the co-stimulatory molecule CD40 is key in mediating robust protection. CD40 signaling was demonstrated to control intrahepatic DC licensing during *Pyfabbf(-)* immunization and seems to be instrumental for establishing robust CD8⁺ effector and memory T-cell responses [189]. Very recently, the immunity mechanisms underlying stage-transcending protection exclusively induced by late-arresting *Pyfabbf(-)* GAP [53] were reported to involve both the cellular and humoral arm of the immune system. Hence, it is proposed that during experimental vaccination with late-arresting GAPs, the immune system gets equipped to establish immunity against both the clinically silent, intrahepatic and pathological, blood-stage phase. As for the humoral response, it is thought that antigenic targets are shared between the hepatic and blood-stage phase and protective antibodies rely on F_c-mediated functions such as F_c receptor binding and complement fixation [190]. Indeed, a study by Keitany et al. showed for the first time that functional antibodies are induced in *Pyfabbf(-)* GAP-immunized mice and inhibit

liver cell invasion as well as the capacity to decrease intrahepatic parasite burden [191]. It is a matter of course that such GAPs not only inform about the underlying immunological mechanisms involved in protection against parasites but will also be ideal to further isolate novel antigens and/or correlates of (transcending) sterile protection. The latter will ideally assist in the development of novel subunit-anti-infective vaccination strategies. Interestingly, a study headed by Chen et al. reported the identification of *PyTm21* as a promising protective antigen candidate that shows the capacity to reduce intrahepatic parasite burden and contribute to sterile protection during *Pyfabbf(-)* GAP experimental immunization [192, 193].

Experimental immunization of rodents with a knockout parasite line for the *Plasmodium*-specific apicoplast protein important for liver merozoite formation (PALM) elicits potent and long-lived protective immunity against sporozoite-induced malaria by only two immunization doses. Indeed *palm(-)* knockout parasites are very similar to the *Plasmodium* late-arresting *fabbf(-)* strain, as PALM seems to be important for the successful transition from liver to blood-stage parasites, i.e., display a growth arrest at a very late time point in intrahepatic development [194]. Indeed, a thorough study of the underlying immunity mechanisms of *palm(-)*-vaccinated rodents revealed persistence of some intrahepatic antigen-specific IFN γ -secreting CD8⁺ memory T cells in aging immunized mice [195, 196].

The involvement of innate immunity in GAP-mediated immune responses still remains to be investigated. GAP immunization of *RAG1 (-/-)* mice that lack adaptive immune cells (which are traditionally considered to mediate memory responses to vaccinations) resulted in a delayed disease outbreak after challenge with viable sporozoites, suggesting that the innate immune system in these mice is sufficient to confer partial protection after vaccination, e.g., through NK memory cells [183]. However, the cellular events and molecular pathways underlying the process of memory generation in NK cells remain largely elusive.

Regarding antigenic specificity, only CSP-specific, IFN γ -secreting CD8⁺ T cells, as well as CSP-specific antibodies, were reported to be elicited by GAP immunizations. Indeed, Kumar et al. measured no significant differences in these CSP-specific responses in Balb/c mice that were either immunized with *PyRAS* or *Pyuis3(-)* and *Pyuis4(-)* genetically attenuated parasites. The authors therefore postulated conserved and similar mechanisms of protection irrespective of the method of attenuation [183, 197].

Recently, the results from a first clinical trial demonstrated safety of *PfGAP* vaccination in humans using *Pfp52(-)/p36(-)* double knockout parasites. Initial immunological profiling showed CSP-specific antibody titers after 200-bite exposure of volunteers and moderate memory responses of CD8⁺ and CD4⁺ T cells to CSP and non-CSP antigens such as LSA-1 or AMA-1. Of importance, a breakthrough infection was observed in one of the *PfGAP*-immunized volunteers suggesting that development of more severely attenuated *PfGAP*, possibly with multiple gene deletions, should be pursued in future developments [42].

