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Natural products as transmission-blocking agents against malaria: a comprehensive review of bioactive compounds and their therapeutic potential

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Abstract

Malaria eradication is hindered by the persistence of transmission stages of *Plasmodium falciparum* that enable parasite transfer from humans to mosquitoes. Current therapeutic strategies, such as artemisinin-based combination therapy (ACT) combined with primaguine, are insufficient due to limited efficacy on mature gametocytes and safety concerns in populations with glucose-6-phosphate dehydrogenase deficiency. This highlights the critical need for innovative, safe, and effective transmission-blocking interventions. This review explores the potential of natural sources, including medicinal plants, marine organisms, and microorganisms—as reservoirs of novel bioactive compounds with anti-malarial properties. A comprehensive literature search identified promising natural products with gametocytocidal and sporontocidal activity, validated through advanced bioassays. The review also evaluates various methodologies, such as colorimetric, microscopy, and flow cytometry assays, for assessing transmission-blocking efficacy. The findings emphasize the potent gametocytocidal effects of certain plant extracts, such as Azadirachta indica and Vernonia amygdalina, and microbial products, including ionophores and proteasome inhibitors. Despite promising in vitro and in vivo data, the transition of these compounds to clinical applications remains limited. Challenges include standardizing assays, addressing resistance to current therapies, and ensuring drug safety for endemic populations. The current review underscores the untapped potential of natural products as transmission-blocking agents and proposes a systematic, stage-specific screening cascade to identify and optimize these compounds. Addressing these gaps could significantly advance global malaria eradication efforts.

Keywords Antimalarial agents, Bioactive compounds, Gametocytocidal activity, Malaria transmission-blocking, *Plasmodium falciparum*

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Background

Malaria is a parasitic disease caused by apicomplexan parasites of the genus *Plasmodium*, which can infect a diverse range of vertebrate hosts and be transmitted by the female anopheles' mosquito. A number species are involved in human malaria, of which Plasmodium falciparum is the most prevalent in sub-Saharan Africa, accounting for 93% of estimated malaria cases in 2018 [1]. In the same year, there were an estimated 228 million cases of malaria occurred worldwide and 405000 deaths from malaria globally compared with 416,000 estimated deaths by the World Health Organization (WHO) in 2017, and 585,000 in 2010 [2]. Children aged under 5 years and pregnant women remains the most vulnerable groups. The life cycle of *P. falciparum* is highly complex, involving several developmental stages in both humans and mosquito. Mature asexual blood stages inside the human host are responsible for all the clinical symptoms of malaria, but only the non-replicating sexual blood stages of the parasite (male and female gametocytes), are able of developing in the mosquito vector and causing onward infection [3]. Though all P. falciparum infections are not severe and some are asymptomatic in highly endemic regions, they can contribute to malaria transmission [4].

The major chemotherapeutic agents to fight against remains artemisinin-based combination therapy (ACT) [5]. However, ACT has no sterilizing effect on gametocytes stages V responsible for malaria transmission [6, 7]. To overcome this gap, ACT is currently combined with primaquine in order to block human to mosquito transmission [5]. Unfortunately, primaquine safety remains a major public health concern due to its haemolytic side effect in patients with glucose-6-phosphate dehydrogenase (G6PD) deficiency [8]. Therefore, the lack of an effective vaccine and the emergence of drug-resistant parasites suggests that interventions targeting the symptom-causing (asexual) stages of the parasite are insufficient to curb malaria transmission and completely eradicate malaria. Also, research on malaria transmission blocking drug discovery targeting gametocytes stages is still lacking. Hence, efforts should be made to encourage the search for new transmission-blocking drugs, by exploring diverse sources of drugs molecules. To achieve this, one promising way to support this effort toward malaria eradication remains herbal-based medicine. Nature-derived inhibitors are the richest source of novel pharmacophores with diverse biological activity as they are known to occupy biologically important chemical space [9]. Indeed, natural products such as medicinal plants, which have been a source of some anti-plasmodial "hits" such as artemisinin extracted from the Chinese herb Artemisia annua and quinine got from the Cinchona trees [10], could be an efficient tools for malaria transmission blocking drug discovery. Indeed, most people in endemic regions heavily rely on medicinal plants for the prevention and treatment of malaria and they have served as a source of bioactive drugs against *P. falciparum* asexual-blood stages. Meanwhile, natural products are still poorly investigated for their efficacy against transmissible gametocyte stages of malaria parasites. Studies report the gametocidal activity of some medicinal plants such as Azadirachta indica, Artemisia afra, and Vernonia amygdalina [11–17] suggesting the need for detailed and depth investigation.

This comprehensive review provides insights into the discovery of malaria transmission-blocking drugs, with a significant emphasis on natural products as a promising and underexplored alternative for future therapies.

Methodology for literature search and selection

The aim of the literature search was to identify studies focused on the discovery of malaria transmission-blocking drugs derived from natural sources, specifically targeting Plasmodium falciparum gametocytes and related transmissible stages. A comprehensive search was conducted across six databases: PubMed, Web of Science, Scopus, Embase, Google Scholar, and Cochrane Library. The search utilized both Medical Subject Headings (MeSH) terms and free-text keywords, including "malaria transmission-blocking," "gametocytocidal agents," "natural products," "medicinal plants," "marinederived compounds," and "microbial bioactive agents." Boolean operators (AND, OR) were applied to combine these terms and optimize the retrieval of relevant studies. Studies were included if they were peer-reviewed articles published in English, investigated natural compounds or extracts for their transmission-blocking potential, and specifically addressed activity against gametocytes or mosquito stages of the Plasmodium life cycle. Articles were also included if they described the development or validation of assays used to evaluate the transmissionblocking efficacy of natural products. Systematic reviews and meta-analyses were considered if they provided substantial information on natural product-based drug discovery. Exclusion criteria were studies focusing solely on asexual blood-stage anti-malarials, those with incomplete data or methodological descriptions, and nonpeer-reviewed materials such as conference abstracts or opinion pieces. Studies unrelated to natural product drug discovery or without clear relevance to malaria transmission-blocking strategies were also excluded. The most representative data were summarized in Tables and Figures.

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Targeting *Plasmodium* gametocytes: biology and transmission-blocking strategies *Plasmodium transmissible stages: gametocytes*

The complete *Plasmodium* life cycle includes the sexual developmental stages in the mosquito as well as the hepatic sporozoite and intra-erythrocytic sexual gametocyte stages in humans [7, 18] (Fig. 1). *Plasmodium* gametocytes are the only parasite stage that can be transmitted to the mosquito and derived from the sexual development in a small per cent of parasites and are void of clinical manifestations [3]. Despite their critical role in malaria transmission, understanding gametocyte biology remains a key focus for developing effective transmission-blocking agents.

Once gametocytes are ingested during a blood meal in the mosquito midgut, they become activated and differentiate into male and female gametes. The male gamete then fertilizes the female gamete, resulting in the formation of a zygote, which further develops into a motile ookinete, penetrates the gut epithelium, and subsequently develops into an oocyst. Maturation of the oocyst leads to the production of sporozoites, which migrate to the salivary gland of the mosquito and are ready for the next round of blood meal. Gametocytogenesis progresses through five morphological stages I-V over the course of 8-10 days [19]. Early gametocytes in stages I to III are susceptible to most classical schizontocidal anti-malarial drugs, whereas mature gametocytes in stages IV and V are insensitive to most available anti-malarial drugs. Only mature stage V gametocytes can develop in the Anopheles mosquito and remain the main stage to be interrupted in the hope of blocking human to mosquito malaria transmission and vice versa. In addition, gametocyte stages I to IV were reported to sequester in deep organs such as bone marrow and spleen and, therefore, difficult to be targeted by anti-malarial drugs whereas gametocyte stage V is the only circulating stage found in the peripheral blood where they take an additional 3 days to become infectious to mosquito [20, 21]. Gametocytes, specifically stage V provide a plausible link in malaria transmission from the human to mosquito, thereby making them a prime target for transmission-blocking strategies.

