Made-to-measure malaria vector control strategies: rational design based on insecticide properties and coverage of blood resources for mosquitoes

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Abstract
Eliminating malaria from highly endemic settings will require unprecedented levels of vector control. To suppress mosquito populations, vector control products targeting their blood hosts must attain high biological coverage of all available sources, rather than merely high demographic coverage of a targeted resource subset, such as humans while asleep indoors. Beyond defining biological coverage in a measurable way, the proportion of blood meals obtained from humans and the proportion of bites upon unprotected humans occurring indoors also suggest optimal target product profiles for delivering insecticides to humans or livestock. For vectors that feed only occasionally upon humans, preferred animal hosts may be optimal targets for mosquito-toxic insecticides, and vapour-phase insecticides optimized to maximize repellency, rather than toxicity, may be ideal for directly protecting people against indoor and outdoor exposure. However, for vectors that primarily feed upon people, repellent vapour-phase insecticides may be inferior to toxic ones and may undermine the impact of contact insecticides applied to human sleeping spaces, houses or clothing if combined in the same time and place. These concepts are also applicable to other mosquito-borne anthroponoses so that diverse target species could be simultaneously controlled with integrated vector management programmes. Measurements of these two crucial mosquito behavioural parameters should now be integrated into programmatically funded, longitudinal, national-scale entomological monitoring systems to inform selection of available technologies and investment in developing new ones.

Keywords: Plasmodium, Anopheles, Vector control, Mosquito, Malaria, Target product profile

Background
While anti-parasitic drugs and vaccines will be essential for the final stages of malaria elimination, their effectiveness as transmission control interventions will rely heavily upon first achieving unprecedented levels of vector control in settings with historically high levels of endemicity [1–4]. The most important malaria parasites of humans are entirely dependent on people as their only secondary, mammalian hosts, so the most potent vector mosquito species are those with highly specialized behaviour adapted to feeding upon humans indoors at night when they are asleep [5–9]. Thus, the majority of the most potent vectors distributed across the tropics predominantly feed upon humans inside houses, where they can be effectively controlled and even eliminated with long-lasting insecticidal nets (LLINs) or indoor residual spraying (IRS) [6–9]. While LLINs and IRS can reduce transmission by these human-specialized, indoor-feeding mosquito species by as much as two orders of magnitude, there are many parts of Africa and the Pacific where malaria transmission can occur at levels four orders of magnitude greater than that required to sustain the parasite population [4]. Much of the residual transmission that persists following scale up of LLINs and/or IRS is sustained by mosquitoes that can evade contact with these
insecticidal interventions by feeding upon humans and animals outdoors [4,6-9]. It will therefore not be possible to eliminate malaria transmission from most of the tropics without developing additional scalable vector control strategies which complement LLINs and IRS by extending intervention coverage of the blood resources that mosquitoes depend upon beyond humans and their houses [4,6-10].

To achieve this laudable goal in practice, product developers, manufacturers and end-users need a manageable short list of ecologically-defined target product profiles to work towards that are based on field-measured behavioural and physiological traits of wild vector populations [11]. From the resulting arsenal of complementary vector control products, malaria control programmes will need to select the most effective subset of these options, based on national or regional surveys of these same key behavioural and physiological traits [8,12]. While quite a long list of underlying parameters of mosquito, parasite and human populations determine the overall level of malaria transmission that occurs in a locality, many of these are difficult or impossible to measure routinely across nationally representative scales and relatively few of them are direct targets of vector control measures [11]. Here, a simple conceptual framework based on mathematical models is described that allows new and existing tools for controlling adult malaria vectors to be prioritized and optimized for specific contexts, by predicting their relative merits based on only two field-measurable behavioural parameters of local mosquito populations and two field-measurable indicators of how those mosquitoes interact with specific vector control products.

