

Gaps in protection: the actual challenge in malaria elimination



Krijn P. Paaijmans^{1,2,3,4,5*} and Neil F. Lobo^{6,7}

Abstract

Progress in reducing both malaria cases and deaths has stalled with regression seen in many geographies. While significant attention is given to the contributing challenges of drug and insecticide resistance, 'residual' malaria is often diminished to transmission resulting from outdoor-biting or zoophagic/opportunistic mosquito vectors. These specific vector bionomic traits are only part of the problem, as residual transmission may be driven by (a combination of) (1) sub-optimal intervention coverage, guality, acceptance, and/or usage, (2) drug resistance, (3) insecticide resistance, (4) refractory, resistant and adaptive vector and human behaviours that lower intervention effectiveness, (5) lack of, limited access to, and/or willingness to use healthcare systems, (6) diagnostic sensitivity along with the parallel issue of hrp2/3 mutations, (7) (inter)national policy, (8) the research and development pipeline, and (9) external factors such as natural disasters and conflict zones. Towards combating the minimization of this extensive and multipronged issue among the scientific community, funding agencies, and public health officials responsible for guiding or developing malaria programmes, an alternative way of describing this transmission is proposed by focusing in on the causative 'gaps in protection'. Defining and wording it as such zeros in on the drivers that result in the observed remaining (or increasing) transmission, allowing the malaria community to focus on solutions by identifying the actual causes. Outlining, defining and quantifying the gaps in protection for a given system is of utmost importance to understand what needs to be done, differentiating what can be done versus what cannot be tackled at that moment, along with delineating the technical and financial capacity required.

*Correspondence:

Krijn P. Paaijmans

- kpaaijma@asu.edu
- ¹ Center for Evolution and Medicine, School of Life Sciences, Arizona State University, Tempe, AZ, USA
- ² The Biodesign Center for Immunotherapy, Vaccines and Virotherapy, Arizona State University, Tempe, AZ, USA
- ³ Simon A. Levin Mathematical, Computational and Modeling Sciences
- Center, Arizona State University, Tempe, AZ, USA

⁴ Centro de Investigação Em Saúde de Manhiça, Fundação Manhiça, Maputo, Mozambique

⁵ ISGlobal, Barcelona, Spain

 $^{\rm 6}$ Eck Institute for Global Health, University of Notre Dame, Notre Dame, IN, USA

⁷ Malaria Elimination Initiative, University of California San Francisco, San Francisco, CA, USA

Background

The scaling up of core malaria interventions to prevent onward transmission has led to large reductions in the global malaria burden since 2000. Diagnosis and treatment of clinical malaria cases with artemisinin-based combination therapy, insecticide-treated nets (ITNs) later replaced by long-lasting insecticidal nets (LLINs) and indoor residual spraying (IRS) are the three core interventions that are estimated to have averted up to 663 million clinical cases between 2000 and 2015, halved *Plasmodium falciparum* infection prevalence in endemic Africa and reduced clinical disease by 40% [1]. However, malaria cases are on the rise again and progress has stalled [2]. In addition, while the use of recommended malaria control interventions has led to large reductions in the malaria burden in several African malaria



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

elimination settings [3, 4], local elimination has not been achieved.

The plateauing of intervention efficacies resulted in the concept of 'residual transmission' gaining significant interest and attention over recent years [5-7]. The World Health Organization (WHO) defined residual malaria transmission in 2016 as 'persistence of transmission after good coverage has been achieved with high-quality vector control interventions to which local vectors are fully susceptible'. In 2018, this definition was updated to 'persistence of malaria transmission following the implementation in time and space of a widely effective malaria programme' [8], indicating a constructive shift in focus away from intervention strategies that only target the mosquito vector. However, the word 'residual transmission' denotes that this is 'remaining (transmission) after the greater part or quantity has gone, which is not necessarily the case. A large portion of transmission may still be present even after the implementation of recommended interventions-evident in the recent increases of transmission seen [2].