Transmission-blocking strategies

For making malaria elimination and eradication a true reality, interruption of parasite transmission is essential. Therefore, the role of transmission-blocking drugs becomes very peremptory [22]. The ability to block transmission relies on identifying (and curing) asymptomatic or semi-immune human hosts carrying transmissible forms of the parasite that represent the major reservoirs of continued infection; eliminating mosquito vectors through multiple and integrated strategies; and eliminating the parasite pool in patients with malaria, which, in the absence of a vaccine, still relies solely on chemotherapy and prophylaxis to prevent new or re-infection [23, 24]. Transmission blocking agent should ideally target and kill gametocyte stages, especially the transmissible mature gametocyte stages V [25], and thereby prevent malaria transmission from human to mosquito by clearing most gametocytes in the human host to render those patients non-infectious to mosquito [26]. In line with

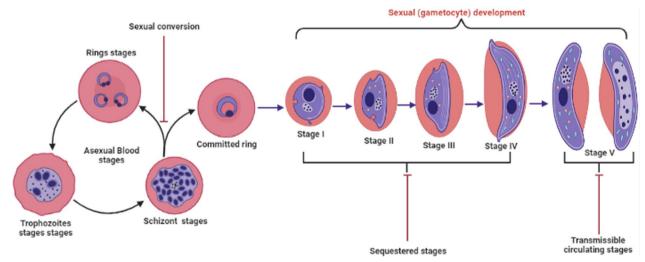


Fig. 1 Developmental stages and transmission potential of *Plasmodium* gametocytes. The diagram depicts the *Plasmodium* gametocyte life cycle, starting with sexual conversion from asexual blood stages to committed rings, which then progress through five gametocyte stages (I–V). Stages I–IV are sequestered in tissues, making them less accessible to drugs. Only mature stage V gametocytes circulate in the blood, becoming transmissible to mosquitoes and critical for malaria spread. This figure was created using BioRender.com elements

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this priority, strategies targeting the vector stages would require a drug to be at pharmacologically relevant concentrations for as long as mature gametocytes circulate (up to 30 days). So, the most appropriate point of intervention remains to target the host gametocytes especially stage V.

Current preclinical assays for transmission-blocking drugs discovery: challenges and prospects Gametocyte production

Gametocytes are currently induced in culture by non-specific stressors, such as changes in hematocrit, addition of mammalian hormones, reticulocytes, nucleic acid synthesis inhibitors (e.g., antifolates), and various drugs like Berenil, chloroquine, and amodiaquine [20, 27–31]. However, standardized, reproducible protocols for gametocyte production are lacking, making it difficult to compare findings across laboratories [32]. Furthermore, newer methods using genetic modification and controlled microenvironments aim to improve consistency in producing high-yield, stage-specific gametocytes for drug screening [33].

Current anti-gametocyte assays

There have been significant investments over the past decade and technological advances in the validation of some standard "methods" for transmission blocking-drug discovery. Several methods for assessing the transmission-blocking potential have been published using a variety of detection methods. These include measurement of metabolic activity such as parasite adenosine triphosphate (ATP) and guanosine triphosphate (GTP) levels [34], expression of a gene reporter [35]; colorimetric assays [36, 37], microscopy and flow cytometry methods.

Colorimetric methods

Colorimetric methods to assess transmission-blocking efficacy of a drug are currently based on metabolic activity such as enzymatic reaction-based spectrophotometry or fluorescence among all *Plasmodium* lactate dehydrogenase (pLDH) based assay [36] relies on the rapid utilization of 3-acetylpyridine adenine dinucleotide (APAD) as a coenzyme by pLDH in the reaction leading to the conversion of lactate to pyruvate [32] and Resazurin/ Alamar blue-based assay using resazurin, a cell-permeant blue dye that functions as a cell viability indicator upon reduced to resorufin (red) by the metabolic activity of living cells [37].

These methods are used for both the early and late stages of transmissible gametocytes. pLDH is a reliable marker for gametocyte viability as the enzyme is present at high levels throughout gametocytogenesis [38, 39]. One major advantage of the pLDH assay is that it is

performed directly on parasite cultures, therefore minimizing the manipulation of gametocytes. However, late-stage gametocytes exhibit decreased expression of genes responsible for glycolysis, protein biosynthesis, and hae-moglobin catabolism [40]. While, redox reactive (oxidoreductive indicators) cell-permeable dyes like Alamar blue have previously been reported as robust assays supporting high-throughput screening, more sensitive than traditionally used tetrazolium dyes [37, 41]. Though these assays enable medium to high-throughput screening against gametocytes, care must be taken especially where compounds target the redox state of the parasite, as it may interfere with the assay readout. Moreover, standard drugs routinely used like methylene blue cannot be used in this assay platform due to colorimetric interference.

Reporter gene methods

This method consists of ATP bioluminescence assay, a luciferase reporter assay, a green fluorescent protein (GFP)-luciferase bioluminescence assay, a luminescenceindirect standard membrane feeding assay (SMFA) and a luminescence-direct SMFA. Luciferase reporter assay consists of both a luciferase reporter enzyme and a detection reagent that provides the enzyme substrate. In this assay, the light emitted is proportional to the reporter gene expression level when the reporter enzyme and detection reagent are combined. Luciferase-expressing *Plasmodium berghei* parasites are currently employed to monitor the parasite's sporogony development in vitro, as well as a tool for the identification of compounds with transmission-blocking activity. This assay has been established to enable accurate, reliable and quantifiable investigations of the stage-specific action of gametocytocidal drugs for each of the early and late gametocyte marker cell lines [42]. However, the major concern with this technique is a higher inter-assay variability between different assays and the problem of ATP interference. However, this method has to be improved in the future by using the recently reported *P. berghei* line expressing the novel luciferase enzyme NanoLuc (PbNLuc) that resulted in delivering a significantly enhanced luminescence signal, enabling single parasite detection in various stages of its life cycle, including in the mosquito vector [43]. Another promising way to improve the assay will be the use of a second reporter gene, placed under the control of a sporozoite-specific promoter, which would enable the specific monitoring of sporozoite formation and maturation [44]. ATP plays a central role in energy exchanges in biological systems (both in eukaryotic and prokaryotic cells), serves as the main donor of free energy, and is produced in all metabolically active cells. The ATP bioluminescent assay is classified as a gold standard for exploring parasite viability. Therefore, this parameter is

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currently used as a suitable tool to assess the functional integrity of P. falciparum gametocytes. Injury and death lead to a rapid decrease in cytoplasmic ATP, [45] providing a reliable platform to test the effect of small molecules on the gametocyte's ability. In addition, transgenic P. falciparum parasites are used to quantify the effects of inhibitors either on asexual blood-stage in vitro using standard assays [46] or on transmission-blocking activity in the mosquito in high-throughput screens using SMFA [47, 48]. For the transmission-blocking assays transgenic P. falciparum (NF54 strain) parasite lines expressing the GFP-luciferase fusion protein under the control of either the strong constitutive P. falciparum hsp70 [47] or the gametocyte-specific pfs16 promoters are currently used [48]. The use of transgenic parasite lines expressing GFPluciferase fusion protein to evaluate the transmissionblocking potential of a given molecule is not without its limitations. First, the promoter is less suitable for high reporter expression in liver stages. Second, when using 'additional copy' chimeric parasites, the gene expression is dependent on the promoter used, which is unlikely to exactly mimic the timing and magnitude of the gene expression [1]. Gametocytes, ookinete, sporozoites, oocytes and liver stages are the main targeted stages in this method. These methods are easy, accurate, reliable, reproducible and quantifiable. However, a higher interassay variability has been noticed.

Microscopy methods

Microscopic examination of Giemsa's stained slides remains the 'gold standard' method for several decades for the detection, quantification, speciation and staging of blood-stage malaria parasites [49]. Today, this technique is currently used in the search for malaria transmission-blocking agents. Microscopy is currently used in association with other techniques in the malaria transmission-blocking drug discovery pipeline. Among these, we have, light microscopy-indirect SMFA- P. falciparum; light microscopy-direct SMFA-P. berghei; automated microscopic analysis and high-content imaging. Light microscopy associated with SMFA is currently used as the confirmatory tool to confirm the transmission-blocking potential of a given molecule. Therefore, P. falciparum SMFA remains the current "gold standard" mosquito-based confirmatory transmission-blocking assay for human malaria. However, owing to its complexity, only selected gametocidal molecules are progressed into SMFA. Predictive tools for the evaluation of transmission-blocking behaviour of compounds in SMFA would be extremely beneficial, but the lack of substantially large data sets from many mosquito feeds preempts the ability to perform correlations between outcomes from in vitro assays and in vivo assay through SMFA [47, 50], and have more overlap with the parasite's life cycle than any of the other gametocytocidal or gametogenesis inhibition assays [51]. The major problem encountered with light microscopy is that this technique is tedious, time-consuming and relies on the skills of technicians of microscopy techniques. The technician must be trained to correctly identify the different parasite species and stages [52, 53]. This final point is of greatest concern, as this inter-reader viability, gives rise to the common criticism that microscopy counts are relatively subjective.