**Biological coverage of all blood resources available to mosquitoes**

Suppression of mosquito populations with vector control products depends on high biological coverage [13], broadly defined as the proportion of all available sources of blood that is effectively modified to kill, deter, contaminate, or incapacitate mosquitoes at times and places when they attempt to use it. The crucial difference between conventional demographic coverage of humans with a protective measure, and biological coverage of blood resources that mosquitoes depend upon, is that the latter is inclusive of all forms of that resource, while the former is merely the subset of that resource that humans represent at the times and places when they can use the intervention. While this definition can be expanded and applied to any resource mosquitoes may exploit (Killeen GF, Seyoum A, Gimnig JE, Corliss G, Kiware SS, Stevenson JC, Drakeley CJ, Chitnis N, personal communication), host attack and blood acquisition are the best understood of all resource utilization behaviours and can be conveniently, passively surveyed by attracting vectors to hosts sampled from within quantifiable populations of humans or animals [10]. These behaviours are also the most obvious and common target for vector control interventions, because they are obligate behaviours for all Anophelinae and determine the rate of pathogen transmission [10]. Biological coverage of all available blood resources with a protective measure \( C_{A,p} \) can therefore be estimated as the product of demographic coverage, defined and surveyed as the proportion of humans protected on a given night \( (C_p) \), and two field-measurable mosquito behavioural parameters: the human blood index \( (Q_h) \) and the proportion of human exposure that occurs indoors \( (\pi_{h,i}) \) [13]:

\[
C_{A,p} = \pi_{h,i} Q_h C_p
\]

Beyond defining coverage of vector control interventions in a measurable and biologically meaningful way, field measurements of these two behavioural parameters can also guide the specification of ideal target product profiles for delivering insecticides to humans or livestock [9,13-16].

**Blood source as a determinant of intervention selection and impact**

Human blood indices are difficult to measure where vector populations are sparse or primarily rest outdoors, and are inevitably prone to bias arising from heterogeneities of sampling efficiency by resting site category [17,18]. Nevertheless, such estimates are remarkably useful as predictors of large-scale variations of pre-existing malaria transmission intensity [19] and are equally important for selecting optimal vector control methods (Figure 1). The vast majority of human malaria infections are caused by *Plasmodium falciparum* and *Plasmodium vivax*, which are both strict anthroposes, so the most efficient vectors in the world are those which predominantly feed upon humans [19]. Fortunately, this dependence upon human blood also renders them vulnerable to population control [7,8] and even elimination [9] with high coverage of people with insecticidal personal protection measures such as LLINs and IRS. The more a vector depends upon human blood, the greater will be the impact of human personal protection measures upon their population density, longevity and transmission potential, and the greater will be the advantage of pesticides which kill rather than repel mosquitoes (Figure 1A) [13,16,20]. For highly efficient, anthropophagic and endophagic vectors that are most readily controlled with indoor use of contact toxins, it is predicted that outdoor repellent use confers no advantage (Figure 1B) and indoor repellent use dramatically underlines the otherwise massive impact of LLIN use (Figure 1C) [15,20].

However, approximately 40% of all *P. falciparum* infections [21] and 95% of *P. vivax* infections [22] occur
outside of sub-Saharan Africa, where diverse primary vectors [23] predominantly feed on animals rather than humans [19]. Where human blood is unimportant to vector survival and reproduction, personal protection of people will have negligible impact upon the mean density, longevity or stability of those mosquito populations but may achieve community-level protection of non-users by simply blocking vector contact with infectious users and vice versa [13,16]. It is therefore irrelevant whether that is achieved through toxicity or repellency (Figure 1B) so personal protection against highly zoophagic vectors should be maximized by whichever mode of action is most practical [16]. Zoophagic vectors usually prefer to feed outdoors where vapour-phase insecticides should have significant advantages because enclosing structures to provide physical protection and application surfaces for solid-phase residual toxins are typically absent, impractical or even undesirable [15]. By definition, any repellent action of an insecticide is manifested at lower, sub-lethal doses of insecticides than those required to kill mosquitoes [24,25] so the former non-lethal mode of action should be optimized to maximize the personal protection afforded by a vapour-phase active ingredient against vectors that primarily feed upon animals.

