

PERSPECTIVE

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Integrating malaria vaccine and CRISPR/Cas9 gene drive: a comprehensive strategy for accelerated malaria eradication

Israel Charles Abraham¹, John Ehi Aboje², Bonaventure Michael Ukoaka³, Kehinde Tom-Ayegunle⁴, Maryam Amjad⁵, Anas Abdulkader⁶, Chinonyelum Emmanuel Agbo⁷, Oluwatosin Ayokunle Akinruli⁸, Taiwo Rebecca Akisanmi⁹, Emmanuel Oyedeji Oyetola¹⁰, Gbolahan Olatunji¹, Emmanuel Kokori¹ and Nicholas Aderinto^{10*}

Abstract

Malaria remains a significant public health challenge, particularly in low- and middle-income countries, despite ongoing efforts to eradicate the disease. Recent advancements, including the rollout of malaria vaccines, such as RTS,S/AS01 and R21/Matrix-M™, offer new avenues for prevention. However, the rise of resistance to anti-malarial medications necessitates innovative strategies. This review explores the potential integration of CRISPR/Cas9 gene drive technology with malaria vaccination efforts to enhance vector control and reduce transmission. By employing gene drive mechanisms for population suppression and replacement of malaria-transmitting *Anopheles* mosquitoes, combined with the immunogenic properties of vaccines, a synergistic approach can be established. This paper discussed the need for integrated strategies to address the biological complexities of malaria and socio-economic factors influencing its prevalence. Challenges such as regulatory hurdles, community acceptance, ecological impacts, and sustainable funding are examined, alongside strategies for implementation within existing malaria control programmes. This integrated approach could significantly contribute to achieving the World Health Organization's targets for malaria reduction by 2030, ultimately enhancing public health outcomes and supporting broader socio-economic development.

Keywords Malaria, Vaccine, CRISPR/Cas9, Gene drive technology

*Correspondence:

Nicholas Aderinto
nicholasoluwayeji@gmail.com

¹ Department of Medicine and Surgery, University of Ilorin, Ilorin, Nigeria

² Benue State University, College of Health Sciences, Makurdi, Nigeria

³ Department of Internal Medicine, Asokoro District Hospital, Abuja, Nigeria

⁴ Dept of Epidemiology & Biostatistics, Johns Hopkins Bloomberg School of Public Health, Maryland, USA

⁵ Liaquat National Hospital and Medical College Karachi, Karachi, Pakistan

⁶ College of Medicine, AlMaarefa University, Riyadh, Saudi Arabia

⁷ Department of Pharmaceutical Sciences, University of Nigeria, Nsukka, Nigeria

⁸ National Health Service, England, UK

⁹ Department of Internal Medicine, State Hospital Abeokuta, Abeokuta, Nigeria

¹⁰ Department of Medicine, Ladoko Akintola University of Technology, Ogbomoso, Nigeria



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Background

The age-long campaign to eradicate malaria has witnessed tremendous progress in recent years. The use of insecticide-treated nets (ITNs) and other vector control strategies has yielded significant hope for malaria elimination [1]. Furthermore, recent advancements, such as the roll-out of novel malaria vaccines, usher in a new era for malaria infection prevention [2]. Despite these efforts, malaria continues to pose a formidable public health challenge globally, disproportionately affecting low- and middle-income countries, particularly in the sub-Saharan African region [3]. According to the World Health Organization (WHO), approximately 597,000 malaria-related deaths were recorded in 2023, alongside over 263 million new cases, with 95% originating from the WHO African Region [4].

In the last three years, the WHO has qualified and recommended two malaria vaccines—RTS,S/AS01 (RTS,S) in 2021 and R21/Matrix-M™ in 2023 [1, 4]. These milestones have heightened hopes for achieving global targets to reduce malaria cases, morbidity, and mortality rates by at least 90% by 2030 [4]. Currently, over 2 million children have received a dose of RTS,S/AS01 through regional vaccine implementation programmes [4, 5]. However, these achievements are threatened by the surge of resistance to anti-malarial medications [6], showing the urgent need for innovative strategies that integrate gene therapy and genomic editing to enhance vector control mechanisms. CRISPR/Cas9 technology has emerged as a potent tool for genetic research, allowing precise DNA modifications through the targeted cutting and inserting of genetic information [7]. Leveraging CRISPR/Cas9 for genetic engineering has the potential to combat diseases like malaria by genetically modifying *Anopheles* mosquitoes, making them unable to transmit the malaria parasite or significantly decreasing their populations [8]. This gene drive technology operates in two main ways: population suppression, which reduces mosquito populations, and population replacement, where non-malaria-transmitting mosquitoes are introduced to the existing population [12, 13].

