<u>REVIEW ARTICLE</u>

Progress in the Study of Epidemiologic Characteristics and Influencing Factors of Asymptomatic Malaria Infection in Africa

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ABSTRACT

Malaria is caused by protozoan parasitic *Plasmodium* infections. *Plasmodium falciparum* is common in Africa; *P ovale, P malaria* and *P vivax* infections are less prevalent and globally confined, contributing to major causes of global mortality and morbidity, particularly in children in sub-Saharan African countries. In 2018, the total incidence of malaria increased from 221 million to 229 million, with an estimated 503 000 deaths reported. Sub-Saharan Africa has the highest number of cases of malaria and highest mortality rate compared with other countries, like southeastern Asia, east Pacific, western, and America with an estimated 213 million cases. In addition, continuous exposure to *Plasmodium* parasites results in the production of partial immunity to guard against more problems, resulting in asymptomatic carriers.

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INTRODUCTION

In several areas of the world, despite a vast number of national and foreign initiatives to tackle the continuing spread of malaria, it remains a significant public health issue. Malaria in tropical and subtropical regions is one of the most common human parasitic diseases, and has particular socioeconomic, public health, and quality of life significance in sub-Saharan Africa and Southeast Asia.^{1,2} Different calculations were made to estimate the global malaria burden. Upwards of 216 million malarial illnesses and 655 000 deaths were recorded by the Global Health Organization in 2010, with 106 malaria-prone countries worldwide at risk, 91 percent of which were in sub-Saharan Africa, 3% in the Eastern Mediterranean Region, and 6% in Southeast Asia.³ The diagnosis of asymptomatic malaria is not simple because of the apparent absence of clinical factors and sometimes low levels of parasites. The most basic concept appears to be parasitemia and a lack of malaria signs, primarily fever (axillary temperature <37.5° C). Thus, a better awareness of asymptomatic malaria epidemiology in affected countries will help improve strategies to reduce the local burden of malaria and its health consequences. Therefore, the objective of this study was to determine the magnitude of asymptomatic malaria pathology and related risk factors with epidemiologic characteristics in individuals on the African continent. (*Altern Ther Health Med.* [E-pub ahead of print.])

While great success has been reported in combating the disease, studies have shown limited success in reducing global malaria cases from 2014 to 2018. In 2018, the incidence of total malaria increased from 221 million to 229 million, with an estimated 503 000 deaths reported. Sub-Saharan Africa has the highest number of cases of malaria and highest mortality rates compared with other countries, like southeastern Asia, east Pacific, western, and America with an estimated 213 million cases of malaria.⁴ Furthermore, 74% of Kenya's population is confirmed to be at risk for malaria and the most heavily affected areas are the coastal areas and the west of the world.⁴ Malaria is caused by protozoan parasite Plasmodium infections. In Africa, P falciparum is common, while P vivax, P ovale and *P* malaria infections are less prevalent and globally limited.^{5,6} *P* falciparum malaria remains a major cause of global death and morbidity, especially among children in sub-Saharan Africa.7,8

Plasmodium parasites are transmitted by *Anopheles* mosquitoes, with *A funestus*, *A arabiensis* and *A gambiae sensu stricto* being the most prevalent in Africa.⁹ In addition, continuous exposure to *Plasmodium* parasites results in partial immunity, which guards against more problems and

thus results in asymptomatic carriers.¹⁰ Studies from Kenya, Uganda and Brazil have shown that asymptomatic parasitemia is 6 to 7 times more prevalent than polymerase chain reaction (PCR) when contrasted with microscopy.¹¹⁻¹³ Many asymptomatic disorders have few to no signs and thus go undetected and untreated. The level of asymptomatic parasitemia is inversely correlated with a clinical diseaseprone community.¹⁴ More specifically, asymptomatic individuals are significant sources of infection.¹⁵ Although the use of PCR is difficult, costly and impractical in most field trials, it is necessary to increase the precision of asymptomatic parasitemia diagnoses.¹⁶ However, scientists and clinicians have developed diagnostic recommendations based on the initiation of clinical events in malaria that helped develop an organized approach to strengthening serious malaria management and treatment.¹⁷ Diagnosis of asymptomatic malaria is not simple because of the apparent absence of clinical events and sometimes low levels of parasites.¹⁸ Thus, a better awareness of asymptomatic malaria epidemiology in affected countries will help improve strategies to reduce the local disease burden and health effects of malaria.¹⁹ Therefore, the objective of this study is to determine the magnitude of asymptomatic malaria pathology and related risk factors and epidemiologic characteristics among individuals on the African continent.

LITERATURE SELECTION

Potentially important English-language only studies were culled from ScienceDirect, Medline, PubMed, Public Library of Science, Mendeley, SpringerLink and Google Scholar. Many keywords were used to scan the literature including "sepsis," "epidemiology of malaria," "malariaassociated hyper inflammation," "mechanism of the pathogenesis of asymptomatic malaria," "involvement of epidemiological characteristic of asymptomatic malaria in African continent," "prevalence of asymptomatic malaria," "asymptomatic malaria and anti-inflammatory immunity," "asymptomatic malaria and antibodies," "factor influencing asymptomatic malaria," "immunosuppressive pathway via Plasmodium falciparum in African continent" and "innate and adaptive immunity in asymptomatic malaria." Reference lists of collected papers were also screened for related articles not found with the initial search strategy.