Toxic insecticides may therefore have substantive advantages over repellents for targeting humans where people represent an important blood source to mosquitoes but not where they primarily rely upon animal blood. This potential advantage of contact toxins over repellents, and its dependence upon the host preferences of the vector, is illustrated in terms of human feeds per mosquito lifetime in Figure 2. For a mosquito such as Anopheles culicifacies, which rarely feeds on human blood but does so often enough to act as a primary vector [16,26], a repellent should achieve community-level suppression of malaria transmission that is equivalent to that of a toxic product conferring the same level of personal protection. This is because feeding upon humans is a relatively rare event, so most transmission is mediated by mosquitoes taking the bare minimum of two human blood meals required to complete the transmission cycle. Mosquitoes that survive after being repelled from a human or human household have a very low chance of ever feeding on another human. High coverage of efficacious repellents can therefore break the transmission cycle by making the possibility of a mosquito feeding on humans twice even more remote, so the epidemiological impact of this mode of action is equivalent to killing mosquitoes outright with toxicants (Figure 2A). However, for a mosquito with a strong or even moderate preference for human blood, such as Anopheles gambiae and Anopheles arabiensis, respectively, diversion away from a protected human user and extension of host-seeking activity undoubtedly increases associated mortality risks, but many will survive and feed on other humans nearby so the toxic product always has a considerable advantage (Figure 2B and Figure 2C) that is very much needed when faced with the massive transmission levels they mediate [2,4,16].

Beyond directly protecting their occasional human victims, mosquito-toxic insecticides may also be applied to...
livestock, to enable population control of zoophilic vectors and achieve greater proportional reductions of transmission where these are the preferred hosts for dominant local vectors [29-31], and transmission is fundamentally easier to manage because zoophilic mosquitoes are less efficient vectors of anthropogenic Plasmodium [16-19]. However, many important vector species in residual transmission systems, such as An. arabiensis in Africa, Anopheles darlingi in Latin America and Anopheles farauti in the Pacific, can readily feed upon either humans or animals [6-8], so that they represent quite efficient vectors requiring a combination of complementary measures to achieve effective intervention coverage of all preferred host types (Figure 3). Furthermore, they express both zoophagy and anthropophagy with remarkable phenotypic plasticity, resulting in spectacular variation in human blood indices across very fine scales [17,18,27,32], so these vectors alone may exhibit behavioural properties encompassing large tracts of the parameter space and associated intervention needs represented by Figure 3.

Current strategic frameworks for developing new vector control tools emphasize the importance of overall human biting rates, expressed using classical Macdonald-Ross models as the product of mosquito population density per human \( (m) \) and the square of the human biting frequency per mosquito \( (a) \) [11]. While both parameters are of central importance to baseline levels of transmission, and therefore to the levels of control that will be required to eliminate it, the human-feeding frequency has far greater influence on local transmission intensity, and therefore geographic distribution of malaria risk [19], because each transmission event requires two blood meals upon humans so vectorial capacity is approximately proportional to its square \( (a^2) \) [33]. The human biting frequency is also proportional to the human blood index \( (Q_h) \) where \( f \) is the mean duration of the feeding cycle length of individual mosquitoes) and is therefore far more relevant to intervention prioritization and optimization (Figures 1, 2 and 3). For example, it is difficult to envisage a situation in which LLINs or IRS would be de-prioritized as the first-choice options for tackling anthropophagic An. gambiae or Anopheles funestus, regardless of their population density.

While human biting frequency \( (a) \), human blood index \( (Q_h) \) or an equivalent term feature in essentially all process-explicit models of malaria transmission [33,34], and are of central importance to selecting and optimizing the most appropriate vector control strategy (Figures 1, 2 and 3), the classical modelling studies that underpinned...
planning of the GMEP [35,36] not only ignored the way in which frequent feeding upon animals ($a$ or $Q_h \rightarrow 0$) attenuates the population suppression effects of human-targeted vector control measures [13,16], these models also omitted parameters to account for the fact that mosquitoes feeding upon humans may not necessarily do so where and when they can be targeted with LLINs or IRS [2,6-8,10,12,13,15,37].

Figure 3 A conceptual illustration of how optimal vector control interventions and intervention combination could be mapped across vector behaviour parameter space, populated by field measurements of diverse target vectors.