It has been demonstrated that immunization of mice with late liver-stage-arresting GAP elicits broader and better CD8⁺ T-cell responses when compared to irradiated (early-arresting) sporozoites, producing superior long-lasting sterile protection

against a malaria infection [53]. These parasites are believed to present a greater diversity and amount of parasite antigen to the host immune system. However, *P. falciparum* parasites lacking FabB/F expression fail to produce sporozoites [198]. Thus, discrepancies in attenuation phenotypes and potency between GAPs of different *Plasmodium* species indicate that the importance of the same gene for liver-stage development may differ between rodent models and *P. falciparum*. Consequently, the identification of *Plasmodium* genes that are critical to mid-/late liver-stage development to create late-arresting *P. falciparum* GAPs is actively being sought, and a number of gene expression studies on rodent and human sporozoite and/or intrahepatic liver stages are currently being mined to identify these genes.

4 Chemoprophylaxis and Sporozoite (CPS) Immunization

While sterile protection has been most elaborately investigated using RAS, induction of protective preerythrocytic immunity is most potent and efficient when sporozoite and liver-stage development remain fully intact, while asexual-stage multiplication is abrogated at the earliest stages [59, 199, 200]. Attempts with targeted deletions of genes encoding late-stage liver antigens are promising, but, as outlined above, safety has thus far not been appropriately ensured in humans because of the occurrence of breakthrough infections with asexual-stage parasites [42, 175, 199]. Successes, however, have been reported with the use of antimalarial drugs in combination with wild-type parasites: chemoprophylaxis and sporozoites (CPS) immunization (or infection-treatment-vaccination (IVT)) can safely induce fully protective immune responses as was first demonstrated in rodents [37, 38, 43, 201]. The first drug used in these *P. berghei* studies to kill parasites at early blood-stage level was chloroquine, which disrupts heme detoxification and does not affect preerythrocytic stages [202, 203]. Other antimalarials have since been successfully used in rodents, including artesunate, azithromycin, clindamycin, mefloquine, piperazine, primaquine, and pyrimethamine with variable effects on parasite liver-stage development [204–208]. Similar results have been obtained in the CHMI model where malaria-naïve adult volunteers receiving mosquito bites from a total of 36–45 *P. falciparum*-infected mosquito under chloroquine prophylaxis (Fig. 4a) show complete protection against a homologous challenge 3–5 months after the last immunization and (partial) protection up to 2.5 years later [44, 45, 56]. Chloroquine and mefloquine are equipotent as prophylactic drugs [209], and other partner drugs are under investigation in ongoing trials [59]. As in murine models [43, 210, 211], induction of sterile CPS-induced protection is dose dependent (Fig. 4b) [209]. Both partial and full protection against a heterologous *P. falciparum* strain after CPS immunization have been demonstrated in a rechallenge trial [212] and are under further investigation (clinicaltrials.gov NCT02098590).

The potency of the CPS-chloroquine regimen is unprecedented compared to radiation-attenuated sporozoite (RAS) immunization and to protection obtained