EPIDEMIOLOGIC CHARACTERISTICS OF ASYMPTOMATIC MALARIA

Recent declines in the incidence of clinical malaria show that asymptomatic parasite carriers are essential to maintaining transmission.²⁰⁻²⁴ In addition, there are claims that asymptomatic carrier parasites are more dangerous than symptomatic carrier parasites.²⁵ The most basic concept appears to be parasitemia and the lack of malaria signs, primarily fever (axillary temperature <37.5° C).²⁶⁻²⁸ Healthy individuals can carry asexual and sexual blood parasites (gametocytes), which are essential for transmission.²⁹ Individuals with persistent asymptomatic infections may contribute to the spread of viruses in the subsequent years.²⁹ Research performed in Mali showed that targeting asymptomatic infections in any chronically symptomatic patient is unlikely to have an impact on clinical malaria prevalence rates.³⁰ Moreover, malaria transmission results in Africa such as in The Gambia, Mali and Senegal suggest that more than 25% of people with sub-microscopic gametocytes are capable of contaminating mosquitoes.³¹ A 2012 study from Laos showed that 20% (175 out of 888) of officially stable individuals had *Plasmodium* infections.³¹ Asymptomatic parasitemia is associated with a greater risk for fever caused by malaria in children up to 18 months of age.³² Older children in areas where malaria is prevalent, however, acquire parasite-specific immunity to the Plasmodium parasite that helps reduce the volume and occurrence of asymptomatic parasite carriers in clinical malaria episodes.³³⁻³⁵ Moreover, schoolchildren with asymptomatic malaria may not be out of school because of a clinical malaria episode but may be a significant reservoir of disease transmission.^{15,36,37} Patients with non-specific signs, such as fever, rigors, and chills, would not require hospital admission. In a minority of patients, extreme malaria can present as fever, diminished awareness, serious anemia, respiratory failure, convulsions, and hypoglycemia, among other symptoms.³⁸ Often such signs are not observed and patients with no recent history of antimalarial care have been diagnosed with infection.29

One important concern remains: Why do P falciparum infections sometimes remain asymptomatic in people with little or poor immunity without producing large parasite densities? A new theory is that it may be that parasites that are less virulent in low-endemic settings are not subject to strong competition from virulent strains, thereby remaining lowdensity, asymptomatic, undetected and untreated, enabling further transmission over a long period.³⁹ However, another hypothesis is that whenever a parasite is present, immune factors are needed to reduce parasite numbers-antiparasite immunity—and antidisease immunity is needed to avoid signs of clinical complications.⁴⁰ Immunity in asymptomatic individuals is adversely affected by antidisease immunity rather than antiparasite immunity. The reasons for this phenomenon remain unknown, and further investigations are needed to clarify how antidisease immunity is triggered, which is explored in the next section of this study (see Figure 1).

MOLECULAR EXPLORATION OF PATHOLOGY OF ASYMPTOMATIC MALARIA

The occurrence and longevity of asymptomatic infections is a dynamic phenomenon linked to degrees of defensive immunity with recurrent malaria exposure and immune system maturity.⁴¹ At the genome level, variations in host gene expression can account for numerous clinical manifestations during host-parasite interactions.⁴² Specifically, gene pathways that control and augment cytokine signals, as well as immunoglobulin output, have been shown to be involved.⁴³ In several studies, an elevated incidence of febrile or cerebral malaria was correlated with an aggressive

pro-inflammatory response, whereas asymptomatic infection was associated with a poor response.44-46 P falciparum infections are distinguished by the co-circulation of many P falciparum clones in acute and chronic infections in malaria-endemic regions, especially in sub-Saharan Africa.47-48 This is considered the anomaly mechanism of injury (MOI). The distribution of some genotypes of MSP 1 was correlated with different clinical malaria manifestations. Msp1 K1 and MAD20 were associated with asymptomatic malaria with minimal chance of fever in young Nigerian children.⁴⁹ Other findings show that high MOI in children with asymptomatic infection is associated with an increased rate of febrile malaria in infants, with a lower risk in older children.^{50,51} Thus, MOI and *P falciparum* genetic variations are significant alternative markers in determining the severity of malaria transmission in various endemic regions. The conventional P falciparum characterization approach utilizes gel electrophorese entangled polymer chair reactions (PCRs) to identify polymorphisms in msp1, msp2 and rich protein in glutamines (GLuRP).52 However, the use of molecular genotyping of polymorphic anti-genetic markers msp1, msp2 and GLuRP for studies of parasite diversity is criticized for providing high immune selection involving these genes.⁵³⁻⁵⁶