Maximizing protective coverage of humans and houses

Feeding upon alternative hosts is by no means the only form of behavioural resilience or resistance [4,8] that limits biological coverage of personal protection measures such as LLINs: some mosquitoes attack humans at times and places where these measures are not realistically applicable. For any product conferring personal protection against mosquito bites, estimating the proportion of human exposure to mosquito bites that occurs at times when it is practical to use it ($n_{h,i}$) is critically important to measuring the maximum amount of protection that can be realistically expected [13,38,39]. In the case of LLINs, this definition can be approximated as the proportion of normal exposure to mosquito bites upon humans lacking LLINs which occurs indoors ($n_{h,i}$) or during sleeping hours ($n_{h,s}$) when it would be practical to use a net [38,39]. These parameters are measured in the field by weighting the observed indoor and outdoor biting rates at each period of the night by the surveyed mean proportion of humans that are indoors and outdoors, respectively, at that time [5,38,39]. While these parameters can be measured for individual people, or strata within human populations [5,40], it is their community-wide mean values as experienced by the mosquito population that determines the magnitude of the mass effect of vector control interventions [2,16,37]. In Africa, consistently high values for this key behavioural parameter, even in some settings with long-established high coverage of LLINs, are primarily driven by the preference of An. gambiae, An. arabiensis and An. funestus for feeding at times when most people are indoors asleep, rather than any strong or consistent preference for feeding indoors per se [38]. These estimates of the proportion of human exposure occurring indoors for net users ($n_{h,i,n}$) suggest at least half of residual transmission now occurs outdoors in African settings with high LLIN coverage [4]. Furthermore, the proportion of human exposure to residual vector populations that occurs indoors has recently dropped in some settings with high coverage of LLINs or IRS [6-8,12]. It has been suggested that these altered patterns of mosquito activity can be explained by the persistence of hungry mosquitoes until people are unprotected at dusk and dawn, a form of behavioural phenotypic plasticity that can be classified as behavioural resilience rather than resistance [7,8]. However, recent modelling analysis suggests shifting distributions of the times when wild mosquito populations actually feed successfully upon human communities using LLINs should not be manifested in the biting rates experienced by unprotected human volunteers because they are fully exposed [42]. Changes in biting patterns observed by
human landing catch may therefore represent genuine emergence of behavioural resistance in the form of altered innate feeding time preferences [6,42].

Regardless of whether observations of outdoor-feeding behaviour reflect pre-existing resilience or emerging resistance, it is clear that they will have to be addressed with insecticide-treated clothing [43,44], vapour-phase insecticides that protect humans outside of their houses [24], or some other intervention that prevents bites from outdoor-feeding mosquitoes [42,45]. While repellents may be ideal for protecting against outdoor exposure to zoophagic vectors [13,15,16], some outdoor-feeding Asian species such as Anopheles dirus [46] and An. farauti [47,48] often feed predominantly upon humans [7] so vapour-phase insecticides that lack repellent properties may be preferred to maximize toxic exposure, mortality rates and population suppression of these species (Figure 3). Furthermore, the persistence and even predominance of indoor-feeding behaviours in vector populations exposed to high coverage of LLINs and/or IRS [4,5,38,48,49] suggests there is still considerable room for improvement upon these technologies for killing mosquitoes that enter houses [42,50]. The proportions of human exposure which occur indoors and outdoors are therefore important and dynamic indicators of vector behaviour that malaria control programmes should survey on a routine basis [6,8,12,13,38] so that they can manage malaria transmission in the same integrated, evidence-based, locally tailored and adaptive manner as agricultural pests [51].

While these indicators are ideal for LLINs, field measurements of the maximum proportion of human exposure which is directly preventable (\(n_h\)) by other personal protection interventions will require more careful consideration, especially for insecticidal clothing or repellent products with usage patterns that are more difficult to survey because they are portable, used outdoors, or require frequent re-application. The issue of where and when protective measures should be applied becomes particularly important for repellents in settings where zoophagic vectors co-exist with anthropophagic counterparts that have already been suppressed with LLINs or IRS applied indoors, or with insecticide-treated clothing or non-repellent, mosquito-toxic, vapour-phase insecticide emanators applied outdoors. Suppressed populations of such potent vectors, that are otherwise behaviourally vulnerable to control, may well rebound if the toxic action of these products is undermined when they are supplemented with repellents [14,15,20]. It is therefore important to note that while both repellent and toxic products may be required in many scenarios where one or more vectors exhibit intermediate or wide-ranging values of the human blood index (Figure 3), these should not be applied in the same time and place but rather combined in a complementary manner, ideally to achieve a “push-pull” strategy similar to those applied to agricultural pests [52].