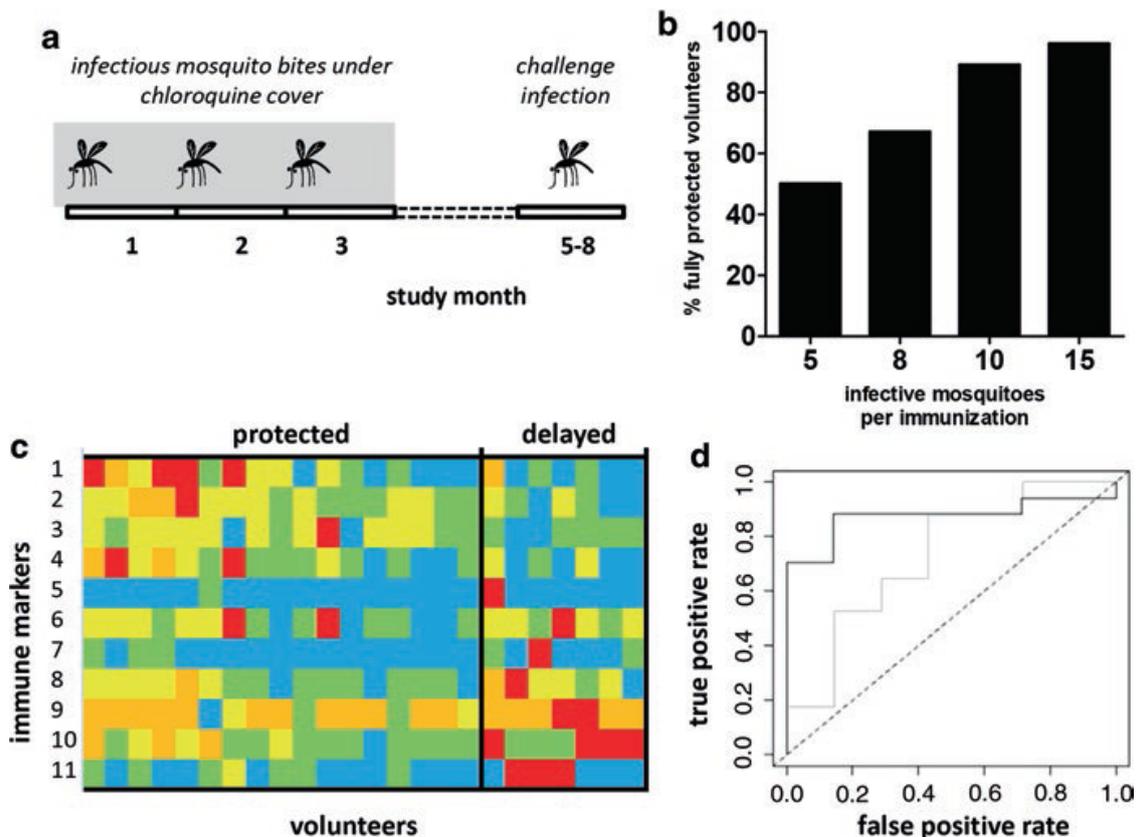


Fig. 4 CPS immunization regime, protection, and cellular immune signature in human volunteers. (a) Malaria-naïve adult volunteers are subjected monthly to bites of *P. falciparum*-infected mosquitoes while taking chemoprophylaxis. Protection is assessed by challenge infection via infected mosquitoes 3–5 months after the last immunization. (b) Sterile protection defined as the absence of parasitemia by microscopy until 21 days after challenge infection is dose dependent as assessed for $n=58$ volunteers enrolled in four clinical trials using either chloroquine or mefloquine prophylaxis [44, 56, 209]. (c) Rows show eleven biomarkers that can distinguish $n=17$ fully protected and $n=7$ partially protected/delayed volunteers in a dose escalation CPS trial [209], identified by penalized logistic regression. Colors indicate the degree of responses, with strongest responses in red. (d) Accuracy of prediction of protection from mosquito bite challenge assessed with the area under the receiver operator curve based on leave-one-out cross validation (LOOCV), using one parameter alone (parasite-specific degranulating CD107a⁺ CD4⁺ T cells, light gray line, AUC 0.73) or in combination with ten additional markers (AUC 0.87, dark gray line) (unpublished data)

after natural infections [59, 213]. While the dogma that protective malaria immunity is difficult to acquire is challenging, possible explanations including the broader antigenic repertoire, higher antigen load, and potential immunomodulatory effects of chloroquine remain speculative [200]. The CPS model, however, opens avenues for a better understanding of immune mechanisms of protective immunity, identification of immune correlates of protection, and novel target antigens for inclusion in subunit vaccines. Moreover, clinical development of a vaccine based on the CPS regime is currently explored with cryopreserved sporozoites as PfSPZ-CVac (clinicaltrials.gov NCT02511054 and NCT02115516 and [59, 214]).