Flow cytometry methods

As a laser-based technology, flow cytometry is largely dependent on the light-scattering properties of cells and particles that make it possible to analyse such characteristics as the size of cells, DNA content within a cell, as well as cell granularity among others. Plasmodium gametocyte quantification based on flow cytometry has been recently proposed with the goal of increasing precision and objectivity in malaria transmission-blocking drug discovery. Flow cytometry is currently used in association with other methods such as microscopy to assess the effect of promising asexual-blood stages inhibitors to distinguish pan-reactive drugs with multistage activity against the malaria parasite. Previous studies on the flow cytometry method allow highly reproducible quantification of specific drug effects on sexual conversion and early sexual development [54]. Flow cytometry in combination with magnetic enrichment has been a useful tool to estimate the inhibitory concentration of known drugs against P. falciparum gametocytes and remains useful to evaluate promising anti-gametocyte drugs. This method is generally based on the detection of Plasmodium double-stranded DNA in the infected erythrocytes [55]. However, this technique is subject to many challenges as far as P. falciparum gametocytes do not go through DNA replication. In addition, flow cytometry cannot distinguish between live and dead parasites [56].

Overall, these methods are used on the later stage and/ or on mature gametocytes, as the immature stages have often been hypothesized to be sensitive to most antimalarial drugs and, hence, of limited concern. To really confirm the transmission-blocking potential of such inhibitors, some of those methods are currently used in association with SMFA known today as a "gold standard" in the field of malaria transmission-blocking drugs discovery. Those available methods already stabilized represent an important starting point for transmission blocking-drug screening. However, each method has its own advantages and limits and thereby required further improvement. Some molecular players such as PfAP2-G could be very useful for inhibition or prevention of sexual

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commitment and therefore limiting human to mosquito malaria transmission and vice versa.

Existing therapeutic approaches

Chemical-based transmission-blocking drugs

Currently, the available drugs with therapeutic activity against late-stage gametocytes is artesunate, artemether, methylene blue, tafenoquine, and primaguine (Table 1) [33, 57]. For its activity on gametocytes, primaquine has been recently recommended at a single low dose (0.25 mg/kg) alongside anti-malarial treatment in disease-endemic countries. Again, ACT is currently used in combination with primaquine in order to treat and block transmission from humans to mosquitoes [5]. To protect the patient from the emergence of artemisinin resistance and reduce malaria transmission, researchers have developed a combination of ACT with methylene blue for malaria eradication [58] relying on the effect of methylene blue on parasite asexual-blood stages and, mature male and female gametocytes. However, ACT recommended for malaria transmission-blocking are not active on mature gametocyte stages V involved in transmission and are all threatened by emerging resistance [6, 11]. Besides, the use of primaquine is severely restricted due to its haemolytic issues in individuals with glucose-6-phosphate-dehydrogenase (G6PD) deficiency [33, 59]. Moreover, very few new agents are in development specifically for malaria transmission-blocking, although some have multistage activity [1]. Overall, new tools to specifically target sexual commitment are urgently needed for malaria control and prevention [3]. Indeed, the transmission of malaria can also be interrupted by targeting other parasitic stages, known as the sporogony stages (gametes, zygotes, ookinetes, oocysts and sporozoites), which occur inside the mosquito. These mosquito stages are also attractive drug targets because parasites in the majority of these stages exception of sporozoites, constitute the bottleneck population [22]. Targeting the parasite's sporogony stages remains an indirect transmission-blocking method of drug delivery due to the presence of these stages, not in the bloodstream of the host but inside the body of the mosquito. Drugs targeting the sporogony stages inside the mosquito vector has to be present in the host's blood circulation over and above the concentration required to kill these mosquito-stage parasites, for a sufficient period [51] making gametocytes a more suitable target for a transmissionblocking drug search. In recent years, many potential new antimalarial transmission-blocking drugs have been discovered. Advanced transmission-blocking drugs in

 Table 1
 Summary on current experimental assays used for transmission-blocking drug discovery

Transmission Blocking assay	Variants	Targeted stages	Turnaround time	Advantages	Limits	References
Colorimetric methods	-pLDH-based assay -Resazurin/Alamar blue based-assay	Gametocytes	48–72 h	-Robust, fast, easy, cheap and reproduc- ible	Colorimetric interference with the drug	[37, 41]
Reporter gene methods	-ATP bioluminescence assay -Luciferase reporter assay -GFP-Luciferase biolu- minescence -Luminescence- Indi- rect SMFA -Luminescence- Direct SMFA	Gametocytes Ookinete Oocyst Sporozites Liver stages	24–72 h	- Minimal interference -Easy, accurate, reproducible, reliable and quantifiable	-Higher inter-assay variability -ATP interference	[47, 48, 99]
Microscopy	-Light microscopy- Indirect SMFA- <i>P.</i> <i>falciparum</i> - Light microscopy- Direct SMFA- <i>P. berghei</i> - Automated microscopic analysis and High content imaging	Oocyst, Sporozo- ites, Male gam- ete, Ookinete	0–72 h	Easy, reliable and reproducible	-Require well-trained microscopist - Inter-reader viability	[44, 100, 101]
Flow cytometry	-Flow cytometry- gametocyte commit- ment assay	Early and late gametocyte stages	48 h	-Accurate and reproducible -Fast, robust and simple -Precise and objective	-Absence of DNA since these cells do not multiply	[54, 56, 102]

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the development pipeline include the synthetic endoperoxides OZ439 that has proven activity against the asexual stages of the parasite [60] and is currently in phase IIa clinical trials, and cipargamin, which has previously demonstrated potent dose-responsive activity against the sexual stages of P. falciparum [61]. Cipargamin, the most advanced candidate in blocking human-to-mosquito transmission and OZ439 scientifically called artefenomel (Fig. 2). Their structural design has been inspired by artemisinin [60] and they are two transmission-reducing drugs under development. The effect of cipargamin on gametocytes and the transmission-reducing potential was investigated in vitro and compared to other antimalarials by Van Pelt-Koops et al. [61]. In this investigation, the authors firstly measured the in vitro activity on asexual P. falciparum parasites from the drug-sensitive NF54 and K1 strains, which showed favourable results for cipargamin compared to lumefantrine, artemether and primaguine. Secondly, gametocytocidal activity was measured, with cipargamin being the most effective inhibitor of early and late gametocyte development. Thirdly, these four compounds were added to blood meals of blood-feeding Anopheles mosquitoes, where cipargamin and lumefantrine both independent yielded reduced oocyst counts [61]. Additionally, the transmission blocking effect of cipargamin have been reported by Upton and colleagues in 2015 [62]. Results from the studies of Charman and colleagues (2011) demonstrated that the endoperoxide, OZ439, has sterilizing effect against

CINHO

Cipargamin

Fig. 2 Chemical structures of Cipargamin and Artefenomel (OZ439)

P. falciparum transmissible gametocytes with several advantages compared to artemisinin derivatives, such as prolonged plasma exposure, higher potency, stability and its synthetic nature [60]. It has shown comparable efficacy in the treatment of *P. falciparum* and *P. vivax* patients [60] are undergoing phase IIa clinical trials and is intended to be used as a single-dose combination therapy for acute malaria. It has previously been demonstrated to completely cure P. berghei-infected mice (bloodstage infection) with a single oral dose of 20 mg/kg [60] but does not affect P. berghei ookinete development in in vitro assays at 10 µM, suggesting it functions before the ookinete stage [63]. It has additionally demonstrated a potent effect in the SMFA with P. falciparum at 10 µM [63], OZ439 significantly reduced oocyst and sporozoite development in the mosquito when administered 24 h pre-feed at the dose of 6.5 mg/kg but did not significantly reduce the number of secondary mouse infections.

In the light of the above limitations encounter by chemical molecules used as transmission-blocking drugs, natural products and their derivative appear as a promising alternative (Fig. 3).