People in areas in which malaria is endemic are often asymptomatic and clinically resistant to diseases from numerous genetically complicated *P* falciparum infections over time.⁵⁷ Children with repeated episodes of malaria often have a modified immune system,^{58,59} which is attributed to enhanced development of B cells, interleukin (IL)-10, activation of neutrophils and CD8 + T cells.60 Immune responses are regulated by cytokines that control inflammation such as tumor necrosis factor alpha (TNF-a), interferon gamma (IFN-y), IL-12, thus providing defensive immunity.⁶¹ However, overstimulation of the immune system results in additional cytokine output and immune cell activation, which further inhibits the host defensive mechanism to parasitemia.^{62,63} Also, anti-inflammatory cytokines like transforming growth factor beta (TGF-β), IL-10 and IL-27 are active in the dampening of pro-inflammatory cytokines to reduce disease intensity.^{61,64} The existence of antiinflammatory cytokines, particularly IL-10, prevents the clearance of parasites, impedes the production of antiparasite immunity and promotes the development of asymptomatic infections.^{65,66} The risk of an infection being asymptomatic rises with age, as frequent malaria exposure contributes to partial antidisease immunity.⁶⁷ Furthermore, increased IL-10 levels have also been correlated with asymptomatic infections in pregnant women.68 In research in young Ugandan children, CD4 + T cytokines were found to be affected by previous malaria infection exposure. CD4 + T cells in more exposed children had higher levels of IL-10, while less exposed children had higher TNF-a levels and, therefore, inflammation. The activation of pro-inflammatory cytokine (TNF α , IL-1 β and IFN- γ) development was reported in a malaria transcriptomic analysis with the least activation of asymptomatic cytokines.²⁷ Also, studies have shown that elevated regulatory T cell levels were related to increased parasitemia, the expression of TGF- β and clinical symptoms⁶⁹⁻⁷¹ while lower levels can reduce the risk of developing symptoms, resulting in immunity to antidisease.72 Thus, antibodies against malaria tend to be short-lived since malaria infection cannot produce a sufficient B-cell response to the antigen.⁷³ In comparison, a report on Swedish travelers formerly treated for malaria preserved long-lasting B memory cells for 16 years with no exposure afterwards.⁷⁴ In asymptomatic patients, higher immunoglobulin G antigenspecific titers were found compared with patients with other findings.^{75,76} In Gabonese children with symptomatic infection with *P* falciparum, elevated antigen-specific antibody responses have been associated with high IFN-a and IL-10 levels, suggesting a protective immune system response.77 Thus, asymptomatic infections were correlated with a lack of TNF-a development.⁴⁴ On the other hand, a 2014 study explored the high activity of V δ 2+ $\gamma\delta$ T cells due to the low level of pro-inflammatory cytokines⁷⁸ and increased immunoregulatory gene expression possibly dampening the effects of associated infections.

FACTORS INFLUENCING ASYMPTOMATIC MALARIA

Various factors including age, sex, education level, use of mosquito netting around the bed, the vector-host ratio during the wet season and recurrent exposure to malaria have been correlated with risk factors in different studies.⁷⁹ The gender variation in asymptomatic malaria may be attributable to samples from various seasons and socio-economic classes, prior access to prevention and recovery and the presence or absence of mosquito reproductive sites.

The investigators suggested that men are more vulnerable to asymptomatic infections than women. Furthermore, various studies have shown that the gender distribution of asymptomatic malaria was 9.8% male and 5.1% female.⁸⁰ The incidence in males was 48.1% and in females 41.4% compared with the findings from a 2012 study carried out in Cameroon.⁸¹ Age is one of the most important variables in malariaendemic areas linked to the defensive immune response. Many previous studies have shown that adults are asymptomatic parasite carriers because they are highly protected due to frequent exposure to malaria parasites, whereas viruses are symptomatic in young children while their protection against malaria continues to grow.^{29,82,83} Thus, discovering that schoolchildren (younger than age 15 years) had an elevated risk of asymptomatic malaria infections appears to refute African results. However, this observation did reinforce the findings of earlier studies that schoolchildren have an increased risk for acute P vivax malaria.84,85

The results of vector-based prevention interventions were noteworthy and the data revealed that not sleeping under mosquito nets resulted in a more than 5 times higher probability of developing asymptomatic infections.⁸⁶ Moreover, in many villages in which the inhabitants did not use mosquito nets, there was an increased chance of developing asymptomatic *Plasmodium* infection compared

with individuals who used mosquito nets. Of note, seasonal heterogeneity in small cities was clearer, due to different vector ecology and efforts to monitor this in small cities and encampments.⁸⁵ Moreover, the materials used for house construction, which are reflective of families' socioeconomic position, also demonstrated major effects on asymptomatic infections; residents of wood or bamboo houses were found to have almost double the risk of *Plasmodium* infections (see Figure 2).⁸⁷⁻⁸⁹

The transmission of malaria is influenced by 2 key variables: vector density and gametocyte infectivity. Effective vector-based malaria prevention strategies such as the utilization of indoor residual spraying and mosquito netting, especially long-lasting insecticide-treated netting, are therefore essential for reducing malaria occurrence.⁹⁰ In addition, hidden asymptomatic infections, which provide a reservoir for local malaria transmission, are one of the key challenges to achieving this goal. Any asymptomatic pathogens may be critically charged with drug-resistant genes.⁹¹