4.1 Parasite Life Cycle Target for CPS-Induced Protection

The mechanism of CPS-induced immunity appears to be determined by specific parasite/host combinations regarding both the immune effector mechanisms involved and the parasite life cycle stage targeted. Most evidence points to preerythrocytic stages as targets for protective immunity: (1) human CPS-immunized volunteers are protected from challenge by mosquito bite, but not after a blood-stage challenge [56], and (2) CPS immunization of C57/Bl6 mice with *P. berghei* or Balb/c mice with *P. yoelii* with 10,000 or more sporozoites intravenously [37, 43, 203, 207, 208, 215, 216] results in preerythrocytic immunity and reduction of liver parasite burden [43]. Intravenous immunization with high numbers of *P. chabaudi* sporozoites also induces preerythrocytic immunity, while presumably low numbers of sporozoites as delivered by mosquito bites primarily induce immune responses against blood stages in this model [211]. Blood-stage immunity induced by CPS immunization has been reported also in the *P. yoelii* model [208, 215], where control of parasite growth was clearly antibody dependent [208, 215]. More strikingly, there is evidence that exposure to blood stages only can also convey preerythrocytic immunity ([201, 211] and Nganou-Makamdop et al. unpublished data). The combined data suggest that antigens shared between liver and blood stages [17] create the possibility for the existence of stage-transcending protection [217].

Since clinical follow-up of prolonged parasitemia in the CHMI model is limited for obvious safety reasons, CPS studies in humans primarily highlight induction of liver-stage immunity [56]. There are nevertheless hints for the presence not only of immune recognition but also of blood-stage immunity: (1) although humoral responses directed against blood-stage antigens are minimal ([44, 218] and Behet et al. unpublished data) and show no growth inhibition activity in vitro [56], IFN γ responses to blood-stage parasites are enhanced both in vivo [56] and in vitro [44, 45, 209, 219], and (2) immunized volunteers show a somewhat reduced peak parasitemia after blood-stage challenge [56]. These data feed the possibility that at least some functional blood-stage immunity might be present which only becomes apparent after a number of asexual multiplication cycles required for boosting. This hypothesis is in line with a correlation between prepatent period, as determined by thick smear, and the interval between a positive qPCR and positive thick smear as a proxy for parasite growth rate (Bijker et al. unpublished data), in partially protected volunteers (24). Since prepatency itself was not dependent on parasite density at the time of positive qPCR, these data suggest that parasite growth becomes slower when parasitemia is longer lasting. Thus, the CPS model efficiently induces sterile protection that is, in principle, accomplished at the preerythrocytic level. In addition, a blood-stage component may be present, which may contribute in case of partial protection and might become more evident in individuals possessing some degree of anti-disease immunity that allows longer follow-up. Albeit insufficient by itself, the short period of low-grade parasitemia as obtained under chloroquine prophylaxis in humans might be an asset for induction of at least some contributing blood-stage immunity. Indeed, low parasitemia has been shown before to efficiently induce blood-stage protection in the CHMI model [220].

4.2 Humoral Immunity After CPS Immunization

Parasite-specific antibodies induced by CPS immunization may interfere at several levels with preerythrocytic parasite development where sporozoite integrity, motility (gliding), hepatocyte invasion, and/or intrahepatic development may be compromised. In the *P. berghei* murine model, the functional contribution of antibodies to CPS-induced preerythrocytic protection is unclear: in CPS-immunized C57/Bl6 mice, antibodies induced against sporozoites and liver stages inhibit hepatocyte infection in vitro [208], but are unable to confer protection against sporozoite challenge in vivo [43, 208]. In human volunteers, CPS immunization induces both a robust plasma antibody and long-lasting memory B-cell response [218]. This humoral response is largely directed against preerythrocytic parasite antigens with CSP being the major target identified thus far [44, 209, 218, 221]. Plasma IgG from CPS-immunized volunteers inhibits sporozoite traversal in vitro, as well as liver-stage infection in human liver-chimeric mice following challenge by *P. falciparum*-infected mosquito bites [222]. However, anti-CSP titers per se appear to be a correlate of exposure rather than protection in the human CPS model [218]. In fact, anti-CSP titers are the lowest in individuals that become already protected during the immunization period [218]. Since CSP is not only expressed in sporozoites but also in liver stages [32, 223, 224], one possible explanation might be that at least a proportion of the parasites is eliminated by CPS-induced immune responses early during liver-stage development. It remains to be investigated to what level human anti-CSP antibodies will be sufficient to prevent sporozoite infectivity. There is evidence that targets other than CSP might be involved to accomplish sterile protection [52, 130]. The use of protein microarrays probing the antibody compartment for proteome-wide antigen reactivity [156, 225] has already revealed novel protection-associated antigen targets in RAS-immunized individuals [155], and similar investigations are currently ongoing in CPS-immunized volunteers.