**Biological coverage of indoor resting sites versus human blood indoors**

If one considers interventions which target resources other than blood, that mosquitoes may use several times in a single gonotrophic or feeding cycle, it is clear that existing definitions for \(Q_h\) and \(n_h\), based on the concept of protecting humans against exploitation by mosquitoes as sources of blood, must be extended and generalized further. Taking IRS of resting sites in human habitations as the most obvious example, models of malaria transmission and vector population dynamics could be parameterized using estimates of the utilization rate of indoor resting sites, quantified as the mean number of times that a typical mosquito rests indoors per gonotrophic cycle (\(\alpha_{r,i}\), where \(\alpha\) represents the mean number of times a mosquito utilizes any given resource during a single gonotrophic cycle, \(r\) represents all resting site resources, and \(i\) represents the subset of resting sites that are indoors (Killeen GF, Seyoum A, Gimnig JE, Corliss G, Kiware SS, Stevenson JC, Drakeley CJ, Chitnis N, personal communication)). However, despite the widespread use and global prioritization of IRS as a frontline malaria vector control tool [53], the only available field measurements of this parameter are undoubtedly underestimated because they rely upon captures of resting mosquitoes at a single point in time in the early morning. Such a temporal snapshot of resting events will obviously fail to detect mosquitoes that rested on the surveyed indoor surfaces but then left again before they were surveyed. Entomological survey methods for dramatically improving the detection efficiency of resting events clearly need to be developed, presumably by exploiting the diversity of marker systems that are now available for labelling insects [54], or the rapidly improving technologies for observing them visually [55,56].

In the absence of direct measurements of \(\alpha_{r,i}\), it is possible to use \(n_{h,i}\) as a reasonable surrogate in many contexts, based on the assumption that many vectors which preferentially feed inside houses usually rest there too. Defining vector control coverage in terms of mosquito dependence upon obtaining blood from humans indoors has therefore proven useful for rationalizing the differential impact of not only LLINs, but also IRS, upon sympatric primary vectors in a variety of settings [9,13]. However, using \(n_{h,i}\) as a surrogate for \(\alpha_{r,i}\) does have major limitations and may be very misleading for many vectors with divergent values of these two parameters because they feed indoors but rest outdoors or *vice versa*. Vector species that combine indoor feeding with natural or insecticide-induced outdoor resting are important contributors to persistent residual malaria transmission, despite high coverage of LLINs or IRS.
sites (fully exploited with vector control using rationally-designed mosquitoes remains to be quantitatively understood, or remarkable that utilization of indoor resting spaces by rates at which mosquitoes utilize subsets (doors, lack of adequate procedures for measuring the indoors but rest outdoors, or feed outdoors but rest in- given that several important vector species either feed a similar manner to the way in which Figures 1 and 3 to map out predicted and observed impacts of IRS, as well as other intervention strategies targeting specific subsets of resting sites, across behavioural parameter space in a similar manner to the way in which Figures 1 and 3 do so for blood resources, which are more readily defined and quantified. However, in practice, the measurement of these parameters is more challenging. Given that several important vector species either feed indoors but rest outdoors, or feed outdoors but rest indoors, lack of adequate procedures for measuring the rates at which mosquitoes utilize subsets (x) of resting sites (r) that can be targeted with insecticides (αx,r), the most important of which are the indoors surfaces (x = i) inside houses and other shelters (αx,i), is clearly a methodological deficit that needs to be urgently addressed.