CONCLUSION

In areas in which malaria is hyper-prevalent, antimosquito initiatives only function effectively because of the reservoir that has developed in susceptible people who do not even realize they bear the parasite. In high-transmission zones, asymptomatic rather than symptomatic malaria prevails. In sub-Saharan Africa, 24 million people are reported to have asymptomatic malaria infections, including 38% to 50% of school-age children in West Kenya. Of the 219 million malaria infections noted globally in 2017, more than 90% were in sub-Saharan Africa. The number of asymptomatic malaria reports, particularly with P falciparum, underscores the underlying problem facing the malaria elimination program in Africa. Besides vector management and the care of symptomatic and asymptomatic cases, active and inactive case identification must be utilized to eliminate malaria. Accurate monitoring via molecular assay and valid serologic methods in asymptomatic individuals and submicroscopic cases of malaria, particularly in areas with seasonal and low transmission rates, can be very helpful. This suggests that the burden of asymptomatic malaria needs to be further measured using a more adaptive approach, taking into account the various age groups and the extent of malaria transmission to improve the elimination and worldwide eradication of malaria.

CONFLICT OF INTEREST

All the investigators state that they have no conflict of interest.

REFERENCES

- 1. Hochman S, Kim K. The Impact of HIV and malaria coinfection: What is known and suggested venues for further study. Interdisciplin *Perspect Infect Dis.* 2009;2009:617954.
- Minakawa N, Omukunda E, Zhou G, Githeko A, Yan G. Malaria vector productivity in relation to the highland environment in Kenya. *Amer J Trop Med Hygiene*. 2006;75:448-453.

- Manuel Ramos J, Reyes F, Tesfamariam A. 2006. Change in epidemiology of malaria infections in a rural area in Ethiopia. J Travel Med. 2006;12:155-156.
- Touray AO, Mobegi VA, Wamunyokoli F, Herren JK. 2020. Diversity and Multiplicity of P. falciparum infections among asymptomatic school children in Mbita, Western Kenya. *Scientific Reports* 2020;10:5924.
- Howes RE, Reiner RC, Jr, Battle KE, et al. 2015. Plasmodium vivax transmission in Africa. PLoS NTD, 2015;9:e0004222.
- Roucher C, Rogier C, Sokhna C, Tall A, Trape JF, A 20-year longitudinal study of Plasmodium ovale and Plasmodium malariae prevalence and morbidity in a West African population. *PloS One*. 2014;9:e87169.
- Liu L, Oza S, Hogan D, et al. Global, regional, and national causes of under-5 mortality in 2000-15: an updated systematic analysis with implications for the Sustainable Development Goals. *Lancet.* 2014;388:3027-3035.
- Snow RW, Omumbo JA, Lowe B, et al. Relation between severe malaria morbidity in children and level of Plasmodium falciparum transmission in Africa. *Lancet*. 1997;349: 1650-1654.
- Sinka ME, Bangs MJ, Manguin S, et al. The dominant Anopheles vectors of human malaria in Africa, Europe and the Middle East: occurrence data, distribution maps and bionomic précis. *Parasits Vectors*. 2010;3:117.
- Bousema JT, Gouagna LC, Drakeley CJ, et al. Plasmodium falciparum gametocyte carriage in asymptomatic children in western Kenya. *Malaria J*. 2004;3:18.
- Alves FP, Durlacher RR, Menezes MJ, Krieger H, Silva LH, Camargo EP. High prevalence of asymptomatic Plasmodium vivax and Plasmodium falciparum infections in native Amazonian populations. *Am J Trop Med Hyg.* 2002;66:641-648.
