



# *Systematic Review* **Advances in Malaria Diagnostic Methods in Resource-Limited Settings: A Systematic Review**

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**Abstract:** Malaria continues to pose a health challenge globally, and its elimination has remained a major topic of public health discussions. A key factor in eliminating malaria is the early and accurate detection of the parasite, especially in asymptomatic individuals, and so the importance of enhanced diagnostic methods cannot be overemphasized. This paper reviewed the advances in malaria diagnostic tools and detection methods over recent years. The use of these advanced diagnostics in lower and lower-middle-income countries as compared to advanced economies has been highlighted. Scientific databases such as Google Scholar, PUBMED, and Multidisciplinary Digital Publishing Institute (MDPI), among others, were reviewed. The findings suggest important advancements in malaria detection, ranging from the use of rapid diagnostic tests (RDTs) and molecular-based technologies to advanced non-invasive detection methods and computerized technologies. Molecular tests, RDTs, and computerized tests were also seen to be in use in resource-limited settings. In all, only twenty-one out of a total of eighty (26%) low and lower-middle-income countries showed evidence of the use of modern malaria diagnostic methods. It is imperative for governments and other agencies to direct efforts toward malaria research to upscale progress towards malaria elimination globally, especially in endemic regions, which usually happen to be resource-limited regions.

**Keywords:** malaria; polymerase chain reaction (PCR); loop-mediated isothermal amplification (LAMP); diagnostic methods

### **1. Introduction**

Malaria elimination has been a focal topic of public health discussions for the past decade or more. Despite being a tropically endemic parasitic infection, the impact of malaria is far reaching and remains a global health concern. The 2023 World Health Organization (WHO) report states that malaria cases rose to an estimated 249 million in 2022, with an increase of 5 million more cases from the year 2021 [\[1\]](#page-13-0). Although relentless efforts are being made and strategies put in place, much more is required to free our globe of the parasitic infection, particularly in indigenous malaria-endemic countries such as a number of sub-Saharan African countries where most cases occur [\[2\]](#page-13-1).



**Citation:** Yalley, A.K.; Ocran, J.; Cobbinah, J.E.; Obodai, E.; Yankson, I.K.; Kafintu-Kwashie, A.A.; Amegatcher, G.; Anim-Baidoo, I.; Nii-Trebi, N.I.; Prah, D.A. Advances in Malaria Diagnostic Methods in Resource-Limited Settings: A Systematic Review. *Trop. Med. Infect. Dis.* **2024**, *9*, 190. [https://doi.org/](https://doi.org/10.3390/tropicalmed9090190) [10.3390/tropicalmed9090190](https://doi.org/10.3390/tropicalmed9090190)

Academic Editors: John Frean and Shigeyuki Kano

Received: 27 June 2024 Revised: 31 July 2024 Accepted: 19 August 2024 Published: 23 August 2024



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Central to eliminating malaria is early, accurate detection, quantification, and differentiation of the parasitic infection, especially among asymptomatic persons. Asymptomatic plasmodium-infected individuals represent a major threat to malaria elimination worldwide as they do not show signs of clinical disease yet serve as parasite reservoirs and significantly contribute to the spread of the infection [\[3\]](#page-13-2). Notably, the majority of these asymptomatic infections are missed by conventional diagnostic techniques. As a result, the need for reliable, sensitive, and specific diagnostic or detection methods arises, which would also be useful for monitoring any decline in malaria transmission [\[4\]](#page-13-3).

Technologies for malaria diagnosis have advanced in recent years; however, certain factors, such as the lack of laboratory infrastructure, operational costs, electricity requirements, and special operation expertise, have impeded the implementation of these advanced techniques in the vast majority of malaria endemic areas. This is especially the case when it comes to molecular testing, as these tests can be particularly expensive in addition to other challenges not only for malaria but for other infectious diseases [\[5\]](#page-13-4). The WHO describes microscopy (thin and thick film) as the primary method of detection [\[6\]](#page-13-5). Though microscopy is extensively used, it is unable to adequately detect low parasitemia, which is essential for effective treatment and subsequent elimination of the parasitic infection [\[7\]](#page-13-6). In addition, it is a laborious process requiring much expertise and experience for accurate diagnosis [\[4](#page-13-3)[,8\]](#page-13-7). Other concerns have been the invasive approach of this technique, where blood samples are collected after a painful pierce of a needle, and yet an accurate diagnosis unassuredly relies solely on the discretion of the laboratory scientist. In several developing countries, there is inadequate expertise, equipment, and supplies required for accurate detection; as such, there are greater risks of contamination and false diagnosis [\[9\]](#page-13-8). Furthermore, it becomes more unreliable and difficult to distinguish low level infections as transmissions decline; hence, there is a need for alternative approaches to detection as elimination is being considered [\[4\]](#page-13-3).

Can there be a faster, more specific, and more sensitive method of detecting malaria that can easily be implemented in resource-limited areas? The question remains among scientists globally. Can malaria be eliminated and many more lives saved by the emergence of technologies that offer early detection and differentiation of very minimal malarial infections? Does mankind stand a chance of advancement towards needle-free malaria detection, point-of-care devices, and personalized malaria medicine? For lower and lowermiddle-income countries, which total 26 and 54, respectively (Table [1\)](#page-2-0), according to the World Bank, will there be access to such effective diagnostic tools [\[10\]](#page-13-9)? It is worthy of note that 11 of these countries, all in sub-Saharan Africa, bear 70% of the global malaria burden, according to the 2023 WHO report [\[1\]](#page-13-0) (Table [1\)](#page-2-0). The above-mentioned questions are but a few that remain on the minds of scientists and thus drive research.

Subsequently, there are several techniques that have been developed over the years to address some of the challenges with the gold standard technique. Rapid diagnostic tests (RDTs) are fast and reliable. Malaria RDTs do not require skilled personnel or constant electricity, but relative to malaria microscopy, they are expensive, have a short shelf life, and only give qualitative results [\[11\]](#page-13-10). Other diagnostic techniques, such as enzyme-linked immunosorbent assay (ELISA), lateral flow immunoassay (LFIA), microarrays, aptamer-based biosensors, genomic sequencing, loop-mediated isothermal amplification (LAMP), nested PCR, real-time PCR, and quantitative nucleic sequence-based amplification, are usually reserved for research and surveillance purposes. These techniques have higher sensitivity and specificity for malaria diagnosis relative to microscopy and RDTs. Notwithstanding, some of them are more laborious and expensive to deploy in resource-limited areas.

<span id="page-2-0"></span>**Table 1.** Countries classified by World Bank as low or lower-middle-income economies in 2024 [\[1,](#page-13-0)[10\]](#page-13-9). The table lists all resource-limited countries divided into low (top portion) and lower-middle (bottom portion) income countries, with special emphasis placed on the 11 countries (right portion) that together bear 70% of the global malaria burden.



This systematic review looks at traditional and modern techniques in light of their main advantages and disadvantages, as well as the countries where they have been used, with emphasis placed on lower and lower-middle-income countries. Also, emphasis would be placed on molecular-based techniques and how common they are in resource-limited settings, which usually happen to be endemic to malaria.

#### **2. Materials and Methods**

In conducting this systematic review, an accurate and authentic outcome was ensured by adherence to the Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) guidelines. The review was registered in the open science framework database [\(https://doi.org/10.17605/OSF.IO/DV6Z3](https://doi.org/10.17605/OSF.IO/DV6Z3) (accessed on 28 June 2024)). Relevant details needed were obtained from articles published in journals and from databases up to 2022 since the search was done in early 2023. Key search phrases used for the search included "malaria detection methods", "emerging technologies in malaria", "recent advances in malaria detection or diagnosis", "emerging methods in malaria diagnosis", "traditional methods of malaria detection", "Point of care devices for malaria detection", "Non-invasive or needle-free malaria detection", and "personalized malaria medicine". Numerous articles were obtained from databases, journals, and other publishing sites, including Google Scholar, PUBMED, and MDPI databases.

From the databases, a total of 327 were identified. After the searches, the publications were sorted out to remove duplicates, and 20 publications were removed. The records were further screened to remove all incomplete, unpublished articles, and ineligible publications. With articles published in recent years under consideration, all accessible publications were considerable options, leaving out articles from journals that needed to be purchased, were restricted, or there was not a PDF version of the complete paper readily available.

Upon abstract screening, articles were selected based on the following general criteria: a traditional or modern method of malaria detection or diagnosis was investigated. A total of 276 articles were obtained, uploaded into Mendeley Reference Manager and Endnote, and carefully reviewed for full text eligibility and results presentation.

Identification

Screening

Eligibility

Included

Of these articles, some investigated traditional methods, while most investigated various modern methods. Some articles were also found useful as they investigated or reviewed "emerging technologies in malaria detection" or "advances in malaria detection", among others, and thus were used in the other parts of the review writing. After a comprehensive document screening, 167 were later removed as it was found that they did not suit the review criteria, leaving 109 to be reviewed. These articles were removed for reasons including the papers being published earlier than 2014 (for those to be analyzed), they did not specifically investigate diagnostic tools for malaria, the article obtained was not the published version, and the research scope and contents were not clear or did not focus on a possible malaria detection method. In several instances, multiple malaria detection methods were identified in a single publication; thus, the number of developed methods identified exceeded the number of publications used. Figure [1](#page-3-0) below shows the sorting-out process.

<span id="page-3-0"></span>

**Figure 1.** A PRISMA flow diagram showing the method of article selection. **Figure 1.** A PRISMA flow diagram showing the method of article selection.