4.3 Cellular Immunity After CPS Immunization

As in other whole-sporozoite vaccination approaches [95], cellular responses induced after CPS immunization are considered to be responsible for the key effector mechanisms driving elimination of infected hepatocytes. In the C57/Bl6 *P. berghei* CPS model, there is a sustained CD8⁺ T-cell IFN γ response [205, 216] correlating with protective efficacy [216], and immunity can be transferred by total splenocytes [203] and CD4⁺ and CD8⁺ T cells [43]. Despite a strong CD8⁺ T-cell response, protection of *P. yoelii* CPS-immunized C57/Bl6 mice, in contrast, was independent of CD8⁺ T cells, suggesting a role for CD4⁺ T cells or antibodies instead. CPS immunization in humans induces long-lasting cellular immune responses [44, 45, 209, 219], which are still detectable even 2.5 years after the initial immunization [45]. In line with a role for both Th1 and cytotoxic responses, CPS-immunized volunteers show increased expression of granzyme B and the Th1

transcription factor T-bet in activated and proliferating T cells during immunization [214]. However, a clear immune correlate of protection has not yet been identified, as neither polyfunctional effector memory T cells that coproduce IFN γ and interleukin-2 [44, 45] nor total IFN γ -producing CD4⁺, CD8⁺, or $\gamma\delta$ T cells that can differentiate between fully and partially protected volunteers [209]. Parasite-specific degranulation by CD4⁺ T cells and granzyme B production by CD8⁺ T cells, in contrast, are significantly higher in fully protected individuals [209]. Nevertheless, these cytotoxic responses also do not allow for a distinct differentiation between the two groups. Given the fact that individuals differ in their “immunological fingerprint” and protection likely relies on the interaction of multiple immune effector pathways, a combination of multiple immune markers in the form of an immune signature may be of more predictive of protection than individual responses (Bijker and Teirlinck et al. unpublished data, Fig. 4c, d). Next to adaptive responses, this may also include innate lymphocytes such as NK cells but particularly $\gamma\delta$ T cells, which are expanded following CHMI, CPS, and high-dose RAS [47, 219, 226] and show memory-like responses following CPS immunization [209, 219].

Based on observations from natural immunity in the field, the current dogma is that acquisition of protective immunity against malaria per se is a lengthy and difficult process involving many exposures and is rapidly lost in the absence of exposure. The efficient induction and persistence of immunity following CPS immunization, however, clearly refutes this statement and indicates that this may only apply to protection from disease, but not infection. A plausible explanation can be found in the composition of immune responses targeting the different parts of the life cycle involving distinct parasite immune regulation and evasion strategies [200]. CPS immunization induces protection from sporozoite challenge, in line with predominantly preerythrocytic targeted antibodies that reduce sporozoite infectivity and long-lasting cellular cytokine and cytotoxic responses. None of these responses, however, are a clear-cut correlate of protection, and further research is necessary to identify which (combination) of immune responses predicts protection. Importantly, while this immunization regimen has been primarily tested against homologous parasites, heterologous challenge studies are underway and indicate at least some protection. Future studies will need to focus on the identification of target antigens of the induced immune responses and their functionality also against heterologous parasites. Another focus should be to elucidate the potential contribution of blood-stage immunity in this model, particularly in preexposed individuals. Thereby, CPS immunization may not only aid future subunit vaccine development but—once satisfactory—also be considered for clinical development as a vaccine to be marketed for special groups including travelers or the military.