The integration of malaria vaccination with CRISPR/Cas9 gene drive technology represents a promising dual approach that could significantly reduce malaria-associated morbidities and mortalities. Despite the emergence of some unfavourable outcomes associated with these technologies [11, 14], the application of CRISPR methodologies has catalyzed scientific investigations into critical pathways and the optimization of vector control strategies. This perspective paper aims to provide an examination of the synergistic potential of malaria vaccines and CRISPR/Cas9 gene drive technology in the global endeavor to eradicate malaria.

The need for integrated approaches

The fight against malaria, a disease that has plagued humanity for centuries, requires innovative and multifaceted strategies [15]. While traditional treatments, such as chemoprophylaxis, vector control, and vaccination, have long been employed in malaria control efforts, the use of *Artemisia annua* and its derivatives for malaria treatment emerged only in the late twentieth century despite the plant's earlier historical applications in traditional medicine. Recent global initiatives, however, show the urgent need for integrated and innovative approaches to combat this disease effectively [16]. Since the second global effort to eradicate malaria began in 2006, significant progress has been made in reducing mortality rates and advancing malaria vaccine development through international collaboration and funding [17]. However, despite these advances, existing malaria vaccines have shown limited effectiveness due to the complex interplay between the *Plasmodium* parasite and the human immune system [18].

The WHO established product characteristics for malaria vaccines, targeting a reduction in malaria infections of at least 90% over 12 months and a minimum of 45% over 32 months for certain categories [19]. Historically, vaccines like RTS,S, which targets the circumsporozoite protein (CSP), achieved 46% efficacy in children and 27% in infants during Phase III trials. However, this vaccine had no significant impact on severe malaria incidence during the trials, and immunity waned quickly, necessitating additional booster doses [20, 21]. Despite these limitations, findings from the 2024 World Malaria Report demonstrated that the introduction of RTS,S resulted in a statistically significant 13% reduction in all-cause mortality (excluding injury) and a 22% reduction in hospitalized severe malaria among age-eligible children, underscoring its real-world impact [4]. Moreover, recent CSP-based vaccines, such as R21, have demonstrated reduced efficacy, while blood-stage vaccines targeting various *Plasmodium* proteins have struggled to meet efficacy expectations, with some, like GM22, achieving only 14% efficacy [22]. These challenges show the need for more effective vaccines that can provide durable immunity against malaria.

Gene drive technology, particularly through CRISPR/Cas9, represents a revolutionary approach to malaria control by allowing targeted modifications of mosquito populations to reduce *Plasmodium* transmission [23, 24]. Despite its potential, gene drive technology faces significant hurdles, including the risk of creating resistant mosquito strains and declining efficacy across generations [25, 26]. Although newer versions of population suppression show no evidence for selection of resistance [27, 28]. Furthermore, ethical concerns regarding irreversible

genetic changes and potential ecological impacts require thorough consideration and active stakeholder engagement to ensure responsible implementation [29–32].

Integrating malaria vaccines with CRISPR/Cas9 gene drive technology represents a transformative strategy in the global fight against malaria, addressing both the biological complexities of the disease and the socio-economic factors that perpetuate its prevalence [33]. The current malaria vaccine landscape is characterized by limited efficacy and short-lived immunity, necessitating innovations that enhance protective responses [17]. Integrating CRISPR/Cas9 gene drive technology offers a complementary approach. Modifying the genetic makeup of mosquito populations to reduce their capacity to transmit malaria presents a viable strategy for decreasing the incidence of malaria infections within vaccinated populations [34]. Gene drives can be designed to introduce genetic modifications that reduce the fertility or lifespan of female *Anopheles* mosquitoes, the primary vectors responsible for transmitting the *Plasmodium* parasite to humans [35]. This strategic reduction in mosquito populations, when integrated with effective vaccination campaigns, has the potential to significantly lower malaria transmission rates, thereby enhancing the impact of vaccination efforts [36].