- Idris ZM, Chan CW, Kongere J, et al. High and heterogeneous prevalence of asymptomatic and sub-microscopic malaria infections on islands in Lake Victoria, Kenya. Sci Rep. 2016;6:36958.
- Nsobya SL, Parikh S, Kironde F, et al. Molecular evaluation of the natural history of asymptomatic parasitemia in Ugandan children. J Infect Dis. 2004;89:2220-2226.
- Mwangi TW, Fegan G, Williams TN, et al. Evidence for over-dispersion in the distribution of clinical malaria episodes in children. *PloS One*. 2018;3:e2196.
- Alves FP, Gil LH, Marrelli MT, et al. Asymptomatic carriers of Plasmodium spp. as infection source for malaria vector mosquitoes in the Brazilian Amazon. J Med Entomol. 2005;42:777-779.
- Kijogi C, Kimura D, Bao LQ, et al. Modulation of immune responses by Plasmodium falciparum infection in asymptomatic children living in the endemic region of Mbita, western Kenya. *Parasitol Intern*. 2018;67:284-293.
- Bottius E, Guanzirolli A, Trape J-F, Rogier C, Konate L, Druilhe P. Malaria: even more chronic in nature than previously thought; evidence for subpatent parasitaemia detectable by the polymerase chain reaction. Trans R Soc Trop Med Hyg. 1996;90:15-19.
- Trape JF, Zoulani A, Quinet MC. Assessment of the incidence and prevalence of clinical malaria in semi-immune cgildren exposed to intense and perennial transmission. *Am J Epidemiol.* 1987;126:193-201.
- Rabinovich RN, Drakeley C, Djimde AA, et al. 2017. malERA: An updated research agenda for characterising the reservoir and measuring transmission in malaria elimination and eradication. *PLoS Med.* 2017;14:e1002452.
- Iwagami M, Keomalaphet S, Khattignavong P, et al. 2017. The detection of cryptic Plasmodium infection among villagers in Attapeu province, Lao PDR. *PLoS* NTD. 2017;11:e0006148.
- Lover AA, Dantzer E, Hongvanthong B, et al. Prevalence and risk factors for asymptomatic malaria and genotyping of glucose 6-phosphate (G6PD) deficiencies in a vivax-predominant setting, Lao PDR: implications for subnational elimination goals. *Malar J.* 2018;17:218.
- Niang M, Thiam LG, Sane R, et al. Substantial asymptomatic submicroscopic Plasmodium carriage during dry season in low transmission areas in Senegal: Implications for malaria control and elimination. *PloS One*. 2017;12:e0182189.
- Sáenz FE, Arévalo-Cortés A, Valenzuela G, et al. Malaria epidemiology in lowendemicity areas of the northern coast of Ecuador: high prevalence of asymptomatic infections. *Malar J.* 2017;16:300.
- von Seidlein L, Dondorp A. Fighting fire with fire: mass antimalarial drug administrations in an era of antimalarial resistance. *Expert Rev Anti Infect Ther*. 2015;13:715-730.
- Laishram DD, Sutton PL, Nanda N, et al. The complexities of malaria disease manifestations with a focus on asymptomatic malaria. *Malaria J.* 2012;11:29.
- Botwe AK, Asante KP, Adjei G, Assafuah S, Dosoo D, Owusu-Agyei S. Dynamics in multiplicity of Plasmodium falciparum infection among children with asymptomatic malaria in central Ghana. *BMC Genet.* 2017;18:67-67.
- Tran TM, Jones MB, Ongoiba A, et al. Transcriptomic evidence for modulation of host inflammatory responses during febrile Plasmodium falciparum malaria. Sci Rep. 2016;6:31291.
- Wamae K, Wambua J, Nyangweso G, et al. Transmission and age impact the risk of developing febrile malaria in children with asymptomatic Plasmodium falciparum parasitemia. J Infect Dis. 2019;219:936-944.
- Lindblade KA, Steinhardt L, Samuels A, Kachur SP, Slutsker L. 2013. The silent threat: asymptomatic parasitemia and malaria transmission. *Exp Rev Anti Infect Ther.* 2013;11:623-639.