Conclusions
For mosquito populations to be successfully suppressed, vector control products targeting their blood hosts need to attain high biological coverage of all available sources, rather than merely high demographic coverage of a targeted subset such as humans indoors [9,13]. Beyond defining biological coverage in a quantifiable manner, the human blood index and the proportion of human exposure that occurs indoors may also be used to define optimal target profiles for diverse products and product combinations to protect humans or livestock against blood-seeking mosquitoes (Figures 1 and 2). For vector mosquitoes that feed only occasionally upon humans, preferred animal hosts may be optimal targets for mosquito-toxic insecticides, and vapour-phase insecticides optimized to maximize repellency, rather than toxicity, may be ideal for directly protecting people against indoor and outdoor exposure. However, for vectors that primarily feed upon people, repellent vapour-phase insecticides may be inferior to toxic ones and may undermine the impact of contact insecticides applied to human sleeping spaces, houses or clothing if combined in the same time and place. The theory of biological coverage may also be extended to other life history parameters, such as indoor resting, to assess the impact of vector control interventions, such as IRS, which target resting mosquitoes. While measurements of the proportion of human blood meals that occur indoors often correlate well with the rates at which vectors utilize indoor resting sites, many important vectors do not rest where they feed. In such cases, the proportion of human blood meals occurring indoors cannot be used a proxy measure of indoor resting site utilization rate, and aspiration capture of resting mosquitoes only gives a snapshot of mosquito distributions at specific points in time, so new entomological methods for detecting all resting events at insecticide-targetable surfaces are urgently needed. The conceptual framework outlined here relates to anthroponotic malaria parasites as specific motivating examples. However, these concepts and strategies should also be directly applicable to other mosquito-borne anthropoposes, such as dengue, urban yellow fever and lymphatic filariasis, or adapted to zoonotic pathogens such as *Plasmodium knowlesi*, Rift Valley fever and West Nile virus, for which their implications should be different but no less rational, so that diverse target species can be simultaneously controlled with integrated vector management programmes [62]. However, rather than stereotyped, hypothetical schematics such as Figure 3, control programmes, policy makers, research funders, and product developers need such maps of vector behaviour parameter space to be populated with real field estimates of these mosquito behaviours and impacts of specific interventions upon those vectors. Only then will they be able to inform the selection of available technologies based on rational expectations of impact, and also prioritize investment in developing new ones. Entomological techniques for measuring these two critical behavioural parameters are well established and have changed little since classic texts were written during, or immediately after, the failed Global Malaria Eradication Programme decades ago [5,17,18,63,64]. Unfortunately, they have only been applied at village or district scales thus far, with inconsistent methodology, and with haphazard distribution across times and locations, because they have been predominantly funded
through sporadic, short-term research projects. Fortunately, recent global policy emphasizes strengthening capacity for routine entomological surveillance at Ministries of Health in particular, rather than just research and academic institutions [65], so perhaps the time has finally come to integrate such behavioural parameter measurements into programmatically funded, longitudinal monitoring systems operating on national and regional scales [8].