In fact, nature remains an attractive, credible, untapped source of new therapeutic candidate compounds since a tremendous chemical diversity is found in medicinal plants, animals, marine organisms, and microorganisms [12, 64]. Despite major scientific and technological progress in combinatorial chemistry, drugs derived from natural products still make a significant contribution to drug discovery today and the role of natural products as a source for remedies has been formally recognized [64] (Fig. 4). In addition, natural products have been used as scaffolds for the development of new products with huge therapeutic and industrial potential due to their unusual chemical features [65]. Therefore, the depth investigation of natural products and their derived phytoconstituents toward transmission-blocking drugs discovery needs a well-defined strategy. Hence, The transmission blockingbased natural product agents research strategies might include (i) the exploration of various sources of natural products (such as plants, marine algae and microorganisms) for gametocidal potency, (ii) ethnopharmacologybased plants selection of plants traditionally used in disease endemic country for malaria prevention; (iii) the repurposing of natural products with activity against asexual-blood stages; (iv) develop and validate more suitable, time effective and easier way to run the assay. Overall, natural products remain underexploited and there is no transmission-blocking natural product in clinical development. This underlines the low rate of progress in identifying gametocidal compounds from a natural source and the key importance of designing and implementing a clear, rationale, and sustainable screening test

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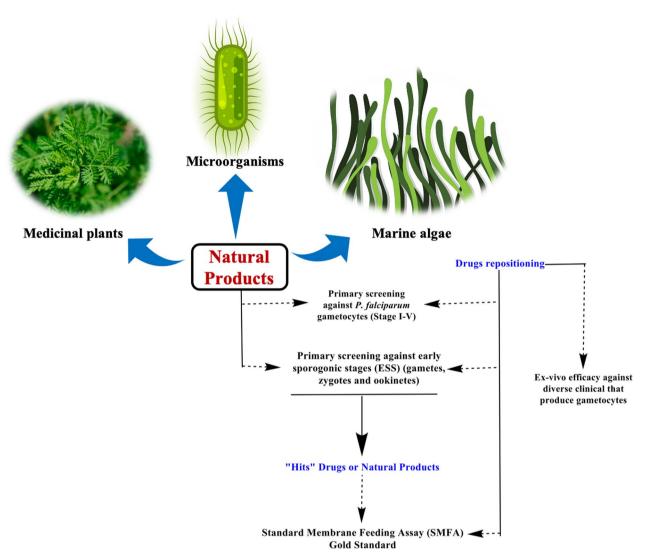


Fig. 3 Diagram with the sources and screening pipeline for identifying natural products with transmission-blocking potential against malaria. Natural products derived from medicinal plants, microorganisms, and marine algae serve as primary sources of bioactive compounds. Initial screening is conducted against Plasmodium falciparum gametocytes (Stages I-V) and early sporogonic stages (ESS), including gametes, zygotes, and ookinetes. Promising compounds, termed "hit" drugs or natural products, undergo further validation in ex vivo efficacy assays and standard membrane feeding assays (SMFA), considered the gold standard for assessing transmission-blocking potential. Additionally, drug repurposing approaches are incorporated to identify established drugs with potential activity against transmissible malaria stages

(See figure on next page.)

Fig. 4 Proposed test cascade approach for screening and isolation of transmission-blocking antimalarial compounds from natural products. Natural products sourced from medicinal plants, microorganisms, and marine algae undergo primary screening against *Plasmodium falciparum* transmissible stages. Promising extracts are identified at 100 μ g/mL and subjected to dose–response testing, with active extracts showing IC₅₀ \leq 10 μ g/mL progressing further. Bioassay-guided purification isolates compounds with IC₅₀ \leq 5 μ g/mL, which then undergo studies of mode of action, including time-kill assays and stage-specific activity (early vs. late stages). Additional testing includes target identification and structure–activity relationship (SAR) studies. Active compounds are further assessed in in vitro male/female gamete assays and finally validated in the Standard Membrane Feeding Assay (SMFA) for transmission-blocking efficacy

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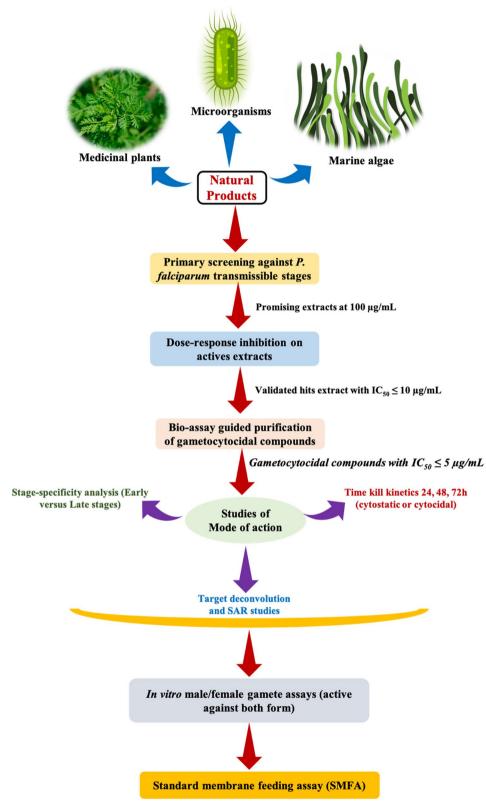


Fig. 4 (See legend on previous page.)

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cascade for the isolation of naturally derived transmission-blocking compounds. Hence, the existing approach used for transmission blocking-drug discovery did not take into consideration a comprehensive cascade strategy in order to discover a promising molecule able to inhibit human gametocytes stages and mosquitoes' stages such as ookinete, gametes and zygote. In that regard, the discovery of transmission-blocking drugs from natural resources or from an existing set of drug molecules should first start with the screening of human transmissible gametocytes in order to identify a "hit" molecule that could screen on mosquitoes stages and in vivo using SMFA.

Role of plant extracts and secondary metabolites in malaria transmission-blocking

Medicinal plants used as alternative drugs are indicative of the vital role that plants play in many developing countries, and are also sources of novel plant-derived phytoconstituents that could be lead to the treatment of malaria and other diseases [66, 67]. The WHO estimated that ~65–80% of the world population ensured their primary health care using predominately plant-derived traditional medicines, existing a lower prevalence in developed countries [68]. Drug discovery approaches using medicinal plants as a tool provide diversity and structurally different compounds targeting multiple P. falciparum life cycles including liver stages, asexual-blood stages in the individual and transmissible gametocytes stages in the human host and mosquito's vector. Moreover, medicinal plants remain a promising alternative to conventional medicine, considering the rich biodiversity of the continent. Moreover, people living in Africa's endemic countries use medicinal plants even when they are not suffering from malaria. Indeed, they use it as a preventive strategy to stay safe from malaria infection and highlight the fact that, medicinal plants used for malaria prevention could contain preventive bioactive phytoconstituents with the potential to reduce malaria transmission as it has worked in the past for symptom-causing asexual blood stages. Among the few available drugs used in the fight against malaria transmission, only artemisinin used in combination with primaquine, comes from a medicinal plant, the Chinese herb Artemisia annua highlighting the fact that medicinal plants have been neglected for transmission-blocking drug discovery. Hence, medicinal plants present in nature remain powerful tools and inexhaustible sources that could accelerate malaria eradication worldwide. However, there is a paucity of studies that demonstrate the transmission-blocking effect of traditionally used medicinal plants involving human malaria parasite P. falciparum. The data in Table 2 summarize medicinal plant species investigated for their potential to inhibit or kill transmissible gametocyte stages in vitro or in vivo. In addition, it is evident from the in vitro assays that the pLDH-based assay was the most used to assess the gametocidal effect and therefore, highlighting the fact that, this method could be the easier method for investigating transmission-blocking anti-malarial compounds.

In vitro findings on gametocytocidal activity

Scientific literature only reports very few in vitro studies (Table 2) investigating the potential of medicinal plants or another natural resource to block human-to-mosquito malaria transmission by inhibiting P. falciparum gametocytes. Moving in that way, a study conducted by [12] reported the inhibition of P. falciparum early and latestage gametocyte viability by extracts from eight traditional South African medicinal plants (Table 2). As a result, the IC₅₀ of tested plant extracts ranged from 5.70 to 35.95 μ g/mL in the early stage and from 3.20 to 35.95 µg/mL in the late gametocytes stages. Among the thirteen plant extracts investigated, three of them viz Artemisia afra, Trichilia emetica and Turraea floribunda were more potent in both stages with IC_{50} < 10 µg/mL. Besides, the three most active extracts in both early and late stages belong to Asteraceae and Meliaceae families. Extracts from Combretaceae family well-known for their pronounced activity against asexual-blood stages of P. falciparum showed moderated activity on both gametocyte stages with IC₅₀ ranging from 28.35 to 32.74 μ g/ mL. In addition, Amoah et al. [14] reported the gametocidal activity of eight Ghanaian herbal medicines on P. falciparum gametocytes. From this investigation, authors showed that two of the herbal-antimalarial plant's products CM (in which active ingredients are Cryptolepis sanguinolenta and Azadirachta indica) and RT (in which active ingredients are Aloe schweinfurthii, Khaya senegalensis, Piliostigma thonningii and Cassia siamea) inhibited the growth of late-stage gametocytes by >80% at 100 μg/mL whilst one, KG (in which active principle are Nauclea latifolia, Phyllanthus fraternus, Cryptolepis sanguinolenta) was the most active on early-stage gametocytes at that same concentration. However, at 1 µg/mL, only YF (mixture of C. sanguinolenta and A. indica) significantly inhibited the survival of late-stage gametocytes although at that same concentration YF barely inhibited the survival of early-stage gametocytes. Soré et al. [15] reported the activity of aqueous, ethanolic, and methanolic extract from leaves and stem bark of five medicinal plants widely used traditionally in western Burkina Faso on P. falciparum gametocytes stages using a luciferase assay (Table 2). In the same vein, Udeinya and colleagues showed in vitro gametocidal effects of neem (A. indica, Meliaceae) leaf and seed extracts on P. falciparum (Table 2). At a dosage of 2.5 g/mL, neem extract kills Fokou *et al. Malaria Journal* (2025) 24:164 Page 11 of 22