- Portugal S, Tran TM, Ongoiba A, et al. Treatment of chronic asymptomatic plasmodium falciparum infection does not increase the risk of clinical malaria upon reinfection. *Clin Infect Dis.* 2017;64:645-653.
- Bousema T, Dinglasan RR, Morlais I, et al. Mosquito feeding assays to determine the infectiousness of naturally infected Plasmodium falciparum gametocyte carriers. *PloS One*. 2012;7:e42821.
- Wamae K, Wambua J, Nyangweso G. Transmission and age impact the risk of developing febrile malaria in children with asymptomatic Plasmodium falciparum parasitemia. J Infect Dis. 2019;219:936-944.
- Garn SM, Leonard WR. Malnutrition and weight loss in patients with AIDS. Nutr Rev. 1989:47:354-356.
- 34. Ben-Tovim N. Cylinder dioptometry. Arch Ophthalmol. 1970;84:260-271.
- Males S, Gaye O, Garcia A. Long-term asymptomatic carriage of Plasmodium falciparum protects from malaria attacks: A prospective study among Senegalese children. *Clin Infect Dis.* 2008;46:516-522.
- Coalson JE, Walldorf JA, Cohee LM, et al. High prevalence of Plasmodium falciparum gametocyte infections in school-age children using molecular detection: patterns and predictors of risk from a cross-sectional study in southern Malawi. *Malaria J.* 2016;15:527.
- Walldorf JA, Cohee LM, Coalson JE, et al. School-age children are a reservoir of malaria infection in Malawi. *PloS One*, 2015;10:e0134061.
- Marsh K, Forster D, Waruiru C, Mwangi I, Winstanley M, Marsh V, Newton C, Winstanley P, Warn P, Peshu N, et al. Indicators of life-threatening malaria in African children. N Engl J Med, 1995;332:1399-1404.
- Björkman AB. Asymptomatic low-density malaria infections: a parasite survival strategy? *The Lancet Infectious Diseases*. 2018;18:485-486.
- Ademolue TW, Awandare GA. Evaluating antidisease immunity to malaria and implications for vaccine design. *Immunology*. 2018;153:423-434.
- Rodriguez-Barraquer I, Arinaitwe E, Jagannathan P, Kamya MR, Rosenthal PJ, Rek J, Dorsey G, Nankabirwa J, Staedke SG, Kilama M, Drakeley C, Ssewanyana I, Smith DL, Greenhouse B. Quantification of anti-parasite and anti-disease immunity to malaria as a function of age and exposure. *eLife* 2018;7.
- 42. Almelli T, Nuel G, Bischoff E, Aubouy A, Elati M, Wang CW, Dillies MA, Coppée JY, Ayissi GN, Basco LK, Rogier C, Ndam NT, Deloron P, Tahar R. Differences in gene transcriptomic pattern of Plasmodium falciparum in children with cerebral malaria and asymptomatic carriers. *PloS one* 2014;9:e114401.
- Boldt ABW, van Tong H, Grobusch MP, Kalmbach Y, Dzeing Ella A, Kombila M, Meyer CG, Kun JFJ, Kremsner PG, Velavan TP. The blood transcriptome of childhood malaria. *EBioMedicine*. 2019;40:614-625.
- 44. Jagannathan P, Eccles-James I, Bowen K, Nankya F, Auma A, Wamala S, Ebusu C, Muhindo MK, Arinaitwe E, Briggs J, Greenhouse B, Tappero JW, Kamya MR, Dorsey G, Feeney ME. IFNγ/IL-10 co-producing cells dominate the CD4 response to malaria in highly exposed children. *PLoS pathogens*. 2014;10:e1003864.
- Othoro C, Lal AA, Nahlen B, Koech D, Orago AS, Udhayakumar V. A low interleukin-10 tumor necrosis factor-alpha ratio is associated with malaria anemia in children residing in a holoendemic malaria region in western Kenya. J Infect Dis. 1999;179:279-282.
- Udomsangpetch R, Chivapat S, Viriyavejakul P, Riganti M, Wilairatana P, Pongponratin E, Looareesuwan S. Involvement of cytokines in the histopathology of cerebral malaria. Am J Trop Med Hyg. 1997;57:501-506.
- Babiker HA, Ranford-Cartwright LC, Currie D, Charlwood JD, Billingsley P, Teuscher T, Walliker D. Random mating in a natural population of the malaria parasite Plasmodium falciparum. *Parasitology*. 1994;109(Pt 4):413-421.
- Bereczky S, Liljander A, Rooth I, Faraja L, Granath F, Montgomery SM, Färnert A. Multiclonal asymptomatic Plasmodium falciparum infections predict a reduced risk of malaria disease in a Tanzanian population. *Microbes Infect* 2007;9:103-110.
- Amodu OK, Adeyemo AA, Ayoola OO, Gbadegesin RA, Orimadegun AE, Akinsola AK, Olumese PE, Omotade OO. Genetic diversity of the msp-1 locus and symptomatic malaria in south-west Nigeria. *Acta tropica*, 2005;95:226-232.
- Felger I, Smith T, Edoh D, Kitua A, Alonso P, Tanner M, Beck HP. Multiple Plasmodium falciparum infections in Tanzanian infants. *Trans R Soc Trop Med Hyg.* 1999;93(Suppl 1):29-34.
- Mayor A, Saute F, Aponte JJ, Almeda J, Gómez-Olivé FX, Dgedge M, Alonso PL. Plasmodium falciparum multiple infections in Mozambique, its relation to other malariological indices and to prospective risk of malaria morbidity. *Trop Med Int Health*. 2003;8:3-11.
- Snounou G. Genotyping of Plasmodium spp. Nested PCR. Methods Mol Med. 2002;72:103-116.
- 53. Chen JT, Li J, Zha GC, Huang G, Huang ZX, Xie DD, Zhou X, Mo HT, Eyi JUM, Matesa RA, Obono MMO, Li S, Liu XZ, Lin M. Genetic diversity and allele frequencies of Plasmodium falciparum msp1 and msp2 in parasite isolates from Bioko Island, Equatorial Guinea. *Malar J.* 2018;17:458.
- Mohd Abd Razak MR, Sastu UR, Norahmad NA, Abdul-Karim A, Muhammad A, Muniandy PK, Jelip J, Rundi C, Imwong M, Mudin RN, Abdullah NR. Genetic Diversity of Plasmodium falciparum Populations in Malaria Declining Areas of Sabah, East Malaysia. *PloS one*. 2016;11:e0152415.
- Nkhoma SC, Trevino SG, Gorena KM, Nair S, Khoswe S, Jett C, Garcia R, Daniel B, Dia A, Terlouw DJ, Ward SA, Anderson TJC, Cheeseman IH. Resolving withinhost malaria parasite diversity using single-cell sequencing. *bioRxiv*. 2018;391268.