5 Other Models of Sporozoite-Based Vaccination

RAS, GAP, and CPS are, by far, the most thoroughly investigated approaches for whole-sporozoite vaccination against malaria. However, alternative strategies for eliciting protective immune responses against human-infective *Plasmodium* parasites have also been proposed and will be briefly addressed here.

5.1 Chemically Attenuated Sporozoites

Chemical attenuation of sporozoites (CAS) has been investigated as an alternative strategy for the production of a liver-stage vaccine against malaria. Similarly to other whole-sporozoite immunization approaches, the concept of CAS relies on the immune responses elicited during hepatic infection by chemically treated sporozoites, which are unable to progress into a patent blood-stage infection and cause disease. The strategy has been evaluated by the use of centamycin [227], a DNA-binding agent that is effective against blood-stage *Plasmodium* parasites and blocks sexual differentiation of the parasites in mosquitoes [228]. Centamycin-treated *P. berghei* sporozoites were shown to display a significantly reduced hepatocyte infection rate and impaired preerythrocytic development and to be unable to generate blood-stage infections in mice [229]. Immunization of mice with *P. berghei* [229] and *P. yoelii* [229] CAS was shown to produce sterile immunity against challenge with fully infectious sporozoites. Vaccination with *P. yoelii* or *P. berghei* CAS induced sporozoite-specific antibodies and CSP-specific IFN γ -producing CD8⁺ T-cell responses at levels similar to those generated by RAS immunization [229]. More recently, it was shown that chemical attenuation of *P. berghei* parasites with isopentaquine during their liver-stage development alters disease outcome and protects mice from experimental cerebral malaria [230]. It has been argued that the CAS vaccine presents certain advantages over the RAS approach, most notably the fact that the chemical attenuation process can be strictly controlled, leading to a vaccine that is reproducibly attenuated and does not present the risks potentially associated with improper parasite irradiation [229]. However, further investigations are clearly necessary before the true potential of CAS vaccination can be fully ascertained.

5.2 Rodent *Plasmodium*-Based Immunization Against Human Malaria

A yet unpublished novel approach for whole-sporozoite vaccination against malaria has recently been proposed, based on the use of genetically modified rodent *Plasmodium* parasites that were genetically engineered to express and present antigens of their human-infective counterparts. The concept relies on the notion that such genetically modified rodent parasites may elicit cross-species immune responses arising from similarities between the antigenic compositions of the two species of parasites (interestingly, immunization with *P. falciparum* CelTOS has previously been shown to confer protection against *P. berghei* [231]), along with antigen-targeted responses resulting from the introduction of specific human *Plasmodium* immunogens on the rodent parasite platform. The proof of concept of this idea has been established through the use of *P. berghei*, a rodent parasite species that is widely recognized as nonpathogenic to humans. *P. berghei* is readily

amenable to genetic modification [232], and strategies to generate transgenic parasites expressing human-infective *Plasmodium* proteins have been developed [233]. A method for transgene expression and gene complementation [234] was employed to generate selectable marker-free *P. berghei* parasites that express *P. falciparum* CSP under the control of the *P. berghei* upregulated in sporozoites 4 (UIS4) promoter [*Pb(PfCS@UIS4)*]. *P. berghei* was shown to effectively infect human hepatocytes of human liver-chimeric mice, a requisite for any whole-sporozoite vaccine against malaria. Immunization of mice with *Pb(PfCS@UIS4)* was shown to elicit the production of antibodies against whole *P. falciparum* sporozoites as well as against *P. falciparum* CSP. Passive transfer of the immune serum of these mice into liver-humanized mice followed by challenge by *P. falciparum*-infected mosquito bites led to a significant blockage of hepatic infection by this human-infective parasite. Following completion of its preclinical validation, the *Pb(PfCS@UIS4)* vaccine candidate is expected to enter Phase I/IIa clinical trials. If successful, the use of “naturally attenuated” rodent parasites for immunization against human malaria could gain further momentum, and next-generation vaccine candidates can be envisaged to protect not only against multiple species of human malaria but also against various stages of the parasite’s life cycle.