Table 2 Current transmission drugs, structure, chemical class, mode of action and preferential gametocytes stages of action

Name of	Chemical class	Structures	Mode of action in I	P. falciparum	Stage targeted	Refs.
compound			Molecular target	Pathway		
Primaquine	8-aminoquinoline	NH ₂	Not yet identified	Precisely unclear but generated metabolites might be interfering with the redox metabolism of the parasite	Early-stage I to III	[103]
Tafenoquine	8-aminoquinoline	$ \begin{array}{c c} & & & \\ & & \\ F & & \\ \end{array} $	Not yet identified	Not yet identified	Early-stage I to III	[104]
Artesunate	Endoperoxide	ОН	Proteasome/Existing proteins/Newly synthesized proteins		Early (I–III) and Late (V) stage	[105]
Methylene blue	Benzothiazine	N S Cr	Monoamine oxidase Glutathione reductase Hemozoin forma- tion	Glutathione reductase or inhibi- tion of hemozoin formation	Late-stage V	[106, 107]
Artemether	Endoperoxide		Proteasome/Existing proteins/Newly synthesized proteins	Inhibition of protein translation	Early (I–III) and Late (V) stage	[85, 105]

IC₅₀ Inhibitory Concentration 50, ND Not Determined, IRBCs Infected Red Blood Cells

more than 90% of *P. falciparum* immature and mature gametocytes in culture [16, 69].

In vivo findings on transmission-blocking

The results from Table 3 summarize a subset of medicinal plants investigated for their transmission-blocking properties in vitro and in vivo. This investigation shows that the studies were mostly conducted in vitro with very few in vivo. Among all investigated plant extracts, Azadirachta indica and Vernonia amygdalina [17] are the only ones investigated in vivo and showed a good reduction of oocyst density in mosquitoes, preventing, therefore, malaria transmission back from mosquitoes to humans very few or no study have been carrying out so far toward the investigation of plant-based medicine for their ability to block human to mosquitoes malaria transmission. An in-depth literature review showed that aqueous, ethanolic plant extracts and fractions from Vernonia amygdalina display in vivo transmission-blocking potential by reducing the P. berghei macrogametocyte density in mice by about 50%. Ver-EtOH reduced P. berghei oocyst prevalence and density by 27 and 90%, respectively and inhibited almost completely (> 90%) early sporogony stages (ESS) development in vitro at 50 μ g/mL. At the same dose, four fractions obtained from the ethyl acetate phase of the methanol extract displayed inhibitory activity > 90% against ESS [13].

commercially available standardized extract from neem seeds, 'NeemAzal' (rich in azadirachtin: azadirachtin A [34%], azadirachtin B to azadirachtin K [16%]) was examined for sporontocidal activity in 'miceto-mosquito and mosquito-to-mice' transmission model [17]. In this study, ookinete formation and oocyst development were completely blocked inside the Anopheles stephensi mosquitoes, when the mosquitoes were fed with blood from the gametocytaemic mice treated with 50 mg/kg "NeemAzal". This complete blockage of transmission was further confirmed when healthy uninfected mice on which these mosquitoes were fed, showed no oocysts. However, upon midgut dissection zygotes and post-zygotic forms were observed but these were less in number and of abnormal morphology. Since no mature ookinetes were observed at 50 mg/kg "NeemAzal", it can be hypothesized that the extract probably interferes

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Table 3 Medicinal plants investigated for their transmission-blocking potential

Plants	Family	Transmission	n blocking po	tential					
species		In vitro activ	ity		In vivo activity				
		IC ₅₀ on early stages (μg/ml	IC ₅₀ on late stages (μg/ ml)	Methods used	Micro gametocytesper 2,500 IRBCs (95% CI)	Micro gametocytes per 2,500 IRBCs (95% CI)	Percentage inhibition Early Sporogonic Stages (EES)	Reduction in Oocyst density	References
Argenome mexicana	Papaver- aceae	35.95 ± 11.1	35.95 ± 11.1	Luciferase assay	ND	ND	ND	ND	[15]
Artemisia afra	Asteraceae	5.7 ± 0.3	3.2 ± 0.2	pLDH-based assay	ND	ND	ND	ND	[108]
Azadirachta indica	Meliaceae	ND	ND	Microscopy	ND	ND	ND	86.0–93.5	[109]
Catha edulis	Celastraceae	> 10	> 10	pLDH-based assay	ND	ND	ND	ND	[13]
Combretum collinum	Combreta- ceae	28.35 ± 3.3	28.35 ± 3.3	Luciferase assay	ND	ND	ND	ND	[15]
Guiera sen- egalensis	Combreta- ceae	ND	ND	Microscopy	ND	ND	ND	No activity	[109]
Khaya anthotheca	Meliaceae	> 10	> 10	pLDH-based assay	ND	ND	ND	ND	[108]
Leonotis leonurus	Lamiaceae	12.8 ± 1.4	13.9 ± 2.6	pLDH-based assay	ND	ND	ND	ND	[108]
Leonotis leonurus	Lamiaceae	> 10	> 10	pLDH-based assay	ND	ND	ND	ND	[108]
Lophira Ianceolata	Ochnaceae	11.35 ± 3.2	11.35 ± 3.2	Luciferase assay	ND	ND	ND	ND	[15]
Olea euro- paea sub sp. Africana	Oleaceae	> 10	> 10	pLDH-based assay	ND	ND	ND	ND	[13]
Terminalia macroptera	Combreta- ceae	32.74 ± 11.9	32.74 ± 11.9	Luciferase assay	ND	ND	ND	ND	[15]
Trichilia emetica Vahl subsp. Emetica	Meliaceae	7.6 ± 0.9	3.8 ± 0.5	pLDH-based assay	ND	ND	ND	ND	[108]
Turraea floribunda	Meliaceae	9.2 ± 0.2	4.6 ± 0.7	pLDH-based assay	ND	ND	ND	ND	[108]
Vernonia amygdalina	Asteraceae	ND	ND	Microscopy	5.4-8.0	32–44.8	93.4–99.1	26-53	[13]
Zanthoxylum zanthoxy- loïdes	Rubiaceae	20.63 ± 2.3	20.63 ± 2.3	Luciferase assay	ND	ND	ND	ND	[15]

with gametogenesis and ookinete development. Also, "NeemAzal" at 50 mg/kg was not found to disrupt oocyst maturation during its incubation with early oocysts, suggesting a lack of any activity against the early oocysts. The observed inhibition of gametogenesis and ookinete formation was attributed to the microtubule-targeted mode of action of neem extracts containing azadirachtin [17]. Moreover, authors equally reported the transmission-blocking potential of an azadirachtin-enriched extract of neem seeds at 50 mg/kg mouse body weight in rodent

malaria in vivo model of *Plasmodium berghei/Anopheles stephensi*. *Azadirachta indica* is a popular medicinal plant used in various Asian and African countries for the treatment of various illnesses including malaria. Furthermore, two *Az.. indica* fractions codified as PNE and ENM were found to be more active than the third fraction ENS at 0.125% v/v and 0.25% v/v against early and late-stage gametocytes [70]. Two fractions (IRDN-A and IRDN-B) obtained from an extract of neem leaves led to complete inhibition of early-stage and late-stage gametocytes

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at 1 and 100 μ g/mL respectively (IC₅₀ for IRDN-A and IRDN-B against both early and late-stage gametocytes = 10^{-3} μ g/mL) [17]. *Azadirachta indica* is one of the important medicinal plants which has been investigated via in vivo experimental trial and showed promising results in malaria transmission-blocking experiments.

Plasmodium transmissible gametocytes Plant-derived compounds

Medicinal plants are investigated as one of the important raw materials which act as excellent malaria transmission-blocking agents. Two compounds 1α,4αdihydroxybishopsolicepolide and yomogiartemin, isolated from Artemisia afra showed good inhibition on P. falciparum gametocytes stages and the two others from Vernonia amygdalina viz vernolide and vernodalol have the potential to inhibit transmissible gametocytes stage V. It was also reported that azadirachtin from neem tree prevents the formation of motile male gametes in P. berghei with an IC₅₀ of 3.5 µM and completely inhibited ex-flagellation in P. falciparum at 100 µM [71]. In addition to the compounds previously described, several other natural products have shown promising transmission-blocking activity. These include Mallotojaponin C, a clerodane diterpenoid from *Mallotus japonicus*, which exhibits gametocytocidal effects [72]; Parthenin and Parthenolide, sesquiterpene lactones from Parthenium hysterophorus and Tanacetum parthenium, respectively, reported to impair gametocyte development [73]; and securinine-related alkaloids [74], which have demonstrated activity against sexual stages of Plasmodium. These compounds, structurally related to artemisinin, may offer potential scaffolds for developing new transmission-blocking drugs.