### **3. Results**

#### *3.1. Traditional Methods Used for Malaria Detection*

Table [2](#page-4-0) below shows a summary of traditionally used methods of malaria detection, elaborating on their approach as well as the pros and cons of using these methods for diagnosis. Though there has been the development of new and innovative methods of detection over the years, microscopy, using thick and thin blood films coupled with Giemsa staining, remains the gold standard for the diagnosis of malaria parasitic infections [\[8](#page-13-7)[,12\]](#page-13-11).

<span id="page-4-0"></span>



### *3.2. Modern Methods Used for Malaria Detection*

The quest to effectively treat malaria while gravitating towards its elimination has driven the development of various tools and assays for the diagnosis of malaria (4). Tables [3](#page-5-0) and [4](#page-7-0) contain recently developed methods used in the diagnosis of malaria and where they have been used. These diagnostic approaches vary greatly, ranging from biosensors and molecular assays down to computerized algorithms and automated analyzers, which have been developed or used over recent years, no earlier than 2014. The advantages and limitations of each diagnostic method are considered, as well as the summarized procedure by which it is conducted.

Table [5](#page-10-0) analyzes evidence of the use of some recently developed detection tools in lower and lower-middle-income countries where there are often resource limitations. The test types that featured most frequently in publications were PCR techniques (eleven), followed by RDT tests (nine), then LAMP techniques and computerized/digital deep machine learning approaches (six each). In all, twenty-one countries had publications featuring modern malaria diagnostic methods.

In Figure [2,](#page-10-1) the various methods of malaria detection reported from the identified studies have been represented graphically, indicating which diagnostic trends are being largely investigated, used more, or have gained much research interest. The chart represents malaria diagnostic developments investigated from 2014 until 2022. PCR-based methods and LAMP-based methods were the most prevalent methods.

<span id="page-5-0"></span>

**Table 3.** Modern (PCR/LAMP-based) methods used for malaria detection and evidence of use in developed countries.



## **Table 3.** *Cont.*

<span id="page-7-0"></span>

**Table 4.** Modern (non-PCR/non-LAMP-based) methods used for malaria detection and evidence of use in developed countries.



## **Table 4.** *Cont.*



**Table 4.** *Cont.*



<span id="page-10-0"></span>**Table 5.** Evidence of use of modern methods of malaria detection in low and lower-middle-income countries.

<span id="page-10-1"></span>

**Figure 2.** Frequencies of malaria detection/diagnosis methods in reviewed publications. **Figure 2.** Frequencies of malaria detection/diagnosis methods in reviewed publications.

### **4. Discussion**

Critical to achieving effective control, treatment, and subsequent elimination of malaria is the timely detection of the parasitic infection. In the face of this threatening infection, continuous progress and innovative research are required, which leads to the development of new tools that will be useful in the fight against malaria [\[117\]](#page-18-10). This article reviewed the recent developments in malaria diagnostic methods and their potential for point-ofcare and personalized malaria care, with special emphasis on the use of these methods in economically challenged countries.

The findings from this review suggest great advancement recently in malaria diagnostics. Research efforts by many scientists around the globe have progressed from developing improved malaria microscopy techniques into enhanced and more accurate molecular, immunological, computerized, digital methods of detection, automated analyzers, and point-of-care devices. Studies suggest that the influence of the old age infection on global health outcomes has urged on the design of more efficient diagnostics, with efforts directed at the development of point-of-care devices useful for resource-limited areas [\[7\]](#page-13-6). For an active drive towards the elimination of malaria, an early detection approach capable of revealing low levels of the parasitic infection is imperative [\[3\]](#page-13-2).

As observed in Tables [2–](#page-4-0)[5,](#page-10-0) the outcome of this review indicates that recent malaria detection methods actively being used or investigated include traditional methods, molecular techniques with polymerase chain reaction (PCR), loop-mediated isothermal amplification (LAMP)-based assays, and machine learning/computerized techniques (that exploit the physical and/or biological properties of Plasmodium-infected erythrocytes to enhance malaria diagnosis), among others. Figure [2](#page-10-1) shows the frequencies of detection methods as identified from various articles published over the last decade. Several other technologies and chemical assays are also being designed to tackle the malaria burden. RDTs were among the commonly used or researched modern methods in resource-limited settings, as seen in Table [5](#page-10-0) and Figure [2.](#page-10-1) This is not surprising since they are relatively low cost and easy to use.

Studies confirm PCR-based techniques as having widespread use globally as they are highly sensitive and capable of detecting very low parasitemia levels [\[3,](#page-13-2)[22,](#page-14-28)[35\]](#page-14-27). Polymerase chain reaction basically makes use of DNA extracted from whole blood or other samples. The process continues with denaturation, amplification, and elongation steps, after which the sensitivity and specificity of the assay can be assessed [\[35\]](#page-14-27). It is proposed that the PCR method, under equal reaction parameters, can diagnose all five species of the plasmodium parasitic infection [\[35\]](#page-14-27). Our findings reveal that a wide range of PCR assays have been developed or used over the past decade, which are less laborious and provide much faster and more accurate results [\[22\]](#page-14-28). Furthermore, PCR-based assays are widely preferred due to several reasons, including simultaneous species-specific detection and quantification, higher sensitivity, higher specificity, less time consuming, easy to use, and capable of diagnosing subclinical infections [\[13,](#page-13-12)[18,](#page-14-0)[22,](#page-14-28)[23](#page-14-24)[,41](#page-15-10)[,42\]](#page-15-11).

Though PCR is an effective approach to malaria detection, it is limited by the requirement of costly laboratory facilities and expertise and thus less beneficial to resource-limited areas and at the point of care [\[3\]](#page-13-2). Despite that, quite a number of studies in resource-limited settings, including some African countries, utilized PCR-based techniques, as shown in Table [5](#page-10-0) and Figure [2](#page-10-1) [\[13](#page-13-12)[,16](#page-14-23)[,18](#page-14-0)[,23](#page-14-24)[,25](#page-14-25)[,29](#page-14-26)[,35–](#page-14-27)[38](#page-15-5)[,40\]](#page-15-6). Other advanced PCR techniques, such as lab chip real-time PCR (LRP) and hair qPCR, were found to be suitable alternatives for point-of-care or resource-limited settings, though no evidence was found of the former currently being used or researched in lower or lower-middle-income countries [\[29,](#page-14-26)[31\]](#page-14-29). Gómez-Luque et al. proposed that due to limitations observed, more research is required to affirm the use of the hair qPCR as an efficient technique for malaria detection [\[29\]](#page-14-26). The one advantage the hair qPCR has over other PCR types is the use of non-invasive samples. LRP, however, being highly sensitive, specific, and less expensive will be beneficial for diagnosis and control in malaria-endemic countries [\[31\]](#page-14-29).

LAMP-based assays have also dominated research on malaria diagnostics. As seen in Table [3,](#page-5-0) studies have shown the development or use of various LAMP assays, which are effective malaria diagnostics [\[118\]](#page-18-11). Rei Yan et al. reviewed LAMP assays and found them easy to use in regions where there is limited access to clinical expertise and molecular biology equipment. Modified LAMP based assays such as multiplex LAMP with dipstick DNA chromatography, high throughput LAMP, 18S rRNA LAMP, mediated LAMP combined with lateral flow detection (LFD), etc., are highly sensitive, easy to use, consistent, convenient, cost effective, and useful in point-of-care situations [\[55](#page-15-12)[,56](#page-15-9)[,58](#page-16-15)[,59\]](#page-16-11), thus enabling an approach towards personalized healthcare. Table [5](#page-10-0) provides evidence of the development and use of LAMP techniques in lower and lower-middle-income countries, including countries in sub-Saharan Africa where malaria is endemic [\[45,](#page-15-7)[48,](#page-15-8)[56–](#page-15-9)[59\]](#page-16-11).

Other molecular methods worthy of note as they double as point-of-care or easyto-use methods include nuclear magnetic resonance (NMR)-based hemozoin detection, ultra-bright SERS nanorattles, recombinase-aided amplification with lateral flow dipstick assay, and dye-coupled aptamer-captured enzyme-catalyzed assay [\[86,](#page-17-21)[104,](#page-17-20)[105,](#page-18-12)[110](#page-18-13)[–112\]](#page-18-14). Though the latter two could be used in resource-limited settings due to their low cost, the study found only a dye-coupled aptamer-captured enzyme-catalyzed assay used in India [\[104\]](#page-17-20). Veiga and Peng identified nuclear magnetic resonance (NMR)-based hemozoin detection as having the potential of enabling personalized malaria medicine (that is, malaria treatment tailored to individual characteristics) with needleless diagnosis foresighted [\[119\]](#page-18-15). This technology may offer the detection of phenotypic variants, which are observable variations in characteristics among parasites of the same species as a result of genetic diversity, host–parasite interactions, or environmental factors, among others [\[120,](#page-18-16)[121\]](#page-18-17). For example, there are drug-resistant variants, those with surface antigen variations, and variants with different clinical presentations, among others [\[121–](#page-18-17)[123\]](#page-18-18). The ability to detect such variants would increase diagnostic accuracy and be considerably useful against parasite drug resistance. Acquiring these time- and patient-specific phenotypic identifiers is a basic step to personalized malaria medicine as variants continually rise [\[119\]](#page-18-15). The one advantage that phenotypic variant determination using NMR technology may have over nucleic acid amplification-based methods for genomic profiling is the extremely fast turnaround time for some of the devices [\[110\]](#page-18-13). Unfortunately, there was no evidence of such methods being used in lower and lower-middle-income countries as per the studied published data in the research articles reviewed.