6 Future Directions and Concluding Remarks

The application of a whole-sporozoite vaccination approach has raised various types of concerns, ranging from safety to deployment in the field. Safety-wise, whole-sporozoite immunization requires ensuring that sporozoites cannot complete their development process in the liver or that merozoites are eliminated as soon as they reach the blood while retaining immunogenicity [60, 235]. On the other hand, the *P. falciparum* sporozoites required for immunization must be obtained from infected mosquitoes, which must be reared under appropriate containment conditions to prevent accidental release. Sporozoites are then obtained from the dissection of mosquito salivary glands and must subsequently be purified and adequately cryopreserved for long-term storage [46]. Finally, application to the field requires knowledge of total dose required for protection and timing of booster immunizations [95], as well as a feasible vaccine administration method that ensures the development of protective immune responses.

Many important aspects to be considered for the design of a next-generation preerythrocytic vaccine may equally account for the development of a blood-stage vaccine. The ultimate question to be answered is what kind of immune response must be induced to confer high-level and long-lived protection and how to accomplish this. This task is both supported and exacerbated by the complex and not always unequivocal data from basic research, preclinical, and clinical studies. In general, it will be vital to improve the knowledge about protective immune

responses, i.e., to properly understand the difference between vaccine-induced protection and naturally acquired immunity, to exploit the data on *P. falciparum* RAS-mediated protection in humans or the large past clinical trials of RTS,S in order to determine which responses are the true mediators of protection.

A substantial amount of data advocates the need to induce antibody response against the sporozoite to prevent liver infection or at least to reduce the number of sporozoites infecting hepatocytes. The latter mechanism is assumed to contribute to a delay in prepatency in RTS,S-vaccinated individuals which consequently limits severe disease. Likewise, it seems unrealistic that complete protection can be mediated solely by humoral responses that were even shown to be dispensable in animal models. In fact, numerous studies have pointed out the essential role of T-cell responses against liver-stage parasites and have explicitly demonstrated the critical function of intrahepatic CD8⁺ T cells in protective immunity. This has important implications for the quality and/or quantity of the CD8⁺ T-cell response but also for the liver as the site of T-cell priming and effector function.

It seems clear that research into the immune responses induced by whole-sporozoite immunization approaches has given first insights into the immune mechanisms involved, but an actual correlate of sterile protective immunity has yet to be revealed. Such a correlate is unlikely to be a single common readout, but instead a signature of multiple immune effector pathways acting in synergy. This will likely include both phenotypic and functional readouts of humoral and cellular immune responses as well as a combination of various antigen targets. Complementing ongoing efforts using the protein microarray for genome-wide screening of antimalarial antibody responses, T cells which target antigen can also be identified by large-scale screen approaches, such as the AtlasTM assay using protein plasmid and expression libraries [225]. Application of this screening technology to samples from RAS- as well as CPS-immunized individuals is currently under way [225] and should be extended to GAP trials in the future. Such comparative analysis will be highly informative to dissect the differences and overlaps in immune signatures between the three main whole-sporozoite immunization approaches to (1) clarify why some approaches are more efficient in inducing protection than others and (2) home in on a minimal signature of antigen targets associated with protection. Target antigen identification will be useful not only for subunit vaccine design but also for generation of next-generation whole-sporozoite immunization regimens using parasites transgenic for variants of key antigens to induce efficient heterologous protection.

It can safely be said that further investigations of the immunological mechanism(s) whereby whole-sporozoite inoculation elicits protective immunity will greatly facilitate whole-sporozoite vaccine development [167]. Such knowledge, along with the expanded use of controlled human malaria infection studies [236, 237] and the envisioned progress in tackling the technical and logistical hurdles that still persist [59], can be expected to lead toward the ultimate goal of obtaining an effective vaccine with a pivotal role in the eventual global eradication of malaria.

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