Moussavi and co-authors in 2015 identified pellitorine as the main insecticidal component in Zanthoxylum heitzii bark against Anopheles gambiae mosquitoes [75], while P. berghei ookinete conversion has been demonstrated to be inhibited by dihydronitidine (0.59 µg/ml) and heitziquinone (6.2 µg/ml) [76]. Similarly, Cox and co-authors in 2024 identified brusatol, a natural product, as a potential transmission blocker through in vitro screening and in vivo analyses [77]. Furthermore, Pulixin, a chromene compound isolated from Artemisia afra, has been identified as a potent inhibitor of gametocyte maturation and parasite transmission [78]. Sterigmatocystin, a fungal metabolite structurally related to aflatoxins, also exhibits notable inhibitory effects on Plasmodium sporogonic stages [78]. These additions highlight the chemical diversity of natural sources with potential transmission-blocking activities. The discussion on plantderived malaria transmission-blocking compounds has been limited here, as this topic has already been comprehensively reviewed by Moyo et al. [79]. Figure 5 shows

Fig. 5 The structures of medicinal plant-derived compounds with anti-gametocyte properties

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the structures of medicinal plant-derived compounds with anti-gametocyte properties.

Microorganisms

Microorganisms such as bacteria and fungi have been invaluable in discovering drugs and lead compounds. In addition, microorganisms produce a wide variety of antimicrobial agents which have evolved to give their hosts an advantage over their competitors in the microbiological world [64]. On the other hand, microorganisms living closely associated with plants present in their neighbourhood and, particularly, their microbiome are also seen as a powerful tool providing a diversity of chemical agents in response to stress and environmental conditions. Therefore, the exploration of new therapeutic approaches based on microorganism-derived phytocompounds or microorganisms living in plants such as bacterial and fungal (endophytes) will be a promising direction toward the discovery of transmission-blocking agents. Amongst the group of some investigated microorganisms-derived natural products, ionophores are one of the very large and important groups of naturally occurring compounds. Ionophores are highly lipophilic substances capable of interacting stoichiometrically with metal ions, thereby serving as a carrier by which these ions can be transported across a bimolecular lipid membrane [80].

Ionophores are toxic to many bacteria, protozoa, fungi, and higher organisms and thus fit the classical definition of antibiotics [81]. Motivated and inspired by the drug repurposing endeavour, D'Alessandro and coworkers (2015) screened in vitro, three ionophores specifically salinomycin, nigericin and monensin isolated from Streptomyces sp. against early and late stage of P. falciparum gametocytes [82]. Results showed that all three compounds displayed good activity against both early and late stages P. falciparum transmissible gametocytes (IC₅₀ < 200 nM) with salinomycin displaying preferential inhibition on late-stage gametocytes [82]. In addition, ionophores showed the potential to inhibit the development of P. berghei gametocytes into early sporogony stages (ESS) in vitro and the transmission-blocking properties of these compounds were confirmed in vivo using the SMFA. In addition, maduramicin, an ionophore produced by actinomycete Actinomadura rubra, has been demonstrated by the works of [83] to have transmission-blocking properties both in vitro and in vivo by killing late-stage P. falciparum gametocytes (IC₅₀< 200 nM). Killing kinetics studies showed that, maduramicin is fast-acting by reducing late-stage gametocyte viability by >90% at 12 h post-treatment with morphological change evident within 1 h following drug exposure. Additionally, in vivo, maduramicin displayed good transmission-blocking activity by significantly inhibiting oocyst development following exposure of gametocytes to the drug for only 90 min before mosquito feed [83].

Adriamycin and plicamycin, two glycosides known as a DNA synthesis inhibitors and RNA synthesis inhibitors respectively, display gametocytocidal activity at submicromolar concentrations [84]. These results underline that natural product-derived inhibitors of nucleic acid synthesis are effective against the P. falciparum transmissible stage of the malaria parasite [79]. This observation can be justified with transcription inhibitors [35, 48, 85], such as puromycin with equipotent in vitro activity against all five development stages of P. falciparum gametocytes [85] and are fast-acting against P. falciparum macro-gametes in kill kinetics assay [50] whilst also blocking P. berghei ookinetes development [86]. In addition, the macrolide chlorotonil A is highly active against late-stage gametocytes [87]. Some antibiotics such tetracycline, fosmidomycin, and deferoxamine have shown their potential against both *P. falciparum* gametocytes (IC₅₀ values > 12.5 μ M) and macro-gametes and are not able to block *P. berghei* ookinete development in vitro.

Epoxomicin, an inhibitor of P. falciparum proteasome is one of the most widely investigated peptides currently used as a reference for in vitro transmission-blocking activity targeting gametocytes stages [50, 83, 88]. Results from some studies showed that epoxomicin displayed a potent in vitro activity against late-stage P. falciparum gametocytes with IC₅₀< 10 nM [34, 36, 89] with sexspecific inhibition on P. falciparum micro-gametes [50, 90]. In vivo studies showed the potential of epoxomicin to completely block P. falciparum oocysts formation in An. stephensi [90]. Additionally, another peptide, carmaphycin B targeting the β5 subunit of the yeast 20 s proteasome have been shown to display a potent inhibition against both asexual-blood and transmissible gametocytes stages of P. falciparum in vitro (IC₅₀ < 1 µM) with 40-fold preferential inhibition on Plasmodium asexualblood stage [91, 92]. The exploration of cyclic oligopeptides including the antibiotic thiostrepton which is known as an inhibitor of P. falciparum asexual-blood stage has been shown to be moderately potent against the five development stages of gametocytes with IC₅₀ ranging from 1.82 to 3.4 µM [85] with a 14-fold enhanced activity against micro-gametes compared to macro-gametes [63]. Moreover, thiostrepton significantly reduces P. berghei oocyst development in An. stephensi mosquito midguts as well as reducing the number of sporozoites per mosquito [44]. In addition, two oligopeptides, dactinomycin and romidepsin displayed gametocytocidal activity at a sub-micromolar concentration in vitro and confirmed in vivo transmission-blocking activity for romidepsin [93]. A library of crude fungal extracts for compounds that disrupt the interaction of fibrinogen-related protein Fokou *et al. Malaria Journal* (2025) 24:164 Page 15 of 22

1 (FREP 1) with *Plasmodium* parasites allowed to identified three active extracts with *Aspergillus niger* (92% inhibition of FREP 1-*Plasmodium* association) being the most potent [94]. From this study, *P*-orlandin was identified as the active principle from this extract and showed in vivo transmission-blocking activity against oocysts formation [94]. In addition, aphidicolin, a mycotoxin from *Cephalosporum aphidicola* exhibited good activity toward *P. falciparum* micro-gametes exflagellation [95] without demonstrating any sign of toxicity.

Natural products from marine drugs

The marine environment has been demonstrated to be an interesting reservoir of compounds with unique chemical features serving as a baseline for the molecular modelling and chemical synthesis of new drugs with greater efficacy and specificity for therapeutics purposes [96]. Moreover, marine organisms do not have a distinguished history of use in traditional medicine compared to terrestrial organisms [96]. However, in the last 50 years, advances in new technologies and engineerings such as scuba diving techniques, manned submersibles, and remotely operated vehicles opened up the marine environment to scientific exploration [68] for drugs discovery. The correlation between several species in these habitats of a limited extent increases their competitiveness and complexity. Therefore, marine organisms such as seaweeds have been demonstrated to be an exceptional source of bioactive chemicals agents, some of them with different structural features from those of terrestrial sources and must be exploited for the discovery of malaria transmission-blocking agents. A set of the seaweed extracts were screened on the liver stage (LS) of the rodent malaria parasite P. berghei by using a transgenic P. berghei parasite expressing the bioluminescent reporter protein luciferase to visualize and quantify parasite development in Huh7 cells, a human hepatoma cell line. Eight seaweed extracts representing all three classes of marine macroalgae [green seaweeds Cladophora rupestris, Codium fragile ssp. tomentosoides, Ulva lactuca; brown seaweeds Cystoseira baccata, C. tamariscifolia; red seaweeds Osmundea pinnatifida, Ceramium virgatum, and Halopitys incurvus], demonstrated moderate activity against hepatic stage parasites with IC₅₀ values ranging from 14.9 to 52.9 µg/mL [97]. Intracellular localization of the liver stage parasites and the effect of the active extracts on the morphology and development of the liver stage parasites in Huh7 cells studied by immunofluorescence analysis. showed that, both extracts severely impaired the development of liver-stage parasites, which was similar to that observed with 15 μ M (= 6.8 μ g/mL) primaquine [97]. In addition, sesquiterpene avarone, its reduced form avarol, isolated from sponge Dysidea avara and thiazoavarone (semisynthetic thiazinoquinone), exhibited good activity against different stages of P. falciparum asexual and sexual forms (gametocytes stage V) (IC₅₀ = 15.53 \pm 5.26, 15.01 ± 3.19 , and 9.30 $\pm 1.90 \, \mu \text{mol/L}$) [98]. The above consideration towards marine algae opens the door for the investigation of marine sources for an anti-parasitic drugs search. Investigating marine sources emerges as a key tool in this direction. Till now, no studies have shown the transmission-blocking potential of marine extracts or derived compounds. Hence, marine algae could have appeared as a novel source of investigations for malaria transmission-blocking drugs search. It is evident from the various findings that natural compounds from plants and marine still need to be explored further whereas a significant case study has proven the potential role of natural compounds for malaria transmission-blocking.