Furthermore, a number of other technologies have emerged capable of point-of-care diagnosis. Unlike the traditional microscopy and commonly used RDTs, some of these methods were found to be highly sensitive, non-invasive as far as sample collection was concerned, and cost effective, even though there was no evidence that cost-effective ones were necessarily being used in economically challenged settings [\[85](#page-17-22)[,95,](#page-17-23)[99–](#page-17-24)[102\]](#page-17-25). In addition to these, Aggarwal et al. classify omics-based diagnostics as another important category to malaria diagnosis and elimination [\[124\]](#page-18-19). Multi-omics combines genomics, proteomics, metabolomics, phenomics, and transcriptomics in the investigation of biomarkers optimal for disease diagnosis and treatment. Though each omics has individual limitations, collectively, multi-omics can lead to a more comprehensive understanding of malaria infections, which can lead to more effective treatments [\[124\]](#page-18-19). In this review, the only struggling economy we found using multi-omics was India. No African nation was indicated.

#### **5. Conclusions**

Given the literature reviewed, there is adequate evidence to suggest that malaria detection or diagnosis will progress significantly in the next decade and beyond towards needleless detection. This advancement will however require increased, detailed, and specified research into the various molecular identifiers and phenotypic variant characteristics of malaria infection while enhancing the accuracy, precision, and specificity of the modernized point-of-care diagnostic tools. With this in view, precedence is duly set for the use of personalized medicine in the treatment of malaria infections. Notwithstanding, the

traditional thin and thick film microscopy and RDTs will continue to play an important role in the accurate detection of malaria infections, especially in resource-limited areas where there is less access to modernized diagnostic tools and little research into advanced malaria detection methods. It is, however, encouraging to see that PCR-based and LAMP-based tests were seen being utilized in these areas, including African countries. However, other modern molecular/point-of-care tests were not being utilized in sub-Saharan Africa. Findings of this study show that approximately a quarter (26%) of a total of eighty countries in low and lower-middle-income settings employ state-of-the-art methods for malaria diagnostics. This underscores the need for governments, non-governmental organizations, and funding bodies to intensify efforts towards malaria diagnostics and research in the fight against malaria.

**Author Contributions:** Conceptualization, A.K.Y.; methodology, A.K.Y., J.O. and J.E.C.; software, A.K.Y. and J.O.; validation, A.K.Y., J.O., J.E.C., N.I.N.-T., E.O., I.K.Y., A.A.K.-K., G.A., I.A.-B. and D.A.P.; formal analysis, A.K.Y. and J.O.; investigation, A.K.Y., J.O. and J.E.C.; resources, A.K.Y., J.O., J.E.C., N.I.N.-T., E.O., I.K.Y., A.A.K.-K., G.A., I.A.-B. and D.A.P.; writing—original draft preparation, A.K.Y., J.O. and J.E.C.; writing—review and editing, A.K.Y., J.O., J.E.C., N.I.N.-T., E.O., I.K.Y., A.A.K.-K., G.A., I.A.-B. and D.A.P.; funding acquisition, A.K.Y., N.I.N.-T., E.O., I.K.Y., A.A.K.-K., G.A., I.A.-B. and D.A.P. All authors have read and agreed to the published version of the manuscript.