- Turner D, Williams S, Heavner J. Pleural permeability to local anesthetics--the influence of concentration, pH, and local anesthetic combinations. *Regional anesthesia*. 1989;14:128-132.
- Bull PC, Marsh K. The role of antibodies to Plasmodium falciparum-infectederythrocyte surface antigens in naturally acquired immunity to malaria. *Trends Microbiol.* 2002;10:55-58.
- Day KP, Marsh K. Naturally acquired immunity to Plasmodium falciparum. Immunol Today. 1991;12:A68-71.
- Greenwood BM, Bradley-Moore AM, Bryceson AD, Palit A. Immunosuppression in children with malaria. *Lancet.* 1972;1:169-172.
- 60. Bediako Y, Adams R, Reid AJ, Valletta JJ, Ndungu FM, Sodenkamp J, Mwacharo J, Ngoi JM, Kimani D, Kai O, Wambua J, Nyangweso G, de Villiers EP, Sanders M, Lotkowska ME, Lin JW, Manni S, Addy JWG, Recker M, Newbold C, Berriman M, Bejon P, Marsh K, Langhorne J. Repeated clinical malaria episodes are associated with modification of the immune system in children. *BMC medicine*. 2019;17:60.
- Malaguarnera L, Musumeci S. The immune response to Plasmodium falciparum malaria. *Lancet Infectious diseases*. 2002;2:472-478.
- Butler NS, Harris TH, Blader IJ. Regulation of immunopathogenesis during Plasmodium and Toxoplasma infections: more parallels than distinctions? *Trends Parasitol.* 2013;29:593-602.
- 63. Lyke KE, Burges R, Cissoko Y, Sangare L, Dao M, Diarra I, Kone A, Harley R, Plowe CV, Doumbo OK, Sztein MB. Serum levels of the proinflammatory cytokines interleukin-1 beta (IL-1beta), IL-6, IL-8, IL-10, tumor necrosis factor alpha, and IL-12(p70) in Malian children with severe Plasmodium falciparum malaria and matched uncomplicated malaria or healthy controls. *Infect Immun.* 2004;72:5630-5637.
- 64. Findlay EG, Greig R, Stumhofer JS, Hafalla JC, de Souza JB, Saris CJ, Hunter CA, Riley EM, Couper KN. Essential role for IL-27 receptor signaling in prevention of Th1-mediated immunopathology during malaria infection. *J Immunol.* 2010;185:2482-2492.
- Kumar R, Ng S, Engwerda C. The Role of IL-10 in Malaria: A Double Edged Sword. Front Immunol. 2019;10:229.
- 66. Portugal S, Moebius J, Skinner J, Doumbo S, Doumtabe D, Kone Y, Dia S, Kanakabandi K, Sturdevant DE, Virtaneva K, Porcella SF, Li S, Doumbo OK, Kayentao K, Ongoiba A, Traore B, Crompton PD. Exposure-dependent control of malaria-induced inflammation in children. *PLoS pathogens*. 2014;10:e1004079.
- Ladeia-Andrade S, Ferreira MU, de Carvalho ME, Curado I, Coura JR. Agedependent acquisition of protective immunity to malaria in riverine populations of the Amazon Basin of Brazil. *Am J Trop Med Hyg.* 2009;80:452-459.
- Wilson NO, Bythwood T, Solomon W, Jolly P, Yatich N, Jiang Y, Shuaib F, Adjei AA, Anderson W, Stiles JK. Elevated levels of IL-10 and G-CSF associated with asymptomatic malaria in pregnant women. Infectious diseases in obstetrics and gynecology. 2010.
- 69. Boyle MJ, Jagannathan P, Farrington LA, Eccles-James I, Wamala S, McIntyre TI, Vance HM, Bowen K, Nankya F, Auma A, Nalubega M, Sikyomu E, Naluwu K, Rek J, Katureebe A, Bigira V, Kapisi J, Tappero J, Muhindo MK, Greenhouse B, Arinaitwe E, Dorsey G, Kamya MR, Feeney ME. Decline of FoxP3+ Regulatory CD4 T Cells in Peripheral Blood of Children Heavily Exposed to Malaria. *PLoS pathogens*. 2015;11:e1005041.
- Minigo G, Woodberry T, Piera KA, Salwati E, Tjitra E, Kenangalem E, Price RN, Engwerda CR, Anstey NM, Plebanski M. Parasite-dependent expansion of TNF receptor II-positive regulatory T cells with enhanced suppressive activity in adults with severe malaria. *PLoS pathogens*. 2009;5:e1000402.
- Walther M, Tongren JE, Andrews L, Korbel D, King E, Fletcher H, Andersen RF, Bejon P, Thompson F, Dunachie SJ, Edele F, de Souza JB, Sinden RE, Gilbert SC, Riley EM, Hill AV. Upregulation of TGF-beta, FOXP3, and CD4+CD25+ regulatory T cells correlates with more rapid parasite growth in human malaria infection. *Immunity*. 2005;23:287-296.
- Frimpong A, Kusi KA, Tornyigah B, Ofori MF, Ndifon W. Characterization of T cell activation and regulation in children with asymptomatic Plasmodium falciparum infection. *Malar J.* 2018;17:263.
- Dorfman JR, Bejon P, Ndungu FM, Langhorne J, Kortok MM, Lowe BS, Mwangi TW, Williams TN, Marsh K. B cell memory to 3 Plasmodium falciparum bloodstage antigens in a malaria-endemic area. *J Infect Dis.* 2005;191:1623-1630.
- Ndungu FM, Lundblom K, Rono J, Illingworth J, Eriksson S, Färnert A. Longlived Plasmodium falciparum specific memory B cells in naturally exposed Swedish travelers. *Eur J Immunol.* 2013;43:2919-2929.
- Braga EM, Barros RM, Reis TA, Fontes CJ, Morais CG, Martins MS, Krettli AU. Association of the IgG response to Plasmodium falciparum merozoite protein (C-terminal 19 kD) with clinical immunity to malaria in the Brazilian Amazon region. *Am J Trop Med Hyg.* 2002;66:461-466.
- Kinyanjui SM, Mwangi T, Bull PC, Newbold CI, Marsh K. Protection against clinical malaria by heterologous immunoglobulin G antibodies against malariainfected erythrocyte variant surface antigens requires interaction with asymptomatic infections. J Infect Dis. 2004;190:1527-1533.
- Guiyedi V, Bécavin C, Herbert F, Gray J, Cazenave PA, Kombila M, Crisanti A, Fesel C, Pied S. Asymptomatic Plasmodium falciparum infection in children is associated with increased auto-antibody production, high IL-10 plasma levels and antibodies to merozoite surface protein 3. *Malar J.* 2015;14:162.