This 'residual' malaria is often coupled with transmission resulting from outdoor-biting vectors, or zoophagic/ opportunistic mosquitoes (i.e. those mosquitoes feeding primarily on animals) [6, 7]-thereby minimizing and relegating this to being a 'vector issue'. This is also not necessarily true or a primary causal factor as demonstrated by human behaviour analysis and the association of primary vectors with indoor transmission in areas with high LLIN coverage [9–12]. A primary, but often ignored paradigm, is that the transmission system intrinsically changes when an intervention strategy is implemented. Present recommended monitoring and evaluation frameworks do not generally adapt to this simple concept and the failure to monitor and understand these shifting drivers of transmission results in the inability to appropriately adapt the intervention strategy towards being continuously effective.

Ironically, mosquito and human behaviours may only be part of the problem, as residual transmission may be driven by (a combination of) (1) sub-optimal intervention access, coverage, quality, acceptance, and/or usage [13, 14], (2) drug resistance [15, 16], (3) insecticide resistance [17], (4) refractory, resistant and adaptive vector and human behaviours that lower intervention effectiveness [18], (5) lack of, limited access to, and/or willingness to use healthcare systems [19], (6) diagnostic sensitivity along with the parallel issue of hrp2/3 mutations [20], (7) (inter)national policy [21], (8) the research and development pipeline [22], and (9) external factors such as natural disasters and conflict zones [23, 24] (Fig. 1). Many of these observed drivers of the transmission presently represent barriers that are harder to combat than just 'providing one LLIN per two household members', since they not just represent requirements in technical and infrastructural capacity, but will also require perceptual, cultural, and behavioural shifts in how industry, policy makers, implementers and end-users approach disease elimination.

To decrease this minimization and possible misunderstanding among the scientific community, funding agencies, and public health officials responsible for guiding or developing malaria programs, an alternative way of describing this transmission is proposed here by focusing on the causative 'gaps in protection'. This term is used to describe a circumstance when (1) an individual's malaria infection is not cleared, and (2) an individual and/or household is potentially exposed to malaria infection (i.e. an infective mosquito bite) due to a lack of an effective and/or adequate protective or preventive intervention in place. These two points address both bi-directional (between humans and mosquitoes) malaria transmission and reservoirs, as well as the intervention frameworks that impact each. Most often, gaps in protection can be directly identified through an assessment of how interventions interact with local human and/or vector populations.

For the current interventions targeting the parasite reservoirs (drugs and vaccines), examples of gaps in protection may include undiagnosed cases due to lack of expertise or failing diagnostics, antimalarial stockouts, counterfeit drugs, non-adherence to the treatment regime, and the exclusion of certain populations (e.g., pregnant women, babies or elderly). For the current core vector control intervention (LLINs), examples of gaps in protection include insecticide resistance (reducing the effectiveness of the protection that the insecticide in LLINs provides), occasions when people are outdoors without protection against potentially infective mosquito bites, suboptimal coverage, and usage. For IRS, gaps in protection may include insecticide resistance, vector species that do not rest on sprayed surfaces, duration of IRS impact, spray quality, people washing walls after spraying, and so on.

Other external drivers of transmission that can also contribute to gaps in protection include natural disasters, such as hurricanes and cyclones, or extreme wet years (e.g., due to El Niño/ La Niña) that affect mosquito abundance and species diversity and typical transmission patterns. Conflict zones and the Covid-19 pandemic also result in systemic drivers of gaps in protection impacting both vector intervention frameworks as well as health system functionality.

Focusing on the gaps in protection allows for zeroing in on the drivers that result in the observed remaining (or increasing) transmission, allowing the malaria

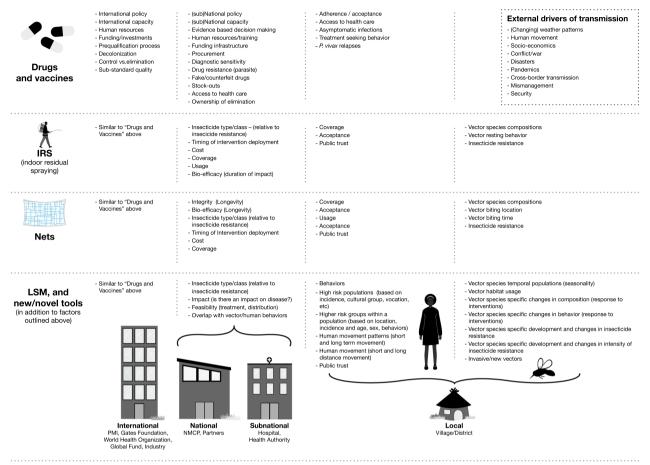


Fig. 1 Potential gaps in protection that can hamper malaria control and elimination efforts, across partners, interventions and geographies