Figure 6 illustrates the intervention points where natural compounds inhibit the development of Plasmodium, thereby blocking malaria transmission at fundamental stages in both human and mosquito hosts.

Table 4 provides a detailed summary of natural products derived from plants, microorganism-associated sources, and marine environments with demonstrated transmission-blocking activity against malaria.

Limitations and clinical gaps in malaria transmission-blocking drug discovery

The current limitations and clinical gaps in malaria transmission-blocking drug discovery pose significant challenges to advancing effective treatments. One of the primary limitations is the lack of comprehensive and standardized screening methods. Assays used to evaluate transmission-blocking efficacy vary widely and are often inconsistent, especially in their ability to measure drug effectiveness across all stages of *Plasmodium* development. This variability hinders comparisons across studies and laboratories, making it difficult to identify reliable transmission-blocking agents.

In addition, there is a scarcity of clinical validation for transmission-blocking compounds, as most research remains at the preclinical stage. Very few transmission-blocking agents have progressed to clinical trials, creating a gap between laboratory findings and real-world applications. Another critical gap lies in the specificity of existing drugs, which predominantly target asexual blood stages, leaving mature gametocytes, particularly stage V, less affected. These later-stage gametocytes are crucial for transmission but are difficult to target effectively without causing adverse effects in patients.

The safety of currently available transmission-blocking drugs, such as primaquine and tafenoquine, is also a concern. Both drugs are associated with haemolytic risks in individuals with glucose-6-phosphate dehydrogenase Fokou *et al. Malaria Journal* (2025) 24:164 Page 16 of 22

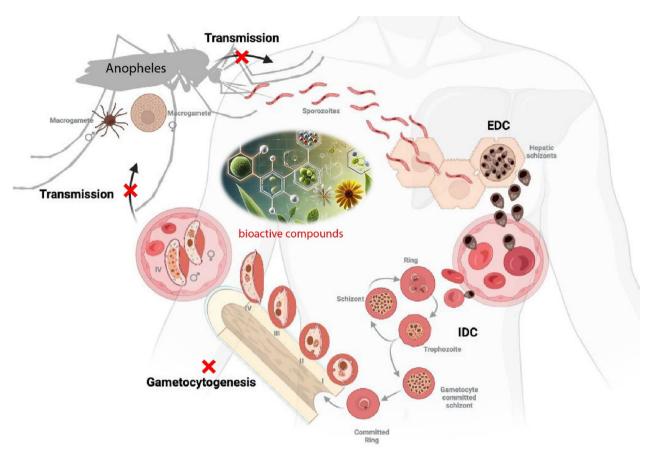


Fig. 6 Targeting malaria lifecycle stages with natural bioactive compounds for transmission blocking. Bioactive compounds derived from medicinal plants, microorganisms, and marine algae act at multiple lifecycle stages. In the hepatic (liver) stage, these compounds inhibit the maturation of hepatic schizonts, which release merozoites into the bloodstream to initiate infection. During the intraerythrocytic developmental cycle (IDC), merozoites infect red blood cells and develop through ring, trophozoite, and schizont stages, with some differentiating into gametocytes. The compounds also prevent gametocytogenesis by inhibiting the maturation of gametocytes within red blood cells, reducing the number of transmissible forms. In the mosquito vector, these compounds target the transmission stages by blocking the development of male and female gametes, thus preventing the formation of infective sporozoites and further transmission. Red X marks indicate stages where bioactive compounds interfere, blocking progression to the next stage and thereby disrupting the malaria transmission cycle

(G6PD) deficiency, which is prevalent in many malariaendemic areas. This limitation reduces the applicability of these drugs in regions where they are most needed, emphasizing the urgent need for safer transmissionblocking alternatives.

Despite the potential of natural products from sources like plants, marine organisms, and microorganisms, there has been limited exploration of these resources for malaria transmission-blocking purposes. Much of the natural world remains untapped, and systematic screening efforts are lacking, which could otherwise identify new candidates with transmission-blocking potential. Furthermore, the emergence of resistance to artemisinin-based therapies, which are central to current malaria treatments, poses a significant threat to ongoing transmission-blocking efforts. The limited availability of alternative drugs with novel mechanisms of action

exacerbates this issue, highlighting the need for new candidates that can overcome resistance.

Lastly, current research focuses primarily on humanstage gametocytes, with minimal emphasis on mosquitostage interventions, such as targeting ookinetes and oocysts. A more comprehensive approach that includes both human and mosquito stages could enhance the overall efficacy of transmission-blocking strategies. Addressing these gaps requires developing standardized, efficient screening protocols, advancing promising preclinical findings to clinical testing, expanding research into underexplored natural sources, and addressing resistance to current therapies. By tackling these limitations, malaria eradication efforts can be significantly strengthened, bringing us closer to effective, sustainable transmission-blocking solutions.

Plany: electived compounds 14,544,047,040,040,040,040,040,040,040,040,0	Category	Compound/source	Targeted stage(s)	Mechanism of action	IC ₅₀ /activity	References
Policycomme early-stage gameto- Inhibit early gametocyte (2 ₁₂ =5.5 µg/ml cyres (5-85% II-III) Policycomme early-stage Gameto- Correct Set (1-10) Policycomme early spoot- Correct Set (1-10) Policycomme early spo	Plant-derived compounds	1a,4a-Dihydroxybishopsolicepolide (Artemisia afa)		Inhibit late gametocyte	IC ₅₀ = 2 µg/ml	[109]
glartemin (Artemisia anfa) Paragraphic anty spore Pa			<i>P. falciparum</i> early-stages gameto-cytes (> 85% II-III)	Inhibit early gametocyte	IC ₅₀ = 5.6 µg/ml	
Placeparum early-stage gameto- cyces (> 85% II-III) gonic stages activity achtin (Azadirachta indica) Male gametes Activity achtin (Azadirachta indica) Male gametes Activity Male gametes Male gametes Activity Action (Azadirachta indica) Male gametes Activity Action mation; blocks exflagellation Action mation; blocks exflagellation Action mation; blocks exflagellation Action mation; blocks exflagellation Action mation for proper exflagellation Action mation for proper exflagellation Action modern in midgut Action modern in		Yomogiartemin (<i>Artemisia afra</i>)	<i>P. falciparum</i> early-stage Gameto-cytes (> 85% IV $-$ V)	Inhibit late gametocyte	IC ₅₀ = 5.3 µg/ml	[109]
idia, Vernodalol (Vernoda) gonic stages gonic stages activity			<i>P. falciparum</i> early-stage gameto-cytes (> 85% II-III)	Inhibit early gametocyte	IC ₅₀ = 4.4 µg/ml	
achtin (<i>Parathenium hysteropho-Petaretan indead pametes</i>) Reduced oocyst intensity in mos-lation) Reduced oocyst intensity in mos-lation Reduced oocyst intensity in mos-lation Reduced oocyst intensity in mos-lation Reduced oocyst intensity in mos-lation and cookinete develop-lation of proper exflagellation Reduced oocyst intensity in mos-completion Reduced oocyst intensity in mos-completion Reduced oocyst intensity in mos-lation of proper exflagellation Reduced oocyst intensity in mos-copying tail tests) Reduced oocyst intensity in mos-lation of proper exflagellation Reduced oocyst intensity in mos-lation of ookinete develop-lation of proper exflagellation Reduced oocyst development Reduced oocyst intensity in mos-lation of ookinete develop-lation of ookin		Vernolide, Vernodalol (<i>Vernonia</i> amygdalina)	Plasmodium berghei early sporo- gonic stages	Showed weak (Vernolide) to moderate (Vernodalol) inhibitory activity	$IC_{50} = 18.7 \mu M$ (Vernodalol)	[109]
noin (Parthenium hysterapho - P. falciparum ocosysts development - P. falciparum stage V gametocytes Inactivation, preventing exflagellation of proper exflagellation of pr		Azadirachtin (Az <i>adirachta indica</i>)	Male gametes	Prevents motile male gamete formation; blocks ex-flagellation	IC_{50} = 3.5 μ M (P. berghei); 100 μ M (P. falciparum)	
P. falciparum stage V gametocytes Inactivation, preventing exflagellar Inon and oocyst development ellation) P. falciparum microgametes (Exflage Inhibition of proper exflagellation and oocyst development parther P. falciparum oocysts Reduced total number of zygotes P. falciparum oocysts Reduced total number of zygotes P. falciparum oocysts Reduced oocyst intensity in mos-falciparum oocysts Reduced oocyst intensity in mos-falciparum stage V gametocytes Inhibition of proper exflagellation P. falciparum ookinetes (Exflage Inhibition of proper exflagellation of ookinete development P. falciparum ookinetes (Exflage Inhibition of proper exflagellation of P. falciparum microgametes (Exflage Inhibition of proper exflagellation of P. falciparum mature stage V gametocytes Inhibition of proper exflagellation of P. falciparum mature stage V gametocytes Inhibition of proper exflagellation of P. falciparum mature stage V plantato ochinetes in vitro on tale to tale to tale to tale to tale to tale tale to tale to tale to tale to tale tale tale tale tale tale tale tale		Parthenin (<i>Parthenium hysteropho-rus</i>)	P. falciparum oocysts	Reduced oocyst intensity in mos- quito midgut	6.25-100 µg/ml	[73]
P. falciparum microgametes (Exflage Inhibition of proper exflagellation 1000 ng/ml			P. falciparum stage V gametocytes	Inactivation, preventing exflagellation and oocyst development	100–1000 ng/ml	
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P. falciparum mature stage V Inhibits viability by allosterically IC ₅₀ = 36.7 µM (viability) binding to Plasmodium falcipa- IC ₅₀ = 33.3 µM (PfdUTPase) rum 2'-deoxyuridine 5'-triphos-phate nucleotidohydrolase (PfdUTPase), enhancing enzyme activity and inhibiting viability of both asexual and sexual stages			P. berghei Zygotes/Ookinete	Reduced total number of zygotes and ookinetes in vitro	Dose-dependent inhibition	
		Securinine (From Natural Product Fragment Library)	<i>P. falciparum</i> mature stage V gametocytes	Inhibits viability by allosterically binding to <i>Plasmodium falciparum 2'</i> -deoxyuridine 5'-triphosphate nucleotidohydrolase (PfdUTPase), enhancing enzyme activity and inhibiting viability of both asexual and sexual stages	IC ₅₀ = 36.7 µM (viability) IC ₅₀ = 33.3 µM (PfdUTPase)	[74]