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References

1. Wenger EA, Eckhoff PA: A mathematical model of the impact of present and future malaria vaccines. Malar J 2013, 12:126.
2. Eckhoff PA: Mathematical models of within-host and transmission dynamics to determine effects of malaria interventions in a variety of transmission settings. Am J Trop Med Hyg 2013, 88:817–827.
3. Okell LC, Griffin JT, Kleinschmidt I, Hollingsworth TD, Churcher TS, White MT, Bousema T, Drakeley CJ, Ghani AC: The potential contribution of mass treatment to the control of Plasmodium falciparum malaria. PLoS One 2011, 6:e20179.
4. Killeen GF: A second chance to tackle African malaria vector mosquitoes that avoid houses and don’t take drugs. Am J Trop Med Hyg 2013, 88:809–816.
5. Elliott R: The influence of vector behaviour upon malaria transmission. Am J Trop Med Hyg 1972, 21:755–763.
6. Russell TL, Beeve NW, Cooper RD, Lobo NF, Burkot TR: Successful malaria elimination strategies require interventions that target changing vector behaviours. Malar J 2013, 12:56.
7. Durnez L, Coosemans M: Residual transmission of malaria: an old issue for new approaches. In Anopheles mosquitoes – New insights into malaria vectors. Edited by Manguin S. Rijeka: Intech; 2013:671–704.
8. Govella NJ, Chaki PP, Killeen GF: Entomological surveillance of behavioural resilience and resistance in residual malaria vector populations. Malar J 2013, 12:24.
9. Killeen GF, Seyoum A, Sikaila CH, Zomboko AS, Gimnig JE, Govella NJ, White MT: Eliminating malaria vectors. Parasitol Vectors 2013, 6:172.
10. Ferguson HM, Domhau A, Beeche A, Borgemeister C, Gottlieb M, Mulla MS, Gimnig JE, Fish D, Killeen GF: Ecology: a prerequisite for malaria elimination and eradication. PLoS Med 2010, 7:e1000303.
11. Vontas J, Moore SJ, Kleinschmidt I, Ranson H, Lindsay SW, Lengeler C, Hamon N, McLean T, Hemingway J: Framework for rapid assessment and adoption of new vector control tools. Trends Parasitol 2014, 30:191–204.
12. Gatton ML, Chitnis N, Chuchter T, Donnelly MJ, Ghani AC, Godfray HC, Gould F, Hastings I, Marshall J, Ranson H, Rowland M, Shaman J, Lindsay SW: The importance of mosquito behavioural adaptations to malaria control in Africa. Evolution 2013, 67:1218–1230.
13. Kiware SS, Chitnis N, Devine GJ, Majambere S, Killeen GF: Biologically meaningful coverage indicators for eliminating malaria transmission. Biol Lett 2012, 8:874–877.
14. Killeen GF, Chitnis N, Moore SJ, Okumu FO: Target product profile choices for intra-domiciliary malaria vector control pesticide products: repel or kill? Malar J 2011, 10:207.
15. Killeen GF, Moore SJ: Target product profiles for protecting against outdoor malaria transmission. Malar J 2012, 11:17.
16. Kiware SS, Chitnis N, Moore SJ, Devine GJ, Majambere S, Killeen GF: Simplified models of vector control impact upon malaria transmission by zoophilic mosquitoes. PLoS One 2012, 7:e37661.
17. Garrett-Jones C: The human blood index of malaria vectors in relation to epidemiological assessment. Bull World Health Organ 1984, 30:241–261.
18. Garrett-Jones C, Boreham P, Pant CP: Feeding habits of anopheles. (Diptera: Culicidae) in 1971–1978, with reference to the human blood index: a review. Bull Entomol Res 1980, 71:165–185.
19. Kowszewski AE, Mellingar A, Spilman A, Malaney P, Sachs SE, Sachs J: A global index representing the stability of malaria transmission. Am J Trop Med Hyg 2004, 70:486–498.
20. Yakob L, Dunning R, Yan G: Indoor residual spray and insecticide-treated bednets for malaria control: theoretical synergisms and antagonisms. J R Soc Interface 2011, 8:799–806.
21. Hay SI, Okto EA, Getting PW, Patil AP, Tamen AJ, Guerra CA, Snow RW: Estimating the global clinical burden of Plasmodium falciparum malaria in 2007. PLoS Med 2010, 7:e1000290.
22. Guerra CA, Howes RE, Patil AP, Getting PW, Van Boeckel TP, Temperley WH, Kabaria CW, Tatem AJ, Marsh H, Elzanjar R, Band A, Snow RW, Hay SI: The international limits and population at risk of Plasmodium vivax transmission in 2009. PLoS Negl Trop Dis 2010, 4:e774.
23. Sinka ME, Bangs MJ, Manguin S, Rubio-Faliso Y, Chareonviriyaphap T, Coetzee M, Mbogo CM, Hemingway J, Patil AP, Tempefre WH, Getting PW, Kabaria CW, Burkot TR, Harbach RE, Hay SI: A global map of dominant malaria vectors. Parasit Vectors 2012, 5:69.
24. Achene NL, Bangs MJ, Farlow R, Killeen GF, Lindsay S, Logan JG, Moore SJ, Rowland M, Sweeney K, Tort SJ, Zwiwelbi L, Grieco JP: Spatial repellents: from discovery and development to evidence-based validation. Malar J 2012, 11:64.
25. Grieco JP, Achene NL, Chareonviriyaphap T, Suwonkerd W, Chauhan K, Sardelis MR, Roberts DR: A new classification system for the actions of IRS chemicals traditionally used for malaria control. PLoS One 2007, 2:e716.
26. Dav V, Sharma VP: The dominant mosquito vectors of human malaria in India. In Anopheles mosquitoes-New insights into new malaria vectors. Edited by Manguin S. Rijeka, Croatia: Intech; 2013:239–271.
27. Killeen GF, McKenzie FE, Foy BD, Bogh C, Beier JC: The availability of potential hosts as a determinant of feeding behaviours and malaria transmission by mosquito populations. Trans R Soc Trop Med Hyg 2001, 95:469–476.
28. Rawlings P, Curtis CF: Tests for the existence of genetic variability in the tendency of Anopheles culicifacies species B to rest in houses and to bite man. Bull World Health Organ 1982, 60:427–432.
29. Rowland M, Dunnari N, Kerrward M, Mohammed N, Urahman H, Hewitt S: Control of malaria in Pakistan by applying deltamethrin insecticide to cattle: a community-randomised trial. Lancet 2001, 357:1837–1841.
Mahende AM, Mosha FW, Mahende JM, Kwesa E: Role of cattle treated with deltamethrin in areas with a high population of Anopheles arabiensis in Moshi, Northern Tanzania. *Malar J* 2007, 6:109.