community to focus on solutions by identifying the actual causes. Outlining, defining and quantifying all gaps for a given system is of utmost importance as it will highlight gaps that can be closed now and gaps for which there is currently no solution, and helps to outline the technical and financial capacity required. Solutions may be as simple as increasing intervention coverage or usage, or the implementation of a supplemental outdoor intervention that combats an outlined gap in protection -a targeted and tailored strategy that addresses local drivers. More systemic and forward-thinking solutions may include a cultural shift within a Ministry of Health or local at-risk communities, and developing novel supply chain mechanisms. It is imperative to outline gaps in protection that simply cannot be closed (i.e. coverage or adherence are at their maximum achievable levels, or there is no present recommended intervention that is suitable for a particular transmission space and time). Understanding the gaps in protection has an additional advantage-enabling appropriate expectations of effect with the implementation of, or change in an intervention strategy. Looking through this 'gaps in protection' lens, the implementor can now develop a focused and catered strategy, stratify responses based on characterized drivers and capacity, while also moderating expectations of impact. Consequently, control and elimination strategies may be optimized by adaptively tackling remaining gaps in protection while expectations are adjusted in parallel.

Outlining a strategic plan with all incident data, as well as the gaps in protection, allows for a realistic approach towards malaria elimination, encompassing what has been done, what should be most appropriate to the site, what is actually feasible, and most importantly, what cannot be done at present. This points to strategies, capacity, funding, the need to improve present or develop novel interventions, which are required towards reaching our goal of eliminating malaria.

Acknowledgements

We thank those who joined the discussion at our poster at the recent American Society of Tropical Medicine and Hygiene meeting (Seattle, 2022). The figure was created by Megan Joyce (Vislab, School of Life Sciences, Arizona State University, USA).

Author contributions

KPP and NFL contributed equally to the conceptualization and writing of the manuscript. Both authors read and approved the final manuscript.

Funding

This work was supported by the Bill and Melinda Gates Foundation (BMGF) and Obra Social "Ia Caixa" Partnership for the Elimination of Malaria in Southern Mozambique (INV-008483). NFL was supported by BMGF funded project A128488, through the UCSF Malaria Elimination Initiative.

Availability of data and materials

Not applicable.

Declarations

Ethics approval and consent to participate Not applicable.

Competing interests

The authors declare that they have no competing interests.

Received: 24 August 2022 Accepted: 29 January 2023 Published online: 07 February 2023