**Funding:** This review work received no external funding.

**Institutional Review Board Statement:** Not applicable.

**Informed Consent Statement:** Not applicable.

**Data Availability Statement:** Not applicable.

**Conflicts of Interest:** The authors declare no conflicts of interest.

### **References**

- <span id="page-13-0"></span>1. WHO. World Malaria Report. 2023. Available online: <https://www.who.int/publications/i/item/9789240086173> (accessed on 19 July 2024).
- <span id="page-13-13"></span><span id="page-13-1"></span>2. Tetteh-Quarcoo, P.B.; Dayie, N.T.; Adutwum-Ofosu, K.K.; Ahenkorah, J.; Afutu, E.; Amponsah, S.K.; Abdul-Rahman, M.; Kretchy, J.-P.; Ocloo, J.Y.; Nii-Trebi, N.I. Unravelling the perspectives of day and night traders in selected markets within a sub-saharan african city with a malaria knowledge, attitude and practice survey. *Int. J. Environ. Res. Public Health* **2021**, *18*, 3468. [\[CrossRef\]](https://doi.org/10.3390/ijerph18073468)
- <span id="page-13-2"></span>3. Zheng, Z.; Cheng, Z. Advances in molecular diagnosis of malaria. *Adv. Clin. Chem.* **2017**, *80*, 155–192.
- <span id="page-13-3"></span>4. malERA Consultative Group on Diagnoses and Diagnostics. A research agenda for malaria eradication: Diagnoses and diagnostics. *PLoS Med.* **2011**, *8*, e1000396.
- <span id="page-13-4"></span>5. Yalley, A.K.; Ahiatrogah, S.; Kafintu-Kwashie, A.A.; Amegatcher, G.; Prah, D.; Botwe, A.K.; Adusei-Poku, M.A.; Obodai, E.; Nii-Trebi, N.I. A systematic review on suitability of molecular techniques for diagnosis and research into infectious diseases of concern in resource-limited settings. *Curr. Issues Mol. Biol.* **2022**, *44*, 4367–4385. [\[CrossRef\]](https://doi.org/10.3390/cimb44100300) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/36286015)
- <span id="page-13-5"></span>6. WHO. Global Malaria Program. 2024. Available online: [https://www.who.int/teams/global-malaria-programme/case](https://www.who.int/teams/global-malaria-programme/case-management/diagnosis/microscopy)[management/diagnosis/microscopy](https://www.who.int/teams/global-malaria-programme/case-management/diagnosis/microscopy) (accessed on 10 June 2024).
- <span id="page-13-6"></span>7. Krampa, F.D.; Aniweh, Y.; Awandare, G.A.; Kanyong, P. Recent progress in the development of diagnostic tests for malaria. *Diagnostics* **2017**, *7*, 54. [\[CrossRef\]](https://doi.org/10.3390/diagnostics7030054) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28925968)
- <span id="page-13-7"></span>8. Singh, K.; Bharti, P.K.; Devi, N.C.; Ahmed, N.; Sharma, A. *Plasmodium malariae* Detected by Microscopy in the International Bordering Area of Mizoram, a Northeastern State of India. *Diagnostics* **2022**, *12*, 2015. [\[CrossRef\]](https://doi.org/10.3390/diagnostics12082015)
- <span id="page-13-8"></span>9. Edwards, H. Tales from the bench: Laboratory diagnosis of malaria. *Trop. Dr.* **2011**, *41*, 106–107. [\[CrossRef\]](https://doi.org/10.1258/td.2010.100199) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/21071486)
- <span id="page-13-9"></span>10. Worldbank. World Bank Country and Lending Groups. 2024. Available online: [https://datahelpdesk.worldbank.org/](https://datahelpdesk.worldbank.org/knowledgebase/articles/906519-world-bank-country-and-lending-groups) [knowledgebase/articles/906519-world-bank-country-and-lending-groups](https://datahelpdesk.worldbank.org/knowledgebase/articles/906519-world-bank-country-and-lending-groups) (accessed on 13 May 2024).
- <span id="page-13-10"></span>11. Azikiwe, C.C.; Ifezulike, C.; Siminialayi, I.; Amazu, L.U.; Enye, J.; Nwakwunite, O. A comparative laboratory diagnosis of malaria: Microscopy versus rapid diagnostic test kits. *Asian Pac. J. Trop. Biomed.* **2012**, *2*, 307–310. [\[CrossRef\]](https://doi.org/10.1016/S2221-1691(12)60029-X) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/23569920)
- <span id="page-13-11"></span>12. Iqbal, J.; Hira, P.R.; Al-Ali, F.; Khalid, N.; Sher, A. Modified Giemsa staining for rapid diagnosis of malaria infection. *Med. Princ. Pract.* **2003**, *12*, 156–159. [\[CrossRef\]](https://doi.org/10.1159/000070751)
- <span id="page-13-12"></span>13. Nadeem, M.F.; Khattak, A.A.; Yaqoob, A.; Awan, U.A.; Zeeshan, N. Assessment of Microscopic detection of Malaria with Nested Polymerase Chain Reaction in War-torn Federally Administered Tribal Areas of Pakistan. *Acta Parasitol.* **2021**, *66*, 1186–1192. [\[CrossRef\]](https://doi.org/10.1007/s11686-021-00374-8)
- <span id="page-14-20"></span><span id="page-14-19"></span><span id="page-14-18"></span><span id="page-14-10"></span><span id="page-14-9"></span><span id="page-14-5"></span><span id="page-14-4"></span><span id="page-14-3"></span>14. Awosolu, O.B.; Yahaya, Z.S.; Farah Haziqah, M.T. Efficacy of *Plasmodium falciparum* histidine-rich protein 2 (Pfhrp 2) rapid diagnostic test (RDT) and microscopy in the detection of falciparum malaria among symptomatic patients in Akure, Nigeria. *Trop. Biomed.* **2022**, *39*, 144–149. [\[CrossRef\]](https://doi.org/10.47665/tb.39.1.019) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/35507937)
- <span id="page-14-21"></span>15. Badiane, A.; Thwing, J.; Williamson, J.; Rogier, E.; Diallo, M.A.; Ndiaye, D. Sensitivity and specificity for malaria classification of febrile persons by rapid diagnostic test, microscopy, parasite DNA, histidine-rich protein 2, and IgG: Dakar, Senegal 2015. *Int. J. Infect. Dis.* **2022**, *121*, 92–97. [\[CrossRef\]](https://doi.org/10.1016/j.ijid.2022.04.060)
- <span id="page-14-23"></span><span id="page-14-6"></span>16. Fontecha, G.; Escobar, D.; Ortiz, B.; Pinto, A.; Serrano, D.; Valdivia, H.O. Field Evaluation of a Hemozoin-Based Malaria Diagnostic Device in Puerto Lempira, Honduras. *Diagnostics* **2022**, *12*, 1206. [\[CrossRef\]](https://doi.org/10.3390/diagnostics12051206) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/35626361)
- <span id="page-14-22"></span><span id="page-14-7"></span>17. Ahmad, A.; Soni, P.; Kumar, L.; Singh, M.P.; Verma, A.K.; Sharma, A.; Das, A.; Bharti, P.K. Comparison of polymerase chain reaction, microscopy, and rapid diagnostic test in malaria detection in a high burden state (Odisha) of India. *Pathog. Glob. Health* **2021**, *115*, 267–272. [\[CrossRef\]](https://doi.org/10.1080/20477724.2021.1893484) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/33634745)
- <span id="page-14-8"></span><span id="page-14-0"></span>18. Ugah, U.I.; Alo, M.N.; Owolabi, J.O.; Okata-Nwali, O.D.G.; Ekejindu, I.M.; Ibeh, N.; Elom, M.O. Evaluation of the utility value of three diagnostic methods in the detection of malaria parasites in endemic area. *Malar. J.* **2017**, *16*, 189. [\[CrossRef\]](https://doi.org/10.1186/s12936-017-1838-4) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28477621)
- <span id="page-14-11"></span><span id="page-14-1"></span>19. Karimi, A.; Navidbakhsh, M.; Haghi, A.M.; Faghihi, S. A morphology-based method for the diagnosis of red blood cells parasitized by *Plasmodium malariae* and *Plasmodium ovale*. *Scand. J. Infect. Dis.* **2014**, *46*, 368–375. [\[CrossRef\]](https://doi.org/10.3109/00365548.2014.880186) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/24568595)
- <span id="page-14-2"></span>20. Mohanty, S.; Sharma, R.; Deb, M. Usefulness of a centrifuged buffy coat smear examination for diagnosis of malaria. *Indian J. Med. Microbiol.* **2015**, *33*, 63–67. [\[CrossRef\]](https://doi.org/10.4103/0255-0857.148380) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/25560003)
- <span id="page-14-12"></span>21. Echeverry, D.F.; Deason, N.A.; Davidson, J.; Makuru, V.; Xiao, H.; Niedbalski, J.; Kern, M.; Russell, T.L.; Burkot, T.R.; Collins, F.H.; et al. Human malaria diagnosis using a single-step direct-PCR based on the Plasmodium cytochrome oxidase III gene. *Malar. J.* **2016**, *15*, 128. [\[CrossRef\]](https://doi.org/10.1186/s12936-016-1185-x)
- <span id="page-14-28"></span><span id="page-14-14"></span><span id="page-14-13"></span>22. Schneider, R.; Lamien-Meda, A.; Auer, H.; Wiedermann-Schmidt, U.; Chiodini, P.L.; Walochnik, J. Validation of a novel FRET real-time PCR assay for simultaneous quantitative detection and discrimination of human *Plasmodium* parasites. *PLoS ONE* **2021**, *16*, e0252887. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0252887) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/34086817)
- <span id="page-14-24"></span><span id="page-14-15"></span>23. Ranjan, P.; Ghoshal, U. Utility of nested polymerase chain reaction over the microscopy and immuno-chromatographic test in the detection of *Plasmodium* species and their clinical spectrum. *Parasitol. Res.* **2016**, *115*, 3375–3385. [\[CrossRef\]](https://doi.org/10.1007/s00436-016-5098-y) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/27147091)
- <span id="page-14-16"></span>24. Freitas, D.R.C.d.; Gomes, L.T.; Fontes, C.J.F.; Tauil, P.L.; Pang, L.W.; Duarte, E.C. Sensitivity of nested-PCR for *Plasmodium* detection in pooled whole blood samples and its usefulness to blood donor screening in endemic areas. *Transfus. Apher. Sci.* **2014**, *50*, 242–246. [\[CrossRef\]](https://doi.org/10.1016/j.transci.2014.01.016) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/24508148)