The integration of vaccines and gene drive technology can create a synergistic effect that addresses both immediate and long-term challenges associated with malaria transmission. Vaccines are capable of priming the immune system to recognize *Plasmodium* antigens, facilitating rapid immune responses upon subsequent exposure [37]. Concurrently, gene drives can systematically diminish the population of malaria-transmitting mosquitoes and exposure to the malaria parasite, by either decreasing the number of vector mosquitoes or the ability of the parasite to develop within the mosquitoes [38]. For instance, a population-level vaccination strategy could focus on high-risk groups, including children under five and pregnant women, while also implementing the deployment of gene drive mosquitoes in endemic regions. As vaccination coverage increases, malaria transmission will decrease due to the combination of decreased exposure to infected mosquito vectors and decreased parasite development in the human host and selective pressure on the mosquito population would be reduced, resulting in decreased transmission rates [39]. This integrated approach promises not only to lower the incidence of malaria infections but also to mitigate the severity of cases, thereby reducing overall morbidity and mortality rates [40, 41].

Achieving the ambitious target set by the WHO of a 90% reduction in malaria incidence and mortality by 2030 necessitates a strategy that incorporates advancements

in both vaccine and gene drive technologies [8]. Furthermore, a broader, coordinated effort is essential. This includes the ongoing development and deployment of effective anti-malarial drugs critical for treating infections and curbing transmission [9]. Community engagement is crucial for educating populations about malaria prevention, vaccination strategies, and the role of gene drive technology. Such engagement fosters trust and enhances the effectiveness of public health initiatives [10]. Moreover, effective vector control strategies must prioritize environmental sustainability, ensuring that the implementation of gene drives minimizes ecological disruption while preserving non-malaria-transmitting mosquito populations [12]. Comprehensive risk assessments, including analyses of potential pathways to harm, are essential to achieve this balance [42–44]. The establishment of robust monitoring systems will be vital for evaluating the effectiveness of these integrated interventions. Continuous assessment will facilitate adaptive management strategies that respond to emerging challenges and shifting epidemiological patterns.

The burden of malaria extends beyond health implications, significantly impacting economic development in endemic regions. Integrating malaria vaccines with gene drive technology can lead to substantial reductions in disease prevalence, resulting in improved health outcomes and increased productivity. Countries heavily burdened by malaria frequently experience impediments to economic growth due to healthcare costs, workforce productivity losses, and diminished educational opportunities. Effectively addressing malaria through integrated strategies not only alleviates direct health impacts but also contributes to broader socio-economic development. This transformation can foster healthier communities, enhance educational outcomes, and create increased economic opportunities, ultimately disrupting the cycle of poverty that malaria exacerbates [45].

Strategy for integration into current programmes

Integrating the combined approach of malaria vaccines and CRISPR/Cas9 gene drive technology into existing malaria control programmes necessitates meticulous planning and coordination. An evaluation of current malaria control initiatives is imperative to identify their strengths, weaknesses, and specific gaps regarding the integration of malaria vaccines and CRISPR/Cas9 technologies. This assessment should involve engaging local health authorities, community stakeholders, and researchers to gain a thorough understanding of how current programmes address malaria transmission. Particular attention should be paid to existing vaccination campaigns and vector control measures, with the aim of determining how CRISPR/Cas9 gene drives can

complement or enhance these efforts by targeting resistant mosquito populations and improving overall transmission dynamics.

Furthermore, fostering collaboration among key stakeholders is vital for creating a unified vision for integrating vaccines and gene drive technologies into established malaria control programmes. This collaboration should encompass government agencies, non-governmental organizations (NGOs), research institutions, and community leaders. Establishing platforms for continuous dialogue and knowledge sharing will facilitate the exchange of best practices, lessons learned, and innovative ideas for leveraging the complementary strengths of vaccines and gene drive technology. Engaging stakeholders early in the integration process will ensure that diverse perspectives inform the development of protocols that optimize the effectiveness of both strategies in reducing malaria transmission [46].