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Category	Compound/source	Targeted stage(s)	Mechanism of action	IC ₅₀ /activity	References
	Pellitorine (Zanthoxylum heitzii)	Adult mosquitoes and larvae	Toxicity on adult mosquitoes and larvae	LD ₅₀ : 50 ng/mg (insect)/13 µg/ml ([9/]
	Dihydronitidine/ Heitziquinone(<i>Zanthoxylum heitzii</i>)	Plasmodium berghei ookinete	Block ookinete conversion	IC ₅₀ : 0.59/6.2 µg/ml (larvae)	[56]
	Brusatol	P. berghei Male Gametes (Stage V)	Gametocidal	IC ₅₀ value of 43.0 nM	[77]
		P. berghei Ookinete Development	Complete blocked likely through inhibition of P25 expression	10 µM (ookinete); 10 nM (p25)	
		P. berghei In Vivo Transmission	Blocks transmission to mosquitoes	0.5 mg/kg	
		<i>P. falciparum</i> gametocyte maturation	Prevents stage progression (I to II/ III, II/III to III/IV) by inhibiting development from early to late stages	100 nM, 500 nM	
		P. falciparum oocyst	Gametocidal	500 nM	
		<i>P. falciparum</i> Male gamete activity (Exflagellation)	Reduces activity	> 80% inhibition at 500 nM	
		P. falciparum Oocyst Development	Reduces burden & prevalence in mosquitoes	500 nM	
	Mallotojaponin C (<i>Mallotus japoni-</i> cus)	Plasmodium falciparum Late and Early gametocyte stages	Showed gametocytocidal activity Prevent gametocytogenesis	3.6 µM (late gametocyte) 0.14 µM (gemetogenisis)	[72]
Microorganism-Derived Com- pounds	Salinomycin, Nigericin, Monensin (<i>Streptomyces</i> sp.)	Early and late gametocytes; sporogony	lonophore action; disrupts ion transport across membranes	IC ₅₀ < 200 nM	[82]
	Maduramicin (A <i>ctinomadura rubra</i>)	Late gametocytes	Fast-acting; kills gametocytes and inhibits oocyst development	IC_{50} < 200 nM; > 90% viability reduction at 12 h	[83]
	Adriamycin, Plicamycin	Gametocytes	Inhibits nucleic acid synthesis	Sub-micromolar activity	[84]
	Epoxomicin	Late gametocytes	Inhibits P. falciparum proteasome	$IC_{50} < 10 \text{ nM}$	[34, 50]
	Carmaphycin B	Asexual stages; gametocytes	Inhibits yeast 20 s proteasome β 5 subunit	IC ₅₀ < 1 µM	[91, 92]
	Thiostrepton	Gametocytes; oocyst	Cyclic oligopeptide targeting gametocytes and mosquito midgut oocysts	$IC_{50} = 1.82 - 3.4 \mu M$	[63, 85]
	Dactinomycin, Romidepsin	Gametocytes	Inhibits transcription	Sub-micromolar activity	[63]
	P-Orlandin (<i>Aspergillus niger</i>)	Oocyst	Disrupts FREP1-Plasmodium interaction	92% inhibition of FREP1 association	[94]
	Pulixin (Purpureocillium lilacinum)	Mature stage V gametocyte	Block transmission of the parasite to mosquitoes	EC ₅₀ : 11 µМ	[78]
	Sterigmatocystin (<i>Purpureocillium lilacinum</i>)	Plasmodium sporogonic stages	Limits sexual parasite infection in mosquitoes	IC ₅₀ : 48 µM	[78]

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References 97 98 $C_{50} = 14.9 - 52.9 \, \mu g/mL$ $C_{50} = 9.30 - 15.53 \, \mu M$ C₅₀/activity mpairs liver-stage parasite devel-Mechanism of action Unknown Asexual stages; gametocytes (Stage **Targeted stage(s)** Hepatic stages Seaweed extracts (e.g., Cladophora Avarone, Avarol (Dysidea avara) rupestris, Cystoseira baccata) Compound/source Marine-derived compounds **Fable 4** (continued) Category

Conclusions and future perspectives

The quest for effective malaria transmission-blocking strategies remains crucial in the global effort to eradicate malaria. Natural products derived from medicinal plants, microorganisms, and marine sources offer a promising reservoir of bioactive compounds that can target various stages of the *Plasmodium* life cycle, especially the gametocyte and mosquito transmission stages. While initial studies have identified some promising candidates, there are still significant gaps in translating these findings into clinically effective drugs. In the future, expanding systematic screening of natural compounds using advanced bioassay-guided purification methods will be essential. Developing standardized protocols for assessing transmission-blocking efficacy across all stages of Plasmodium development can help address current inconsistencies and enhance reproducibility across laboratories. Furthermore, a focus on identifying compounds that can target mature gametocytes, particularly stage V, as well as mosquito stages (such as ookinetes and oocysts), could provide a more comprehensive approach to blocking malaria transmission. Additionally, exploring safe and effective alternatives to drugs like primaquine and tafenoquine, which have limitations due to toxicity in G6PD-deficient populations, should remain a research priority. New technologies, such as high-throughput screening and the integration of machine learning to predict active compounds, could expedite the discovery and development process. Drug repurposing, which involves identifying existing drugs with potential transmissionblocking activity, presents another valuable avenue for finding effective treatments faster. Advances in genomics and proteomics may further aid in understanding the specific mechanisms through which these natural compounds act, facilitating the development of compounds with targeted mechanisms against the malaria parasite's transmissible stages. Ultimately, a concerted effort involving multidisciplinary research, collaborations between academia, industry, and public health organizations, and increased funding will be required to move these promising findings from the laboratory to clinical use. With these advancements, the development of effective, safe, and accessible transmissionblocking agents can become a reality, strengthening global malaria eradication efforts and improving health outcomes in endemic regions.

Author contributions

PVTF, MBTT, CDJM, LRTY, JS-R, DC, R, MK, JCT, FFB made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis, and interpretation, or all these areas. That is revising or critically reviewing the article; giving final approval of the

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version to be published; agreeing on the journal to which the article has been submitted; and, confirming to be accountable for all aspects of the work.

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