- <span id="page-14-25"></span>25. Li, P.; Zhao, Z.; Wang, Y.; Xing, H.; Parker, D.M.; Yang, Z.; Baum, E.; Li, W.; Sattabongkot, J.; Sirichaisinthop, J.; et al. Nested PCR detection of malaria directly using blood filter paper samples from epidemiological surveys. *Malar. J.* **2014**, *13*, 175. [\[CrossRef\]](https://doi.org/10.1186/1475-2875-13-175)
- <span id="page-14-17"></span>26. Pomari, E.; Silva, R.; Moro, L.; Marca, G.L.; Perandin, F.; Verra, F.; Bisoffi, Z.; Piubelli, C. Droplet digital PCR for the detection of *Plasmodium falciparum* DNA in whole blood and serum: A comparative analysis with other molecular methods. *Pathogens* **2020**, *9*, 478. [\[CrossRef\]](https://doi.org/10.3390/pathogens9060478) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32560386)
- 27. Srisutham, S.; Saralamba, N.; Malleret, B.; Rénia, L.; Dondorp, A.M.; Imwong, M. Four human *Plasmodium* species quantification using droplet digital PCR. *PLoS ONE* **2017**, *12*, e0175771. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0175771) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28423028)
- 28. Xu, W.; Morris, U.; Aydin-Schmidt, B.; Msellem, M.I.; Shakely, D.; Petzold, M.; Björkman, A.; Mårtensson, A. SYBR green real-time PCR-RFLP assay targeting the *Plasmodium* cytochrome B gene—A highly sensitive molecular tool for malaria parasite detection and species determination. *PLoS ONE* **2015**, *10*, e0120210. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0120210) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/25774805)
- <span id="page-14-26"></span>29. Gómez-Luque, A.; Parejo, J.C.; Clavijo-Chamorro, M.Z.; López-Espuela, F.; Munyaruguru, F.; Lorenzo, S.B.; Monroy, I.; Gómez-Nieto, L.C. Method for malaria diagnosis based on extractions of samples using non-invasive techniques: An opportunity for the nursing clinical practice. *Int. J. Environ. Res. Public Health* **2020**, *17*, 5551. [\[CrossRef\]](https://doi.org/10.3390/ijerph17155551)
- 30. Chua, K.H.; Lee, P.C.; Chai, H.C. Development of insulated isothermal PCR for rapid on-site malaria detection. *Malar. J.* **2016**, *15*, 134. [\[CrossRef\]](https://doi.org/10.1186/s12936-016-1183-z)
- <span id="page-14-29"></span>31. Kim, J.; Lim, D.H.; Mihn, D.-C.; Nam, J.; Jang, W.S.; Lim, C.S. Clinical usefulness of labchip real-time PCR using lab-on-a-chip technology for diagnosing malaria. *Korean J. Parasitol.* **2021**, *59*, 77. [\[CrossRef\]](https://doi.org/10.3347/kjp.2021.59.1.77)
- 32. Barbosa, L.R.; da Silva, E.L.; de Almeida, A.C.; Salazar, Y.E.; Siqueira, A.M.; Alecrim, M.d.G.C.; Vieira, J.L.F.; Bassat, Q.; de Lacerda, M.V.; Monteiro, W.M. An Ultra-Sensitive Technique: Using Pv-mtCOX1 qPCR to Detect Early Recurrences of *Plasmodium vivax* in Patients in the Brazilian Amazon. *Pathogens* **2020**, *10*, 19. [\[CrossRef\]](https://doi.org/10.3390/pathogens10010019)
- 33. Obaldía, N.; Barahona, I.; Lasso, J.; Avila, M.; Quijada, M.; Nuñez, M.; Marti, M. Comparison of *PvLAP5* and *Pvs25* qRT-PCR assays for the detection of *Plasmodium vivax* gametocytes in field samples preserved at ambient temperature from remote malaria endemic regions of Panama. *PLoS Neglected Trop. Dis.* **2022**, *16*, e0010327. [\[CrossRef\]](https://doi.org/10.1371/journal.pntd.0010327) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/35394999)
- 34. Bouzayene, A.; Zaffaroullah, R.; Bailly, J.; Ciceron, L.; Sarrasin, V.; Cojean, S.; Argy, N.; Houzé, S.; Joste, V.; Angoulvant, A.; et al. Evaluation of two commercial kits and two laboratory-developed qPCR assays compared to LAMP for molecular diagnosis of malaria. *Malar. J.* **2022**, *21*, 204. [\[CrossRef\]](https://doi.org/10.1186/s12936-022-04219-1) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/35761324)
- <span id="page-14-27"></span>35. Sazed, S.A.; Kibria, M.G.; Alam, M.S. An optimized real-time qpcr method for the effective detection of human malaria infections. *Diagnostics* **2021**, *11*, 736. [\[CrossRef\]](https://doi.org/10.3390/diagnostics11050736)
- 36. Grignard, L.; Nolder, D.; Sepúlveda, N.; Berhane, A.; Mihreteab, S.; Kaaya, R.; Phelan, J.; Moser, K.; van Schalkwyk, D.A.; Campino, S.; et al. A novel multiplex qPCR assay for detection of *Plasmodium falciparum* with histidine-rich protein 2 and 3 (pfhrp2 and pfhrp3) deletions in polyclonal infections. *EBioMedicine* **2020**, *55*, 102757. [\[CrossRef\]](https://doi.org/10.1016/j.ebiom.2020.102757)
- <span id="page-15-1"></span><span id="page-15-0"></span>37. Aimeé, K.K.; Lengu, T.B.; Nsibu, C.N.; Umesumbu, S.E.; Ngoyi, D.M.; Chen, T. Molecular detection and species identification of *Plasmodium* spp. infection in adults in the Democratic Republic of Congo: A populationbased study. *PLoS ONE* **2020**, *15*, e0242713. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0242713)
- <span id="page-15-5"></span><span id="page-15-2"></span>38. Canier, L.; Khim, N.; Kim, S.; Eam, R.; Khean, C.; Loch, K.; Ken, M.; Pannus, P.; Bosman, P.; Stassijns, J.; et al. Malaria PCR detection in Cambodian low-transmission settings: Dried blood spots versus venous blood samples. *Am. Soc. Trop. Med. Hyg.* **2015**, *92*, 573–577. [\[CrossRef\]](https://doi.org/10.4269/ajtmh.14-0614)
- <span id="page-15-3"></span>39. Phuong, M.; Lau, R.; Ralevski, F.; Boggild, A.K. Sequence-based optimization of a quantitative real-time PCR assay for detection of *Plasmodium ovale* and *Plasmodium malariae*. *J. Clin. Microbiol.* **2014**, *52*, 1068–1073. [\[CrossRef\]](https://doi.org/10.1128/JCM.03477-13)
- <span id="page-15-6"></span><span id="page-15-4"></span>40. Leski, T.A.; Taitt, C.R.; Swaray, A.G.; Bangura, U.; Reynolds, N.D.; Holtz, A.; Yasuda, C.; Lahai, J.; Lamin, J.M.; Baio, V.; et al. Use of real-time multiplex PCR, malaria rapid diagnostic test and microscopy to investigate the prevalence of *Plasmodium* species among febrile hospital patients in Sierra Leone. *Malar. J.* **2020**, *19*, 84. [\[CrossRef\]](https://doi.org/10.1186/s12936-020-03163-2) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32085711)
- <span id="page-15-10"></span>41. Murillo, E.; Muskus, C.; Agudelo, L.A.; Vélez, I.D.; Ruiz-Lopez, F. A new high-resolution melting analysis for the detection and identification of *Plasmodium* in human and Anopheles vectors of malaria. *Sci. Rep.* **2019**, *9*, 1674. [\[CrossRef\]](https://doi.org/10.1038/s41598-018-36515-9)
- <span id="page-15-11"></span>42. Amaral, L.C.; Robortella, D.R.; Guimarães, L.F.F.; Limongi, J.E.; Fontes, C.J.F.; Pereira, D.B.; De Brito, C.F.A.; Kano, F.S.; De Sousa, T.N.; Carvalho, L.H. Ribosomal and non-ribosomal PCR targets for the detection of low-density and mixed malaria infections. *Malar. J.* **2019**, *18*, 154. [\[CrossRef\]](https://doi.org/10.1186/s12936-019-2781-3)
- 43. Frickmann, H.; Wegner, C.; Ruben, S.; Behrens, C.; Kollenda, H.; Hinz, R.; Rojak, S.; Schwarz, N.G.; Hagen, R.M.; Tannich, E. Evaluation of the multiplex real-time PCR assays RealStar malaria S&T PCR kit 1.0 and FTD malaria differentiation for the differentiation of *Plasmodium* species in clinical samples. *Travel Med. Infect. Dis.* **2019**, *31*, 101442. [\[CrossRef\]](https://doi.org/10.1016/j.tmaid.2019.06.013) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31255712)
- 44. Lee, P.C.; Chong, E.T.J.; Anderios, F.; Lim, Y.A.L.; Chew, C.H.; Chua, K.H. Molecular detection of human *Plasmodium* species in Sabah using PlasmoNex™ multiplex PCR and hydrolysis probes real-time PCR. *Malar. J.* **2015**, *14*, 28. [\[CrossRef\]](https://doi.org/10.1186/s12936-015-0542-5)
- <span id="page-15-7"></span>45. Hayashida, K.; Kajino, K.; Simukoko, H.; Simuunza, M.; Ndebe, J.; Chota, A.; Namangala, B.; Sugimoto, C. Direct detection of *falciparum* and non-*falciparum* malaria DNA from a drop of blood with high sensitivity by the dried-LAMP system. *Parasites Vectors* **2017**, *10*, 26. [\[CrossRef\]](https://doi.org/10.1186/s13071-016-1949-8)
- 46. Colbert, A.J.; Co, K.; Lima-Cooper, G.; Lee, D.H.; Clayton, K.N.; Wereley, S.T.; John, C.C.; Linnes, J.C.; Kinzer-Ursem, T.L. Towards the use of a smartphone imaging-based tool for point-of-care detection of asymptomatic low-density malaria parasitaemia. *Malar. J.* **2021**, *20*, 380. [\[CrossRef\]](https://doi.org/10.1186/s12936-021-03894-w) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/34563189)
- 47. Imai, K.; Tarumoto, N.; Misawa, K.; Runtuwene, L.R.; Sakai, J.; Hayashida, K.; Eshita, Y.; Maeda, R.; Tuda, J.; Murakami, T.; et al. A novel diagnostic method for malaria using loop-mediated isothermal amplification (LAMP) and MinION™ nanopore sequencer. *BMC Infect. Dis.* **2017**, *17*, 621. [\[CrossRef\]](https://doi.org/10.1186/s12879-017-2718-9) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28903726)
- <span id="page-15-8"></span>48. Azam, M.; Upmanyu, K.; Gupta, R.; Sruthy, K.S.; Matlani, M.; Savargaonkar, D.; Singh, R. Development of two-tube loop-mediated isothermal amplification assay for differential diagnosis of *Plasmodium falciparum* and *Plasmodium vivax* and its comparison with loopamp™ malaria. *Diagnostics* **2021**, *11*, 1689. [\[CrossRef\]](https://doi.org/10.3390/diagnostics11091689) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/34574030)