Investing in the training of healthcare workers, field staff, and community members is essential to ensure they possess the requisite knowledge and skills to implement the integrated strategies effectively. Training programmes should focus on the scientific principles underlying both vaccination and gene drive technologies, emphasizing how these approaches work synergistically to enhance malaria control. Additionally, training should highlight the distinct implementation strategies of the two tools: vaccination, which is delivered on an individual basis through healthcare systems, and gene drive mosquitoes, which are deployed on an area-wide basis to impact mosquito populations across regions. Development of educational materials aimed at raising awareness about the benefits and safety of the integrated approach is crucial. These materials should specifically address potential misconceptions regarding gene drive technology and highlight its role in complementing vaccination efforts, thereby fostering community acceptance and support for the integrated strategy.

Moreover, the initiation of pilot projects with gene drive mosquitoes trials that will be done in the context of other malaria control methods currently used at the trial site in select malaria-endemic regions is recommended to test the feasibility, effectiveness, and community acceptance of the integrated strategy before broader deployment [8, 34]. These pilot projects should be designed to assess the simultaneous deployment of vaccines and gene drive mosquitoes, with a focus on evaluating how the two interventions interact in real-world settings. Data collected during these projects will yield valuable insights that can refine implementation protocols, assess logistical requirements, and establish robust monitoring frameworks to track progress and outcomes. Additionally, findings from these pilot projects will be instrumental in

informing subsequent larger-scale interventions, ensuring that both vaccination and gene drive technologies are integrated in a way that maximizes their combined impact.

Establishing robust monitoring and evaluation (M&E) systems is critical for assessing the impact of the integrated approach on malaria transmission, community health outcomes, and ecological implications. Real-time data collection mechanisms should be employed to monitor vaccination coverage, mosquito population dynamics, and transmission rates in areas where gene drives are deployed. The M&E framework should include specific indicators that assess the effectiveness of the combined approach, such as reductions in malaria incidence and changes in mosquito behavior. This dynamic evaluation process will enable adaptive management, allowing for prompt adjustments to strategies based on emerging evidence and stakeholder feedback. By continuously assessing the integration of vaccines and gene drive technologies, the approach can remain responsive to evolving challenges and opportunities in malaria control.

Challenges and future considerations

While the integration of malaria vaccines and CRISPR/Cas9 gene drive technology presents a promising strategy for combating malaria transmission, several significant challenges must be addressed to ensure successful implementation and sustainability of this combined approach. Navigating complex regulatory frameworks for gene drive technology is a prominent barrier to integration [47]. The regulatory landscape varies significantly across countries and regions, often encompassing rigorous safety assessments, ethical considerations, and environmental impact studies [8, 47, 48]. To facilitate compliance, it is essential to work closely with regulatory bodies to ensure that all aspects of the integrated strategy meet established guidelines [49]. This requires an ongoing dialogue with policymakers, who play a critical role in shaping regulations that can adapt to emerging biotechnologies. Building a collaborative relationship with these stakeholders will be crucial for promoting timely approvals while ensuring the safety and efficacy of interventions [50]. Additionally, transparency in the regulatory process can help alleviate public concerns, fostering a more conducive environment for the adoption of innovative solutions.

Community concerns about genetic modification and its implications for ecosystems can significantly hinder the acceptance of gene drive technology. Many communities harbour misconceptions regarding the safety and efficacy of genetic interventions, often fueled by misinformation and fear of unintended consequences [51]. To mitigate stakeholder resistance, it is essential to engage in transparent communication, fostering community

education and actively involving local populations in decision-making processes. Strategies should include public forums, workshops, and information campaigns that address common concerns, clarify the scientific basis of the technologies, and highlight potential benefits. Building trust in scientific innovations through open dialogue and community involvement will be key to ensuring broad-based support for the integrated strategy.