Lyimo IN, Ng'habi KR, Mpanga MW, Daraja AA, Mwashehe DD, Nichimbi NS, Lwetoijire DW, Mvunya LI: Does cattle milling provide a potential target to control wild exophilic Anopheles arabiensis (Diptera: Culicidae) with entomopathogenic fungi? A bio-entomopathogenic zoophrophylaxis strategy for vector control. *J Parasitol Res* 2012, 2012:280583.

Kay BH, Boreham PFL, Edman JD: Application of the "Feeding Index" concept to studies of mosquito host-seeking patterns. *Mosquito News* 1979, 39:88–73.

Smith DL, Battle KE, Ayala JM, Scott TW, McKenzie FE: A new tool for malaria prevention? Results of a trial of permethrin-impregnated mosquito bed nets: a modelling study. *PLoS Pathog* 2012, 8:e1002588.

Reiner RCJ, Perkins TA, Barker CM, Niui T, Chaves LF, Ellis AM, George DB, Le Menach A, Pulliam JRC, Baisancho D, Buckee C, Chiyaka C, Cummings DAT, Garcia AJ, Gatton MG, Geithing PW, Hartley DM, Johnston G, Klein EY, Michael E, Lindsay SW, Lloyd AL, Pigott DM, Reisen WK, Ruktanchanai N, Singh BK, Tatem AJ, Kinton U, Hay SI, Scott TW, et al: A systematic review of mathematical models of mosquito-borne pathogen transmission: 1970–2010. *J R Soc Interface* 2013, 10:20120921.

Garrett-Jones C: Prognosis for interruption of malaria transmission through assessment of the mosquito’s vectorial capacity. *Nature* 1964, 204:1173–1175.

Macdonald G: *The epidemiology and control of malaria*. London: Oxford University Press; 1957.

Griffin JT, Hollingsworth TD, Okell LC, Churcher TS, White M, Hinsley W, Macdonald G: Methods for marking insects: current techniques and future prospects. *Annu Rev Entomol* 2014, 59:529–553.

Boutil S, Manoukis N, Diallo M, Ribeiro JM, Paley DA: Reconstructing the flight kinematics of swarming and mating in wild mosquitoes. *J R Soc Interface* 2012, 9:2624–2638.

Spitzen J, Spoor CW, Gi Erica F, Ter Braak C, Beuwekes J, Van Bruggen SP, Kranenburg S, Noldus LP, Van Leeuwen JL, Takken W: A 3D analysis of flight behavior of Anopheles gambiense s.s. and other malaria mosquitoes in response to human odor and heat. *PLoS One* 2013, 8:e62995.

Muirhead-Thomson RC: MOSQUITO BEHAVIOR IN RELATION TO MALARIA TRANSMISSION AND CONTROL IN THE TROPICS. London: Edward Arnold & Co.; 1951.

Kitau J, Osbrink RM, Tunu PNG, Matowa J, Malima RC, Magesa SM, Bruce J, Mafura F, Rowland MW: Species shifts in the Anopheles gambiae complex: Do LLINs successfully control Anopheles arabiensis? *Malar J* 2012, 11:3481.

Okumu FO, Kiware SS, Moore SJ, Killeen GF: Mathematical evaluation of community level impact of combining bed nets and indoor residual spraying upon malaria transmission in areas where the main vectors are Anopheles arabiensis mosquitoes. *Parasit Vectors* 2013, 6:177.

Okumu FO, Mbeje EA, Ligamba G, Moore J, Ntamurugiro AJ, Kavishe DR, Kinyanga MG, Turner E, Lerenza LM, Moore SJ: Comparative evaluation of combinations of long lasting insecticidal nets and indoor residual spraying, relative to either method alone, for malaria vector control in an area dominated by Anopheles arabiensis. *Parasit Vectors* 2013, 6:46.

Pluess B, Tanser FC, Lengeler C, Sharp BM: Indoor residual spraying for preventing malaria. *Cochrane Database Syst Rev* 2010, 4:CD006657.

WHO: *Global Strategic Framework for Integrated Vector Management*. Geneva: World Health Organization; 2004.

Elliot R: Studies on man-vector contact in some malariasis areas in Colombia. *Geneva: World Health Organization; 1967:32.*

Garrett-Jones C: A method for estimating the man-biting rate. *Geneva: World Health Organization; 1964:22.*

WHO: *Guidance note on capacity building in malaria entomology and vector control*. Geneva: Geneva Global Malaria Programme, World Health Organization; 2013:3.