- 49. Jang, W.S.; Lim, D.H.; Choe, Y.; Jee, H.; Moon, K.C.; Kim, C.; Choi, M.; Park, I.S.; Lim, C.S. Development of a multiplex loop-mediated isothermal amplification assay for diagnosis of *Plasmodium* spp., *Plasmodium falciparum* and *Plasmodium vivax*. *Diagnostics* **2021**, *11*, 1950. [\[CrossRef\]](https://doi.org/10.3390/diagnostics11111950)
- 50. Mohon, A.N.; Getie, S.; Jahan, N.; Alam, M.S.; Pillai, D.R. Ultrasensitive loop mediated isothermal amplification (US-LAMP) to detect malaria for elimination. *Malar. J.* **2019**, *18*, 350. [\[CrossRef\]](https://doi.org/10.1186/s12936-019-2979-4)
- 51. Viana, G.M.R.; Silva-Flannery, L.; Barbosa, D.R.L.; Lucchi, N.; do Valle, S.C.N.; Farias, S.; Barbalho, N.; Marchesini, P.; Rossi, J.C.N.; Udhayakumar, V.; et al. Field evaluation of a real time loop-mediated isothermal amplification assay (RealAmp) for malaria diagnosis in Cruzeiro do Sul, Acre, Brazil. *PLoS ONE* **2018**, *13*, e0200492. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0200492) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/29995953)
- 52. Cuadros, J.; Martin Ramírez, A.; González, I.J.; Ding, X.C.; Perez Tanoira, R.; Rojo-Marcos, G.; Gómez-Herruz, P.; Rubio, J.M. LAMP kit for diagnosis of non-falciparum malaria in *Plasmodium ovale* infected patients. *Malar. J.* **2017**, *16*, 20. [\[CrossRef\]](https://doi.org/10.1186/s12936-016-1669-8) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28061871)
- 53. Lau, Y.L.; Lai, M.Y.; Fong, M.Y.; Jelip, J.; Mahmud, R. Loop-mediated isothermal amplification assay for identification of five human *Plasmodium* species in Malaysia. *Am. J. Trop. Med. Hyg.* **2016**, *94*, 336–339. [\[CrossRef\]](https://doi.org/10.4269/ajtmh.15-0569)
- 54. Patel, J.C.; Lucchi, N.W.; Srivastava, P.; Lin, J.T.; Sug-Aram, R.; Aruncharus, S.; Bharti, P.K.; Shukla, M.M.; Congpuong, K.; Satimai, W.; et al. Field evaluation of a real-time fluorescence loop-mediated isothermal amplification assay, realamp, for the diagnosis of Malaria in Thailand and India. *J. Infect. Dis.* **2014**, *210*, 1180–1187. [\[CrossRef\]](https://doi.org/10.1093/infdis/jiu252) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/24795480)
- <span id="page-15-12"></span>55. Moonga, L.C.; Hayashida, K.; Kawai, N.; Nakao, R.; Sugimoto, C.; Namangala, B.; Yamagishi, J. Development of a multiplex loop-mediated isothermal amplification (LAMP) method for simultaneous detection of spotted fever group rickettsiae and malaria parasites by dipstick DNA chromatography. *Diagnostics* **2020**, *10*, 897. [\[CrossRef\]](https://doi.org/10.3390/diagnostics10110897)
- <span id="page-15-9"></span>56. Aydin-Schmidt, B.; Morris, U.; Ding, X.C.; Jovel, I.; Msellem, M.I.; Bergman, D.; Islam, A.; Ali, A.S.; Polley, S.; Gonzalez, I.J.; et al. Field evaluation of a high throughput loop mediated isothermal amplification test for the detection of asymptomatic *Plasmodium* infections in Zanzibar. *PLoS ONE* **2017**, *12*, e0169037. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0169037) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/28095434)
- 57. Lucchi, N.W.; Gaye, M.; Diallo, M.A.; Goldman, I.F.; Ljolje, D.; Deme, A.B.; Badiane, A.; Ndiaye, Y.D.; Barnwell, J.W.; Udhayakumar, V.; et al. Evaluation of the *Illumigene* Malaria LAMP: A Robust Molecular Diagnostic Tool for Malaria Parasites. *Sci. Rep.* **2016**, *6*, 36808. [\[CrossRef\]](https://doi.org/10.1038/srep36808) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/27827432)
- <span id="page-16-15"></span><span id="page-16-2"></span><span id="page-16-1"></span><span id="page-16-0"></span>58. Sharma, S.; Kumar, S.; Ahmed, M.Z.; Bhardwaj, N.; Singh, J.; Kumari, S.; Savargaonkar, D.; Anvikar, A.R.; Das, J. Advanced multiplex loop mediated isothermal amplification (mLAMP) combined with lateral flow detection (LFD) for rapid detection of two prevalent malaria species in india and melting curve analysis. *Diagnostics* **2022**, *12*, 32. [\[CrossRef\]](https://doi.org/10.3390/diagnostics12010032)
- <span id="page-16-11"></span>59. Aninagyei, E.; Boakye, A.A.; Tettey, C.O.; Ntiri, K.A.; Ofori, S.O.; Tetteh, C.D.; Aphour, T.T.; Rufai, T. Utilization of 18s ribosomal RNA LAMP for detecting *Plasmodium falciparum* in microscopy and rapid diagnostic test negative patients. *PLoS ONE* **2022**, *17*, e0275052. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0275052) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/36201568)
- 60. Lai, M.Y.; Ooi, C.H.; Jaimin, J.J.; Lau, Y.L. Evaluation of WarmStart colorimetric loop-mediated isothermal amplification assay for diagnosis of Malaria. *Am. J. Trop. Med. Hyg.* **2020**, *102*, 1370–1372. [\[CrossRef\]](https://doi.org/10.4269/ajtmh.20-0001)
- 61. Barazorda, K.A.; Salas, C.J.; Braga, G.; Ricopa, L.; Ampuero, J.S.; Siles, C.; Sanchez, J.F.; Montano, S.; Lizewski, S.E.; Joya, C.A.; et al. Validation study of Boil & Spin Malachite Green Loop Mediated Isothermal Amplification (B&S MG-LAMP) versus microscopy for malaria detection in the Peruvian Amazon. *PLoS ONE* **2021**, *16*, e0258722. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0258722)
- 62. Vincent, J.P.; Komaki-Yasuda, K.; Iwagami, M.; Kawai, S.; Kano, S. Combination of PURE-DNA extraction and LAMP-DNA amplification methods for accurate malaria diagnosis on dried blood spots 11 Medical and Health Sciences 1108 Medical Microbiology. *Malar. J.* **2018**, *17*, 373. [\[CrossRef\]](https://doi.org/10.1186/s12936-018-2527-7)
- <span id="page-16-3"></span>63. Cordray, M.S.; Richards-Kortum, R.R. A paper and plastic device for the combined isothermal amplification and lateral flow detection of *Plasmodium* DNA. *Malar. J.* **2015**, *14*, 472. [\[CrossRef\]](https://doi.org/10.1186/s12936-015-0995-6) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/26611141)
- <span id="page-16-6"></span><span id="page-16-4"></span>64. Aninagyei, E.; Abraham, J.; Atiiga, P.; Antwi, S.D.; Bamfo, S.; Acheampong, D.O. Evaluating the potential of using urine and saliva specimens for malaria diagnosis in suspected patients in Ghana. *Malar. J.* **2020**, *19*, 349. [\[CrossRef\]](https://doi.org/10.1186/s12936-020-03427-x)
- <span id="page-16-7"></span>65. Turnbull, L.B.; Ayodo, G.; Knight, V.; John, C.C.; McHenry, M.S.; Tran, T.M. Evaluation of an ultrasensitive HRP2–based rapid diagnostic test for detection of asymptomatic *Plasmodium falciparum* parasitaemia among children in western Kenya. *Malar. J.* **2022**, *21*, 337. [\[CrossRef\]](https://doi.org/10.1186/s12936-022-04351-y)
- <span id="page-16-8"></span>66. Briand, V.; Cottrell, G.; Tuike Ndam, N.; Martiáñez-Vendrell, X.; Vianou, B.; Mama, A.; Kouwaye, B.; Houzé, S.; Bailly, J.; Gbaguidi, E.; et al. Prevalence and clinical impact of malaria infections detected with a highly sensitive HRP2 rapid diagnostic test in Beninese pregnant women. *Malar. J.* **2020**, *19*, 188. [\[CrossRef\]](https://doi.org/10.1186/s12936-020-03261-1)
- 67. Wardhani, P.; Butarbutar, T.V.; Adiatmaja, C.O.; Betaubun, A.M.; Hamidah, N.; Aryati. Performance comparison of two malaria rapid diagnostic test with real time polymerase chain reaction and gold standard of microscopy detection method. *Infect. Dis. Rep.* **2020**, *12*, 8731. [\[CrossRef\]](https://doi.org/10.4081/idr.2020.8731) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32874462)
- <span id="page-16-9"></span>68. Naeem, M.A.; Ahmed, S.; Khan, S.A. Detection of asymptomatic carriers of malaria in Kohat district of Pakistan. *Malar. J.* **2018**, *17*, 44. [\[CrossRef\]](https://doi.org/10.1186/s12936-018-2191-y) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/29357890)
- <span id="page-16-10"></span>69. Maltha, J.; Guiraud, I.; Lompo, P.; Kaboré, B.; Gillet, P.; Van Geet, C.; Tinto, H.; Jacobs, J. Accuracy of PfHRP2 versus Pf-pLDH antigen detection by malaria rapid diagnostic tests in hospitalized children in a seasonal hyperendemic malaria transmission area in Burkina Faso. *Malar. J.* **2014**, *13*, 20. [\[CrossRef\]](https://doi.org/10.1186/1475-2875-13-20) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/24418119)
- 70. Fedele, P.L.; Wheeler, M.; Lemoh, C.; Chunilal, S. Immunochromatographic antigen testing alone is sufficient to identify asymptomatic refugees at risk of severe malaria presenting to a single health service in Victoria. *Pathology* **2014**, *46*, 551–554. [\[CrossRef\]](https://doi.org/10.1097/PAT.0000000000000149) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/25158813)
- <span id="page-16-5"></span>71. Kim, J.; Cao, X.E.; Finkelstein, J.L.; Cárdenas, W.B.; Erickson, D.; Mehta, S. A two-colour multiplexed lateral flow immunoassay system to differentially detect human malaria species on a single test line. *Malar. J.* **2019**, *18*, 313. [\[CrossRef\]](https://doi.org/10.1186/s12936-019-2957-x) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31533756)
- 72. Abubakar, A.; Ajuji, M.; Yahya, I.U. Deepfmd: Computational analysis for malaria detection in blood-smear images using deep-learning features. *Appl. Syst. Innov.* **2021**, *4*, 82. [\[CrossRef\]](https://doi.org/10.3390/asi4040082)