The release of genetically modified mosquitoes raises potential ecological concerns, including unintended effects on non-target species and ecosystem dynamics. Comprehensive ecological assessments must be conducted to evaluate these impacts prior to large-scale implementation. These assessments should encompass both short-term and long-term studies to understand how gene drive technology interacts with existing ecosystems [52, 53]. Additionally, establishing monitoring frameworks to track ecological changes post-implementation will be essential. By proactively assessing and addressing potential ecological risks, stakeholders can ensure that malaria control efforts do not inadvertently harm biodiversity or disrupt ecosystem services, ultimately supporting the sustainability of the integrated strategy.

Securing sustainable funding for integrated programmes poses a significant challenge, particularly in low-resource settings where malaria prevalence is often highest [4, 54]. The reliance on external funding can lead to programme discontinuity and hinder long-term planning. To overcome this challenge, partnerships with international organizations, government agencies, and private sector stakeholders will be necessary to support sustained investments in malaria control initiatives. Innovative financing mechanisms, such as public–private partnerships, could be explored to ensure continuous funding and resource allocation. Additionally, leveraging existing funding streams for malaria control to incorporate new technologies can enhance the overall efficiency and impact of resource utilization.

Continuous monitoring of both *Plasmodium* and mosquito populations is vital to detect and address emerging resistance to interventions. The potential for resistance development poses a significant threat to the long-term effectiveness of integrated malaria control strategies. Adaptation strategies must be developed to counteract potential resistance mechanisms, including the integration of complementary interventions that can enhance the durability of both vaccines and gene drive technologies. Ongoing research to understand the dynamics of resistance, including genetic and environmental factors that contribute to it, will be critical in maintaining the effectiveness of integrated strategies. By proactively addressing resistance, stakeholders can enhance the

resilience of malaria control efforts and ensure sustained reductions in transmission rates.

Conclusion

The integration of malaria vaccines with CRISPR/Cas9 gene drive technology presents a transformative opportunity to combat malaria, addressing the multifaceted challenges associated with its transmission and impact. As the global burden of malaria continues to weigh heavily on public health systems, particularly in low- and middle-income countries, the necessity for innovative and synergistic strategies has never been more critical. Leveraging the strengths of both vaccination and gene drive technologies can create a robust framework for malaria control that enhances efficacy and sustainability.

The collaborative potential of vaccines and gene drives could revolutionize malaria prevention efforts, yielding substantial reductions in morbidity and mortality. Vaccines can prime the immune system to recognize and respond to *Plasmodium* infections, reducing parasite development in the human host, while gene drives can systematically diminish exposure to the malaria parasite by either decreasing the populations of malaria-transmitting mosquitoes or reducing the parasite's ability to develop within mosquitoes. This dual approach not only targets the disease at different stages but also offers a more resilient strategy against emerging challenges, such as anti-malarial resistance and ecological disruptions. However, realizing this vision requires careful consideration of regulatory, ethical, and ecological challenges. Engaging with communities to foster understanding and acceptance of genetic technologies is essential for ensuring the success of integrated interventions. Furthermore, robust monitoring and evaluation systems will be critical to assess the effectiveness and impact of these strategies on malaria transmission dynamics and ecological health. As the World Health Organization's goal of reducing malaria incidence and mortality by 90% by 2030 approaches, the integration of innovative technologies alongside traditional methods must be prioritized. Establishing strong collaborations among stakeholders, securing sustainable funding, and continuously adapting to emerging evidence will forge a path toward a future where malaria is no longer a pervasive threat to global health. The combined application of malaria vaccines and CRISPR/Cas9 gene drive technology has the potential to reshape approaches to malaria eradication and serves as a model for future public health interventions against complex infectious diseases.

Abbreviations

CRISPR	Clustered Regularly Interspaced Short Palindromic Repeats
DNA	Deoxyribonucleic Acid
mRNA	Messenger Ribonucleic Acid

RNA	Ribonucleic Acid
WHO	World Health Organization
GMO	Genetically Modified Organism
IGR	Insect Growth Regulator
TDR	Target Product Profile
WHO-TAG	World Health Organization's Technical Advisory Group
LMICs	Low- and Middle-Income Countries
CDC	Centers for Disease Control and Prevention
SP	Sulfadoxine-Pyrimethamine (an antimalarial drug)
RDT	Rapid Diagnostic Test
PEI	Pathogen-Environment Interactions
SNP	Single Nucleotide Polymorphism
MDA	Mass Drug Administration

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JEA conceptualised the study; All authors were involved in the literature review; JEA & NA extracted the data from the reviews studies; All authors wrote the final and first drafts. All authors read and approved the final manuscript.