- 78. Jagannathan P, Kim CC, Greenhouse B, Nankya F, Bowen K, Eccles-James I, Muhindo MK, Arinaitwe E, Tappero JW, Kamya MR, Dorsey G, Feeney ME. Loss and dysfunction of V $\delta 2^+ \gamma \delta$ T cells are associated with clinical tolerance to malaria. *Sci Transl Med.* 2014;6:251ra117.
- Maziarz M, Nabalende H, Otim I, Legason ID, Kinyera T, Ogwang MD, Talisuna AO, Reynolds SJ, Kerchan P, Bhatia K, Biggar RJ, Goedert JJ, Pfeiffer RM, Mbulaiteye SM. A cross-sectional study of asymptomatic Plasmodium falciparum infection burden and risk factors in general population children in 12 villages in northern Uganda. *Malar J*. 2018;17:240.
- Worku L, Damtie D, Endris M, Getie S, Aemero M. Asymptomatic Malaria and Associated Risk Factors among School Children in Sanja Town, Northwest Ethiopia. *International Scholarly Research Notices*. 2014:303269.
- Kimbi H, Ajeagah H, Keka F, Lum E, Nyabeyeu H, Tonga C, Gah A, Lehman L. An Update of Asymptomatic Falciparum Malaria in School Children in Muea, Southwest Cameroon. J Bacteriol Parasitol. 2012;3.
- Doolan DL, Dobaño C, Baird JK. Acquired immunity to malaria. *Clin Microbiol Rev.* 2009;22:13-36.
- Jagannathan P, Muhindo MK, Kakuru A, Arinaitwe E, Greenhouse B, Tappero J, Rosenthal PJ, Kaharuza F, Kamya MR, Dorsey G. Increasing incidence of malaria in children despite insecticide-treated bed nets and prompt anti-malarial therapy in Tororo, Uganda. *Malar J.* 2012;11:435.
- Li N, Parker DM, Yang Z, Fan Q, Zhou G, Ai G, Duan J, Lee MC, Yan G, Matthews SA, Cui L, Wang Y. Risk factors associated with slide positivity among febrile patients in a conflict zone of north-eastern Myanmar along the China-Myanmar border. *Malar J.* 2013;12:361.
- Zhou G, Lo E, Zhong D, Wang X, Wang Y, Malla S, Lee MC, Yang Z, Cui L, Yan G. Impact of interventions on malaria in internally displaced persons along the China-Myanmar border: 2011-2014. *Malar J.* 2016;15:471.
- Zhao Y, Zeng J, Zhao Y, Liu Q, He Y, Zhang J, Yang Z, Fan Q, Wang Q, Cui L, Cao Y. Risk factors for asymptomatic malaria infections from seasonal cross-sectional surveys along the China–Myanmar border. *Malar J*. 2018;17:247.
- Alemu A, Tsegaye W, Golassa L, Abebe G. Urban malaria and associated risk factors in Jimma town, south-west Ethiopia. *Malar J.* 2011;10:173.
- Hiscox A, Khammanithong P, Kaul S, Sananikhom P, Luthi R, Hill N, Brey PT, Lindsay SW. Risk factors for mosquito house entry in the Lao PDR. *PloS one*. 2013;8:e62769.
- Lwetoijera DW, Kiware SS, Mageni ZD, Dongus S, Harris C, Devine GJ, Majambere S. A need for better housing to further reduce indoor malaria transmission in areas with high bed net coverage. *Parasit Vectors*. 2013;6:57.
- 90. Bhatt S, Weiss DJ, Cameron E, Bisanzio D, Mappin B, Dalrymple U, Battle K, Moyes CL, Henry A, Eckhoff PA, Wenger EA, Briët O, Penny MA, Smith TA, Bennett A, Yukich J, Eisele TP, Griffin JT, Fergus CA, Lynch M, Lindgren F, Cohen JM, Murray CLJ, Smith DL, Hay SI, Cibulskis RE, Gething PW. The effect of malaria control on Plasmodium falciparum in Africa between 2000 and 2015. *Nature.* 2015;526:207-211.
- Nyunt MH, Shein T, Zaw NN, Han SS, Muh F, Lee SK, Han JH, Thant KZ, Han ET, Kyaw MP. Molecular Evidence of Drug Resistance in Asymptomatic Malaria Infections, Myanmar, 2015. *Emerg Infect Dis.* 2017;23:517-520.