- 73. Kassim, Y.M.; Palaniappan, K.; Yang, F.; Poostchi, M.; Palaniappan, N.; Maude, R.J.; Antani, S.; Jaeger, S. Clustering-Based Dual Deep Learning Architecture for Detecting Red Blood Cells in Malaria Diagnostic Smears. *IEEE J. Biomed. Health Inform.* **2021**, *25*, 1735–1746. [\[CrossRef\]](https://doi.org/10.1109/JBHI.2020.3034863)
- 74. Sriporn, K.; Tsai, C.F.; Tsai, C.E.; Wang, P. Analyzing malaria disease using effective deep learning approach. *Diagnostics* **2020**, *10*, 744. [\[CrossRef\]](https://doi.org/10.3390/diagnostics10100744)
- <span id="page-16-12"></span>75. Nakasi, R.; Mwebaze, E.; Zawedde, A. Mobile-aware deep learning algorithms for malaria parasites and white blood cells localization in thick blood smears. *Algorithms* **2021**, *14*, 17. [\[CrossRef\]](https://doi.org/10.3390/a14010017)
- 76. Yang, F.; Poostchi, M.; Yu, H.; Zhou, Z.; Silamut, K.; Yu, J.; Maude, R.J.; Jaeger, S.; Antani, S. Deep Learning for Smartphone-Based Malaria Parasite Detection in Thick Blood Smears. *IEEE J. Biomed. Health Inform.* **2020**, *24*, 1427–1438. [\[CrossRef\]](https://doi.org/10.1109/JBHI.2019.2939121) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31545747)
- <span id="page-16-13"></span>77. Manescu, P.; Shaw, M.J.; Elmi, M.; Neary-Zajiczek, L.; Claveau, R.; Pawar, V.; Kokkinos, I.; Oyinloye, G.; Bendkowski, C.; Oladejo, O.A.; et al. Expert-level automated malaria diagnosis on routine blood films with deep neural networks. *Am. J. Hematol.* **2020**, *95*, 883–891. [\[CrossRef\]](https://doi.org/10.1002/ajh.25827) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32282969)
- 78. Islam, M.R.; Nahiduzzaman, M.; Goni, M.O.F.; Sayeed, A.; Anower, M.S.; Ahsan, M.; Haider, J. Explainable Transformer-Based Deep Learning Model for the Detection of Malaria Parasites from Blood Cell Images. *Sensors* **2022**, *22*, 4358. [\[CrossRef\]](https://doi.org/10.3390/s22124358)
- 79. Abdurahman, F.; Fante, K.A.; Aliy, M. Malaria parasite detection in thick blood smear microscopic images using modified YOLOV3 and YOLOV4 models. *BMC Bioinform.* **2021**, *22*, 112. [\[CrossRef\]](https://doi.org/10.1186/s12859-021-04036-4)
- <span id="page-16-14"></span>80. Chibuta, S.; Acar, A.C. Real-time Malaria Parasite Screening in Thick Blood Smears for Low-Resource Setting. *J. Digit. Imaging* **2020**, *33*, 763–775. [\[CrossRef\]](https://doi.org/10.1007/s10278-019-00284-2) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31974686)
- 81. Ashraf, S.; Khalid, A.; de Vos, A.L.; Feng, Y.; Rohrbach, P.; Hasan, T. Malaria Detection Accelerated: Combing a High-Throughput NanoZoomer Platform with a ParasiteMacro Algorithm. *Pathogens* **2022**, *11*, 1182. [\[CrossRef\]](https://doi.org/10.3390/pathogens11101182) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/36297240)
- <span id="page-17-6"></span><span id="page-17-5"></span><span id="page-17-4"></span><span id="page-17-3"></span><span id="page-17-2"></span><span id="page-17-1"></span><span id="page-17-0"></span>82. Kongklad, G.; Chitaree, R.; Taechalertpaisarn, T.; Panvisavas, N.; Nuntawong, N. Discriminant Analysis PCA-LDA Assisted Surface-Enhanced Raman Spectroscopy for Direct Identification of Malaria-Infected Red Blood Cells. *Methods Protoc.* **2022**, *5*, 49. [\[CrossRef\]](https://doi.org/10.3390/mps5030049)
- 83. Wang, W.; Dong, R.L.; Gu, D.; He, J.A.; Yi, P.; Kong, S.K.; Ho, H.P.; Loo, J.F.C.; Wang, W.; Wang, Q. Antibody-free rapid diagnosis of malaria in whole blood with surface-enhanced Raman Spectroscopy using Nanostructured Gold Substrate. *Adv. Med. Sci.* **2020**, *65*, 86–92. [\[CrossRef\]](https://doi.org/10.1016/j.advms.2019.11.004)
- 84. Heraud, P.; Chatchawal, P.; Wongwattanakul, M.; Tippayawat, P.; Doerig, C.; Jearanaikoon, P.; Perez-Guaita, D.; Wood, B.R. Infrared spectroscopy coupled to cloud-based data management as a tool to diagnose malaria: A pilot study in a malaria-endemic country. *Malar. J.* **2019**, *18*, 348. [\[CrossRef\]](https://doi.org/10.1186/s12936-019-2945-1) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31619246)
- <span id="page-17-22"></span><span id="page-17-7"></span>85. McBirney, S.E.; Chen, D.; Scholtz, A.; Ameri, H.; Armani, A.M. Rapid Diagnostic for Point-of-Care Malaria Screening. *ACS Sens.* **2018**, *3*, 1264–1270. [\[CrossRef\]](https://doi.org/10.1021/acssensors.8b00269) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/29781606)
- <span id="page-17-21"></span><span id="page-17-8"></span>86. Ngo, H.T.; Gandra, N.; Fales, A.M.; Taylor, S.M.; Vo-Dinh, T. Sensitive DNA detection and SNP discrimination using ultrabright SERS nanorattles and magnetic beads for malaria diagnostics. *Biosens. Bioelectron.* **2016**, *81*, 8–14. [\[CrossRef\]](https://doi.org/10.1016/j.bios.2016.01.073) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/26913502)
- <span id="page-17-9"></span>87. Yoon, J.; Jang, W.S.; Nam, J.; Mihn, D.C.; Lim, C.S. An automated microscopic Malaria parasite detection system using digital image analysis. *Diagnostics* **2021**, *11*, 527. [\[CrossRef\]](https://doi.org/10.3390/diagnostics11030527) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/33809642)
- 88. Kim, J.D.; Nam, K.M.; Park, C.Y.; Kim, Y.S.; Song, H.J. Automatic detection of malaria parasite in blood images using two parameters. *Technol. Health Care* **2015**, *24*, S33–S39. [\[CrossRef\]](https://doi.org/10.3233/THC-151049)
- 89. Linder, N.; Turkki, R.; Walliander, M.; Mårtensson, A.; Diwan, V.; Rahtu, E.; Pietikäinen, M.; Lundin, M.; Lundin, J. A malaria diagnostic tool based on computer vision screening and visualization of *Plasmodium falciparum* candidate areas in digitized blood smears. *PLoS ONE* **2014**, *9*, e104855. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0104855) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/25144549)
- <span id="page-17-18"></span><span id="page-17-11"></span><span id="page-17-10"></span>90. Post, A.; Kaboré, B.; Reuling, I.J.; Bognini, J.; Van Der Heijden, W.; Diallo, S.; Lompo, P.; Kam, B.; Herssens, N.; Lanke, K.; et al. The XN-30 hematology analyzer for rapid sensitive detection of malaria: A diagnostic accuracy study. *BMC Med.* **2019**, *17*, 103. [\[CrossRef\]](https://doi.org/10.1186/s12916-019-1334-5)
- 91. Dumas, C.; Bienvenu, A.L.; Girard, S.; Picot, S.; Debize, G.; Durand, B. Automated *Plasmodium* detection by the Sysmex XN hematology analyzer. *J. Clin. Pathol.* **2018**, *71*, 594–599. [\[CrossRef\]](https://doi.org/10.1136/jclinpath-2017-204878)
- <span id="page-17-13"></span><span id="page-17-12"></span>92. Racsa, L.D.; Gander, R.M.; Southern, P.M.; McElvania TeKippe, E.; Doern, C.; Luu, H.S. Detection of intracellular parasites by use of the CellaVision DM96 analyzer during routine screening of peripheral blood smears. *J. Clin. Microbiol.* **2015**, *53*, 167–171. [\[CrossRef\]](https://doi.org/10.1128/JCM.01783-14) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/25378575)
- <span id="page-17-19"></span>93. Pillay, E.; Khodaiji, S.; Bezuidenhout, B.C.; Litshie, M.; Coetzer, T.L. Evaluation of automated malaria diagnosis using the Sysmex XN-30 analyser in a clinical setting. *Malar. J.* **2019**, *18*, 15. [\[CrossRef\]](https://doi.org/10.1186/s12936-019-2655-8) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/30670023)
- <span id="page-17-15"></span><span id="page-17-14"></span>94. Hashimoto, M.; Yokota, K.; Kajimoto, K.; Matsumoto, M.; Tatsumi, A.; Yamamoto, K.; Hyodo, T.; Matsushita, K.; Minakawa, N.; Mita, T.; et al. Quantitative detection of *Plasmodium falciparum* using, luna-fl, a fluorescent cell counter. *Microorganisms* **2020**, *8*, 1356. [\[CrossRef\]](https://doi.org/10.3390/microorganisms8091356) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32899795)
- <span id="page-17-23"></span><span id="page-17-16"></span>95. Costa, M.S.; Baptista, V.; Ferreira, G.M.; Lima, D.; Minas, G.; Veiga, M.I.; Catarino, S.O. Multilayer thin-film optical filters for reflectance-based malaria diagnostics. *Micromachines* **2021**, *12*, 890. [\[CrossRef\]](https://doi.org/10.3390/mi12080890)
- <span id="page-17-17"></span>96. Orbán, Á.; Longley, R.J.; Sripoorote, P.; Maneechai, N.; Nguitragool, W.; Butykai, Á.; Mueller, I.; Sattabongkot, J.; Karl, S.; Kézsmárki, I. Sensitive detection of *Plasmodium vivax* malaria by the rotating-crystal magneto-optical method in Thailand. *Sci. Rep.* **2021**, *11*, 18547. [\[CrossRef\]](https://doi.org/10.1038/s41598-021-97532-9) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/34535695)
- 97. Pukáncsik, M.; Molnár, P.; Orbán, Á.; Butykai, Á.; Marton, L.; Kézsmárki, I.; Vértessy, B.G.; Kamil, M.; Abraham, A.; Aly, A.S.I. Highly sensitive and rapid characterization of the development of synchronized blood stage malaria parasites via magneto-optical hemozoin quantification. *Biomolecules* **2019**, *9*, 579. [\[CrossRef\]](https://doi.org/10.3390/biom9100579) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31591333)
- 98. Orbán, Á.; Butykai, Á.; Molnár, A.; Pröhle, Z.; Fülöp, G.; Zelles, T.; Forsyth, W.; Hill, D.; Müller, I.; Schofield, L.; et al. Evaluation of a novel magneto-optical method for the detection of malaria parasites. *PLoS ONE* **2014**, *9*, e96981. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0096981)