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Declarations

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The authors declare that they have no competing interests

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References

- Aderinto N, Olatunji G, Kokori E, Sikirullahi S, Aboje JE, Ojabo RE. A perspective on Oxford's R21/Matrix-M™ malaria vaccine and the future of global eradication efforts. *Malar J*. 2024;23:16.
- Anwar M. An overview of malaria and *Plasmodium*. In: Tarique M (ed.). *Drug targets for Plasmodium falciparum: historic to future perspectives*. Springer, Singapore. 2024.
- Olatunji G, Kokori E, Kwape JM, Olatunji D, Anthony CS, Ogieuhi JJ, et al. *Anopheles stephensi* and the impending challenge to malaria eradication in Africa. *New Microbes New Infect*. 2024;58:101232.
- WHO. World malaria report 2024. Geneva, World Health Organization, 2024. <https://www.who.int/teams/global-malaria-programme/reports/world-malaria-report-2024>. (Accessed 14 Dec 2024)
- Osoro CB, Ochodo E, Kwambai TK, Otieno JA, Were L, Sagam CK, et al. Policy uptake and implementation of the RTS, S/AS01 malaria vaccine in sub-Saharan African countries: status 2 years following the WHO recommendation. *BMJ Glob Health*. 2024;9: e014719.
- Ishizaki T, Hernandez S, Paoletta MS, Sanderson T, Bushell ESC. CRISPR/Cas9 and genetic screens in malaria parasites: small genomes, big impact. *Biochem Soc Trans*. 2022;50:1069–79.
- Xu Y, Li Z. CRISPR-Cas systems: overview, innovations and applications in human disease research and gene therapy. *Comput Struct Biotechnol J*. 2020;18:2401–15.
- WHO. Guidance framework for testing of genetically modified mosquitoes. 2nd Edn. Geneva, World Health Organization, 2021. <https://www.who.int/publications/i/item/9789240025233>. (Accessed 15 Dec 2024)
- Vinayak S, Pawlowic MC, Sateriale A, Brooks CF, Studstill CJ, Bar-Peled Y, Cipriano MJ, Striepen B. Genetic modification of the diarrhoeal pathogen *Cryptosporidium parvum*. *Nature*. 2015;523:477–80.
- Sidik SM, Huet D, Ganesan SM, Huynh MH, Wang T, Nasamu AS, et al. A genome-wide CRISPR screen in *Toxoplasma* identifies essential Apicomplexan genes. *Cell*. 2016;166:1423–1435.e12.
- D'Amato R, Taxiarchi C, Galardini M, Trusso A, Minuz RL, Grilli S, et al. Anti-CRISPR *Anopheles* mosquitoes inhibit gene drive spread under challenging behavioural conditions in large cages. *Nat Commun*. 2024;15:952.
- Gantz VM, Jasinskiene N, Tatarenkova O, Fazekas A, Macias VM, Bier E, et al. Highly efficient Cas9-mediated gene drive for population modification of the malaria vector mosquito *Anopheles stephensi*. *Proc Natl Acad Sci USA*. 2015;112:E6736–43.
- Hammond A, Galizi R, Kyrkou K, Simoni A, Siniscalchi C, Katsanos D, et al. A CRISPR-Cas9 gene drive system targeting female reproduction in the malaria mosquito vector *Anopheles gambiae*. *Nat Biotechnol*. 2016;34:78–83.