References

- 1. Bhatt S, Weiss DJ, Cameron E, Bisanzio D, Mappin B, Dalrymple U, et al. The effect of malaria control on *Plasmodium falciparum* in Africa between 2000 and 2015. Nature. 2015;526:207–11.
- WHO. World malaria report 2022. Geneva: World Health Organization; 2022.
- Galatas B, Saúte F, Martí-Soler H, Guinovart C, Nhamussua L, Simone W, et al. A multiphase program for malaria elimination in southern Mozambique (the Magude project): a before-after study. PLoS Med. 2020;17:e1003227.
- 4. Hsiang MS, Ntuku H, Roberts KW, Dufour M-SK, Whittemore B, Tambo M, et al. Effectiveness of reactive focal mass drug administration and reactive focal vector control to reduce malaria transmission in the low malaria-endemic setting of Namibia: a cluster-randomised controlled, open-label, two-by-two factorial design trial. Lancet. 2020;395:1361–73.
- Durnez L, Coosemans M. Residual transmission of malaria: an old issue for new approaches. In: Manguin S, editor. *Anopheles* mosquitoes—new insights into malaria vectors. InTech Online: Rijeka; 2013.
- 6. Carnevale P, Manguin S. Review of issues on residual malaria transmission. J Infect Dis. 2021;223:S61–80.
- Killeen G. Characterizing, controlling and eliminating residual malaria transmission. Malar J. 2014;13:330.
- WHO. Malaria terminology, 2021 update. Geneva: World Health Organization; 2021.
- Killeen GF, Monroe A, Govella NJ. Simplified binomial estimation of human malaria transmission exposure distributions based on hard classification of where and when mosquitoes are caught: statistical applications with off-the-shelf tools. Parasit Vectors. 2021;14:384.
- Yaro JB, Tiono AB, Sanou A, Toe HK, Bradley J, Ouedraogo A, et al. Risk factors associated with house entry of malaria vectors in an area of Burkina Faso with high, persistent malaria transmission and high insecticide resistance. Malar J. 2021;20:397.
- Monroe A, Moore S, Okumu F, Kiware S, Lobo NF, Koenker H, et al. Methods and indicators for measuring patterns of human exposure to malaria vectors. Malar J. 2020;19:207.
- Bayoh MN, Walker ED, Kosgei J, Ombok M, Olang GB, Githeko AK, et al. Persistently high estimates of late night, indoor exposure to malaria vectors despite high coverage of insecticide treated nets. Parasit Vectors. 2014;7:380.
- Koenker H, Taylor C, Burgert-Brucker CR, Thwing J, Fish T, Kilian A. Quantifying seasonal variation in insecticide-treated net use among those with access. Am J Trop Med Hyg. 2019;101:371–82.

- Opiyo M, Sherrard-Smith E, Malheia A, Nhacolo A, Sacoor C, Nhacolo A, et al. Household modifications after the indoor residual spraying (IRS) campaign in Mozambique reduce the actual spray coverage and efficacy. PLoS Glob Public Health. 2022;2:e0000227.
- Imwong M, Suwannasin K, Kunasol C, Sutawong K, Mayxay M, Rekol H, et al. The spread of artemisinin-resistant *Plasmodium falciparum* in the greater Mekong subregion: a molecular epidemiology observational study. Lancet Infect Dis. 2017;17:491–7.
- Wasakul V, Disratthakit A, Mayxay M, Chindavongsa K, Sengsavath V, Thuy-Nhien N, et al. Malaria outbreak in Laos driven by a selective sweep for *Plasmodium falciparum kelch13* R539T mutants: a genetic epidemiology analysis. Lancet Infect Dis. 2022;6:81.
- Hemingway J, Ranson H, Magill A, Kolaczinski J, Fornadel C, Gimnig J, et al. Averting a malaria disaster: will insecticide resistance derail malaria control? Lancet. 2016;387:1785–8.
- Govella N, Chaki P, Killeen G. Entomological surveillance of behavioural resilience and resistance in residual malaria vector populations. Malar J. 2013;12:124.
- O'Meara WP, Noor A, Gatakaa H, Tsofa B, McKenzie FE, Marsh K. The impact of primary health care on malaria morbidity-defining access by disease burden. Trop Med Int Health. 2009;14:29–35.
- Golassa L, Messele A, Amambua-Ngwa A, Swedberg G. High prevalence and extended deletions in *Plasmodium falciparum* hrp2/3 genomic loci in Ethiopia. PLoS ONE. 2020;15:e0241807.
- 21. Ruwanpura V, Neukom J, Grietens KP, Price RN, Thriemer K, Lynch CA. Opening the policy blackbox: unravelling the process for changing national diagnostic and treatment guidelines for vivax malaria in seven countries. Malar J. 2021;20:428.
- 22. Mwangoka G, Ogutu B, Msambichaka B, Mzee T, Salim N, Kafuruki S, et al. Experience and challenges from clinical trials with malaria vaccines in Africa. Malar J. 2013;12:86.
- 23. Grillet ME, Villegas L, Oletta JF, Tami A, Conn JE. Malaria in Venezuela requires response. Science. 2018;359:528.
- Kouadio IK, Aljunid S, Kamigaki T, Hammad K, Oshitani H. Infectious diseases following natural disasters: prevention and control measures. Expert Rev Anti Infect Ther. 2012;10:95–104.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