- <span id="page-17-24"></span>99. Lukianova-Hleb, E.Y.; Campbell, K.M.; Constantinou, P.E.; Braam, J.; Olson, J.S.; Ware, R.E.; Sullivan, D.J.; Lapotko, D.O. Hemozoin-generated vapor nanobubbles for transdermal reagent- and needle-free detection of malaria. *Proc. Natl. Acad. Sci. USA* **2014**, *111*, 900–905. [\[CrossRef\]](https://doi.org/10.1073/pnas.1316253111)
- 100. Gandarilla, A.M.D.; Glória, J.C.; Barcelay, Y.R.; Mariuba, L.A.M.; Brito, W.R. Electrochemical immunosensor for detection of *Plasmodium vivax* lactate dehydrogenase. *Mem. Inst. Oswaldo Cruz* **2022**, *117*, e220085. [\[CrossRef\]](https://doi.org/10.1590/0074-02760220085) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/36043597)
- 101. de la Serna, E.; Arias-Alpízar, K.; Borgheti-Cardoso, L.N.; Sanchez-Cano, A.; Sulleiro, E.; Zarzuela, F.; Bosch-Nicolau, P.; Salvador, F.; Molina, I.; Ramírez, M.; et al. Detection of *Plasmodium falciparum* malaria in 1 h using a simplified enzyme-linked immunosorbent assay. *Anal. Chim. Acta* **2021**, *1152*, 338254. [\[CrossRef\]](https://doi.org/10.1016/j.aca.2021.338254) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/33648654)
- <span id="page-17-25"></span>102. Hemben, A.; Ashley, J.; Tothill, I.E. Development of an Immunosensor for Pf HRP 2 as a biomarker for malaria detection. *Biosensors* **2017**, *7*, 28. [\[CrossRef\]](https://doi.org/10.3390/bios7030028)
- 103. Jang, I.K.; Jiménez, A.; Rashid, A.; Barney, R.; Golden, A.; Ding, X.C.; Domingo, G.J.; Mayor, A. Comparison of two malaria multiplex immunoassays that enable quantification of malaria antigens. *Malar. J.* **2022**, *21*, 176. [\[CrossRef\]](https://doi.org/10.1186/s12936-022-04203-9)
- <span id="page-17-20"></span>104. Singh, N.K.; Jain, P.; Das, S.; Goswami, P. Dye coupled aptamer-captured enzyme catalyzed reaction for detection of pan malaria and p. Falciparum species in laboratory settings and instrument-free paper-based platform. *Anal. Chem.* **2019**, *91*, 4213–4221. [\[CrossRef\]](https://doi.org/10.1021/acs.analchem.9b00670) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/30793883)
- <span id="page-18-12"></span><span id="page-18-7"></span><span id="page-18-6"></span><span id="page-18-5"></span><span id="page-18-4"></span><span id="page-18-3"></span><span id="page-18-2"></span><span id="page-18-1"></span><span id="page-18-0"></span>105. Lin, H.; Zhao, S.; Liu, Y.; Shao, L.; Ye, Y.; Jiang, N.; Yang, K. Rapid Visual Detection of *Plasmodium* Using Recombinase-Aided Amplification With Lateral Flow Dipstick Assay. *Front. Cell. Infect. Microbiol.* **2022**, *12*, 922146. [\[CrossRef\]](https://doi.org/10.3389/fcimb.2022.922146) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/35811679)
- <span id="page-18-8"></span>106. Yang, D.; Subramanian, G.; Duan, J.; Gao, S.; Bai, L.; Chandramohanadas, R.; Ai, Y. A portable image-based cytometer for rapid malaria detection and quantification. *PLoS ONE* **2017**, *12*, e0179161. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0179161)
- 107. Liu, Q.; Nam, J.; Kim, S.; Lim, C.T.; Park, M.K.; Shin, Y. Two-stage sample-to-answer system based on nucleic acid amplification approach for detection of malaria parasites. *Biosens. Bioelectron.* **2016**, *82*, 1–8. [\[CrossRef\]](https://doi.org/10.1016/j.bios.2016.03.050)
- 108. Shah, J.; Mark, O.; Weltman, H.; Barcelo, N.; Lo, W.; Wronska, D.; Kakkilaya, S.; Rao, A.; Bhat, S.T.; Sinha, R.; et al. Fluorescence In Situ hybridization (FISH) assays for diagnosing malaria in endemic areas. *PLoS ONE* **2015**, *10*, e0136726. [\[CrossRef\]](https://doi.org/10.1371/journal.pone.0136726) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/26333092)
- 109. Shah, J.; Poruri, A.; Mark, O.; Khadilka, U.; Mohring, F.; Moon, R.W.; Ramasamy, R. A dual colour fluorescence in situ hybridization (FISH) assay for identifying the zoonotic malaria parasite *Plasmodium knowlesi* with a potential application for the specific diagnosis of knowlesi malaria in peripheral-level laboratories of Southeast Asia. *Parasites Vectors* **2017**, *10*, 342. [\[CrossRef\]](https://doi.org/10.1186/s13071-017-2273-7)
- <span id="page-18-13"></span>110. Peng, W.K.; Kong, T.F.; Ng, C.S.; Chen, L.; Huang, Y.; Bhagat, A.A.S.; Nguyen, N.-T.; Preiser, P.R.; Han, J. Micromagnetic resonance relaxometry for rapid label-free malaria diagnosis. *Nat. Med.* **2014**, *20*, 1069–1073. [\[CrossRef\]](https://doi.org/10.1038/nm.3622)
- 111. Thamarath, S.S.; Xiong, A.; Lin, P.-H.; Preiser, P.R.; Han, J. Enhancing the sensitivity of micro magnetic resonance relaxometry detection of low parasitemia *Plasmodium falciparum* in human blood. *Sci. Rep.* **2019**, *9*, 2555. [\[CrossRef\]](https://doi.org/10.1038/s41598-019-38805-2)
- <span id="page-18-14"></span>112. Fook Kong, T.; Ye, W.; Peng, W.K.; Wei Hou, H.; Marcos; Preiser, P.R.; Nguyen, N.-T.; Han, J. Enhancing malaria diagnosis through microfluidic cell enrichment and magnetic resonance relaxometry detection. *Sci. Rep.* **2015**, *5*, 11425. [\[CrossRef\]](https://doi.org/10.1038/srep11425)
- 113. Tomescu, O.A.; Mattanovich, D.; Thallinger, G.G. Integrative omics analysis. A study based on *Plasmodium falciparum* mRNA and protein data. *BMC Syst. Biol.* **2014**, *8*, S4. [\[CrossRef\]](https://doi.org/10.1186/1752-0509-8-S2-S4)
- <span id="page-18-9"></span>114. Awasthi, G.; Tyagi, S.; Kumar, V.; Patel, S.K.; Rojh, D.; Sakrappanavar, V.; Kochar, S.K.; Talukdar, A.; Samanta, B.; Das, A. A proteogenomic analysis of haptoglobin in malaria. *PROTEOMICS–Clin. Appl.* **2018**, *12*, 1700077. [\[CrossRef\]](https://doi.org/10.1002/prca.201700077)
- 115. Lindner, S.E.; Swearingen, K.E.; Shears, M.J.; Walker, M.P.; Vrana, E.N.; Hart, K.J.; Minns, A.M.; Sinnis, P.; Moritz, R.L.; Kappe, S.H. Transcriptomics and proteomics reveal two waves of translational repression during the maturation of malaria parasite sporozoites. *Nat. Commun.* **2019**, *10*, 4964. [\[CrossRef\]](https://doi.org/10.1038/s41467-019-12936-6) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/31673027)
- 116. Gardinassi, L.G.; Arévalo-Herrera, M.; Herrera, S.; Cordy, R.J.; Tran, V.; Smith, M.R.; Johnson, M.S.; Chacko, B.; Liu, K.H.; Darley-Usmar, V.M. Integrative metabolomics and transcriptomics signatures of clinical tolerance to *Plasmodium vivax* reveal activation of innate cell immunity and T cell signaling. *Redox Biol.* **2018**, *17*, 158–170. [\[CrossRef\]](https://doi.org/10.1016/j.redox.2018.04.011)
- <span id="page-18-10"></span>117. Commonwealth. The Commonwealth Malaria Report. 2022. Available online: [https://reliefweb.int/report/world/](https://reliefweb.int/report/world/commonwealth-malaria-report-2022) [commonwealth-malaria-report-2022](https://reliefweb.int/report/world/commonwealth-malaria-report-2022) (accessed on 15 June 2024).
- <span id="page-18-11"></span>118. Yan, S.L.R.; Wakasuqui, F.; Wrenger, C. Point-of-care tests for malaria: Speeding up the diagnostics at the bedside and challenges in malaria cases detection. *Diagn. Microbiol. Infect. Dis.* **2020**, *98*, 115122.
- <span id="page-18-15"></span>119. Veiga, M.I.; Peng, W.K. Rapid phenotyping towards personalized malaria medicine. *Malar. J.* **2020**, *19*, 68. [\[CrossRef\]](https://doi.org/10.1186/s12936-020-3149-4) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/32046739)
- <span id="page-18-16"></span>120. Su, X.-Z.; Zhang, C.; Joy, D.A. Host-malaria parasite interactions and impacts on mutual evolution. *Front. Cell. Infect. Microbiol.* **2020**, *10*, 587933. [\[CrossRef\]](https://doi.org/10.3389/fcimb.2020.587933)
- <span id="page-18-17"></span>121. Laishram, D.D.; Sutton, P.L.; Nanda, N.; Sharma, V.L.; Sobti, R.C.; Carlton, J.M.; Joshi, H. The complexities of malaria disease manifestations with a focus on asymptomatic malaria. *Malar. J.* **2012**, *11*, 29. [\[CrossRef\]](https://doi.org/10.1186/1475-2875-11-29) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/22289302)
- 122. Leski, T.A.; Taitt, C.R.; Colston, S.M.; Bangura, U.; Holtz, A.; Yasuda, C.Y.; Reynolds, N.D.; Lahai, J.; Lamin, J.M.; Baio, V. Prevalence of malaria resistance-associated mutations in *Plasmodium falciparum* circulating in 2017–2018, Bo, Sierra Leone. *Front. Microbiol.* **2022**, *13*, 1059695. [\[CrossRef\]](https://doi.org/10.3389/fmicb.2022.1059695) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/36532481)
- <span id="page-18-18"></span>123. Bull, P.C.; Berriman, M.; Kyes, S.; Quail, M.A.; Hall, N.; Kortok, M.M.; Marsh, K.; Newbold, C.I. *Plasmodium falciparum* variant surface antigen expression patterns during malaria. *PLoS Pathog.* **2005**, *1*, e26. [\[CrossRef\]](https://doi.org/10.1371/journal.ppat.0010026) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/16304608)
- <span id="page-18-19"></span>124. Aggarwal, S.; Peng, W.K.; Srivastava, S. Multi-omics advancements towards *Plasmodium vivax* malaria diagnosis. *Diagnostics* **2021**, *11*, 2222. [\[CrossRef\]](https://doi.org/10.3390/diagnostics11122222) [\[PubMed\]](https://www.ncbi.nlm.nih.gov/pubmed/34943459)

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