- Macias VM, McKeand S, ChaverraRodriguez D, Hughes GL, Fazekas A, Pujhari S, et al. Cas9-mediated gene-editing in the malaria mosquito *Anopheles stephensi* by ReMOT Control. *G3 Bethesda*. 2020;10:1353–60.
- Fikadu M, Ashenafi E. Malaria: an overview. *Infect Drug Resist*. 2023;16:3339–47.
- Moxon CA, Gibbins MP, McGuinness D, Milner DA Jr, Marti M. New insights into malaria pathogenesis. *Annu Rev Pathol*. 2020;15:315–43.
- Nevagi RJ, Good MF, Stanisic DI. *Plasmodium* infection and drug cure for malaria vaccine development. *Expert Rev Vaccines*. 2021;20:163–83.
- Stanisic DI, McCall MB. Correlates of malaria vaccine efficacy. *Expert Rev Vaccines*. 2021;20:143–61.
- WHO. Malaria vaccines: preferred product characteristics and clinical development considerations. Geneva: World Health Organization; 2022.
- Duffy PE, Patrick GJ. Malaria vaccines since 2000: progress, priorities, products. *NPJ Vaccines*. 2020;5:48.
- RTS, S Clinical Trials Partnership. Efficacy and safety of the RTS, S/AS01 malaria vaccine during 18 months after vaccination: a phase 3 randomized, controlled trial in children and young infants at 11 African sites. *PLoS Med*. 2014;2014(11): e1001685.
- Stanisic DI, Good MF. Malaria vaccines: progress to date. *BioDrugs*. 2023;37:737–56.
- Makuvura Z, Magano SR, Mugumbate G. Malaria situation and elimination in Africa with specific focus on Zimbabwe: a systematic review. *Cogent Public Health*. 2024;11:2376945.
- Wagnew Y, Hagos T, Weldegerima B, Debie A. Willingness to pay for childhood malaria vaccine among caregivers of under-five children in Northwest Ethiopia. *Clinicoecon Outcomes Res*. 2021;13:165–74.
- Tajudeen YA, Oladipo HJ, Oladunjoye IO, Oladipo MK, Shittu HD, Abdulmumeen IF, et al. Transforming malaria prevention and control: the prospects and challenges of gene drive technology for mosquito management. *Ann Med*. 2023;55:2302504.
- Mudziwapasi R, Changara MC, Ndudzo A, Kaseke T, Godobo F, Mtemeli FL, et al. Gene drives in malaria control: what we need to know. *Biotechnol Biotechnol Equip*. 2021;35:1623–31.
- Hammond A, Pollegioni P, Persampieri T, North A, Minuz R, Trusso A, et al. Gene-drive suppression of mosquito populations in large cages as a bridge between lab and field. *Nat Commun*. 2021;12:4589.
- Lanzaro GC, Sánchez C, Collier TC, Marshall JM, James AA. Population modification strategies for malaria vector control are uniquely resilient to observed levels of gene drive resistance alleles. *BioEssays*. 2021;43: e2000282.
- Hammond AM, Kyrkou K, Bruttini M, North A, Galizi R, Karlsson X, et al. The creation and selection of mutations resistant to a gene drive over multiple generations in the malaria mosquito. *PLoS Genet*. 2017;13: e1007039.
- Sykes N, Bigirwenkya J, Coche I, Drabo M, Dzokoto D, O'Loughlin S, et al. Procedural legitimacy: co-developing a community agreement model for genetic approaches research to malaria control in Africa. *Malar J*. 2024;23:359.

31. Pare Toe L, Dicko B, Linga R, Barry N, Drabo M, Sykes N, et al. Operationalizing stakeholder engagement for gene drive research in malaria elimination in Africa—translating guidance into practice. *Malar J*. 2022;21:225.
32. Kormos A, Nazaré L, dos Santos AA, Lanzaro GC. Practical application of a relationship-based model to engagement for gene-drive vector control programs. *Am J Trop Med Hyg*. 2024;111:341–60.
33. Duffy PE, Gorres JP, Healy SA, Fried M. Malaria vaccines: a new era of prevention and control. *Nat Rev Microbiol*. 2024;22:756–72.
34. Orok AB, Ajibaye O, Aina OO, Iboma G, Adagyo Oboshi S, Iwalokun B. Malaria interventions and control programmes in sub-Saharan Africa: a narrative review. *Cogent Med*. 2021;8:1940639.
35. Arora N, Anbalagan CL, Pannu AK. Towards eradication of malaria: is the WHO's RTS, S/AS01 vaccination effective enough? *Risk Manag Healthc Policy*. 2021;14:1033–9.
36. Hammond AM, Galizi R. Gene drives to fight malaria: current state and future directions. *Pathog Glob Health*. 2017;111:412–23.
37. Bottino-Rojas V, James AA. Population modification using gene drive for reduction of malaria transmission. In: Benedict, Scott MJ (eds). *Transgenic insects: techniques and applications*. Chapt 11. CABI Publ. 2022.
38. Sinden RE. Targeting the parasite to suppress malaria transmission. *Adv Parasitol*. 2017;97:147–85.
39. James S, Collins FH, Welkhoff PA, Emerson C, Godfray HC, Gottlieb M, et al. Pathway to deployment of gene drive mosquitoes as a potential biocontrol tool for elimination of malaria in sub-Saharan Africa: recommendations of a scientific working group. *Am J Trop Med Hyg*. 2018;98(Suppl 6):1–49.
40. Arama C, Troye-Blomberg M. The path of malaria vaccine development: challenges and perspectives. *J Intern Med*. 2014;275:456–66.
41. Hancock PA, North A, Leach AW, Winskill P, Ghani AC, Godfray HCJ, et al. The potential of gene drives in malaria vector species to control malaria in African environments. *Nat Commun*. 2024;15:8976.
42. Teem JL, Ambali A, Glover B, Ouedraogo J, Makinde D, Roberts A. Problem formulation for gene drive mosquitoes designed to reduce malaria transmission in Africa: results from four regional consultations 2016–2018. *Malar J*. 2019;18:347.
43. Connolly JB, Mumford JD, Fuchs S, Turner G, Beech C, North AR, et al. Systematic identification of plausible pathways to potential harm via problem formulation for investigational releases of a population suppression gene drive to control the human malaria vector *Anopheles gambiae* in West Africa. *Malar J*. 2021;20:170.
44. Kormos A, Dimopoulos G, Bier E, Lanzaro GC, Marshall JM, James AA. Conceptual risk assessment of mosquito population modification gene-drive systems to control malaria transmission: preliminary hazards list workshops. *Front Bioeng Biotechnol*. 2023;11:1261123.
45. World Economic Forum. Here's how reducing malaria can add \$16 billion to Africa's GDP every year. 2024. <https://www.weforum.org/stories/2024/06/malaria-global-health-economy-africa/>. (Accessed 19 Dec 2024)
46. Finda MF, Juma EO, Kahamba NF, Mthawani RS, Sambo M, Emidi B, et al. Perspectives of African stakeholders on gene drives for malaria control and elimination: a multi-country survey. *Malar J*. 2023;22:384.
47. James SL, Dass B, Quemada H. Regulatory and policy considerations for the implementation of gene drive-modified mosquitoes to prevent malaria transmission. *Transgenic Res*. 2023;32:17–32.
48. Nateghi RM. CRISPR/Cas9 gene drive technology to control transmission of vector-borne parasitic infections. *Parasite Immunol*. 2020;42: e12762.
49. Carballar-Lejarazú R, Ogaugwu C, Tushar T, Kelsey A, Pham TB, Murphy J, et al. Next-generation gene drive for population modification of the malaria vector mosquito, *Anopheles gambiae*. *Proc Natl Acad Sci USA*. 2020;117:22805–14.
50. Cha J, Ranweiler JS, Lane PA. Stability studies. *Separation. Sci Technol*. 2001;3:445–83.
51. James SL, Marshall JM, Christophides GK, Okumu FO, Nolan T. Toward the definition of efficacy and safety criteria for advancing gene drive-modified mosquitoes to field testing. *Vector-Borne Zoonotic Dis*. 2020;20:237–51.
52. Subsidiary Body on Scientific, Technical and Technological Advice. Additional voluntary guidance materials to support case-by-case risk assessments of living modified organisms containing engineered gene drives. Twenty-sixth meeting; 2024. Nairobi.
53. Conference of the parties to the convention on biological diversity serving as the meeting of the parties to the Cartagena protocol on biosafety. review of effectiveness of processes under the convention and its protocols. Eleventh meeting; 2024 Nov 1; Cali, Colombia
54. Van Meer PJ, Graham ML, Schuurman HJ. The safety, efficacy and regulatory triangle in drug development: Impact for animal models and the use of animals. *Eur J Pharmacol*. 2015;759:3–13.

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