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THE POTENTIAL IMPACT OF INTEGRATED MALARIA TRANSMISSION CONTROL ON ENTOMOLOGIC INOCULATION RATE IN HIGHLY ENDEMIC AREAS

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Abstract

We have used a relatively simple but accurate model for predicting the impact of integrated transmission control on the malaria entomologic inoculation rate (EIR) at four endemic sites from across sub-Saharan Africa and the southwest Pacific. The simulated campaign incorporated modestly effective vaccine coverage, bed net use, and larval control. The results indicate that such campaigns would reduce EIRs at all four sites by 30- to 50-fold. Even without the vaccine, 15- to 25-fold reductions of EIR were predicted, implying that integrated control with a few modestly effective tools can meaningfully reduce malaria transmission in a range of endemic settings. The model accurately predicts the effects of bed nets and indoor spraying and demonstrates that they are the most effective tools available for reducing EIR. However, the impact of domestic adult vector control is amplified by measures for reducing the rate of emergence of vectors or the level of infectiousness of the human reservoir. We conclude that available tools, including currently neglected methods for larval control, can reduce malaria transmission intensity enough to alleviate mortality. Integrated control programs should be implemented to the fullest extent possible, even in areas of intense transmission, using simple models as decision-making tools. However, we also conclude that to eliminate malaria in many areas of intense transmission is beyond the scope of methods which developing nations can currently afford. New, cost-effective, practical tools are needed if malaria is ever to be eliminated from highly endemic areas.

INTRODUCTION

Malaria transmission intensity is best expressed as the entomologic inoculation rate (EIR) which directly reflects the exposure of humans to pathogenic *Plasmodium* parasites.^{1–3} In malaria-endemic parts of sub-Saharan Africa and the southwest Pacific, transmission intensity can vary from undetectable levels to more than 1,000 infective bites per year.^{1,2} Entomologic inoculation rate levels of one infective bite per year, or less, can readily sustain prevalence in

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excess of 40% for *Plasmodium falciparum*, the most pathogenic of the species of human malaria parasites.² Although the clinical outcome of malaria infection depends on the specific immunological and pathological interactions of parasite strains and individual humans,^{4–9} most, if not all, strains of *P. falciparum* are potentially life-threatening.^{10,11} Stable, endemic malaria, at any level of transmission, is inevitably associated with a heavy disease burden.^{2, 12–14} Clearly, very large and sustainable reductions of transmission intensity would be required to negate the public health impact of malaria in endemic areas of sub-Saharan Africa and the southwest Pacific. Nevertheless, despite considerable recent debate,^{15–18} the balance of existing evidence strongly supports the view that effective malaria transmission-control interventions^{19,20} will always be beneficial with respect to all-cause mortality.^{2,12,14,20} Malaria transmission control can reduce morbidity and mortality,^{12,14,20} but large reductions in EIR are required in most endemic settings.^{2,12,14,20}

Transmission control and clinical treatment are separate components of any malaria control program and should be treated as such.^{2,11} Rapid and appropriate treatment of clinical cases can drastically reduce the number of deaths resulting from malaria but cannot reduce the incidence of attacks and has a limited impact on morbidity.^{11,21} What is not clear is whether wide-scale chemoprophylaxis reduces or increases the infectiousness of the human reservoir in endemic areas.^{22–25} Even if the infectiousness of the human reservoir can be reduced, it is difficult to justify the use of antimalarial drugs for transmission control. This is because the widespread administration of drugs is not only expensive and difficult to implement, but also promotes resistance in the parasite population and makes life-threatening cases more difficult to treat.²⁶ Clearly we should not compromise the safety net of effective treatment and cure for the sake of partial prevention. Therefore, drugs should not be used as a transmission control tool in endemic areas although they are useful for controlling epidemics in areas of unstable transmission.²⁷ Furthermore, recent modeling studies have suggested that the reversal of these roles may be useful: suppression of transmission intensity may help slow the evolution of drug resistance.²⁸

Entomologic inoculation rate can be estimated as the product of the human reservoir infectiousness (κ), the lifetime transmission potential of individual mosquitoes (L) and the rate at which they emerge from larval breeding sites (E) relative to human population size (E/N_h):

$$EIR = \kappa LE / N_h \quad (1)$$

The structure of this equation directly implies that measures which reduce any of these contributors will amplify each other's effects when combined. These three contributors are also discrete targets for transmission control that are reduced by quite different interventions. The only intervention we can envisage which could usefully reduce κ , and which is likely to be available in the foreseeable future is a malaria vaccine.²⁹ Tools for the reduction of L include indoor spraying, bed nets, and zooprophylaxis^{12,19,20} whereas source reduction and other forms of larval control represent well-established methods for reducing E/N_h .^{30–32}

Here we estimate the potential impact of these measures on EIR and assess their potential contributions to integrated transmission control in highly endemic areas. The impact of generalized control programs on EIR is estimated at four malaria-endemic sites in Nigeria, Tanzania, and Papua New Guinea for which site-specific predictors of EIR have been reported.¹ Overall, we conclude that existing tools can reduce malaria transmission intensity enough to considerably reduce mortality, even in highly endemic areas, and should be applied in a concerted manner as components of integrated transmission control programs. However, the model also predicts that the elimination of malaria in areas of intense transmission is beyond

the scope of methods that are available and affordable in developing nations. New tools are needed if this goal is ever to be achieved.

METHODS

Modeling malaria transmission intensity at four endemic sites

All terminology and abbreviations are consistent with Table 1 of the companion paper, which describes the model.¹ This model has been tested using published data from four *P. falciparum*-endemic sites where values for EIR, the sporozoite rate (S), human biting rate (H_{Bt}), human reservoir infectiousness (κ), human blood-feeding preference (Q), survival rate per feeding cycle (P_f) and sporogonic incubation period (n) have all either been measured directly or calculated where reported values for other parameters allow it.¹ For details of these parameters, see Table 2 of the accompanying paper.¹ The model predicted EIR values for the four sites, which were 1.17 ± 0.37 (range 0.84–1.59) times those measured in the field and all outputs were consistent with field measurements and previous models.¹ The four sites, namely Kankiya and Kaduna in Nigeria, Namawala in Tanzania, and Butelgut in Papua New Guinea are foci of intense *P. falciparum* transmission, morbidity, and mortality.¹ These sites span a ten-fold range of EIR and a wide range of environmental conditions from across the African continent and the southwest Pacific.¹ Briefly, Kankiya³³ and Kaduna^{34,35} are in the dry Savannah region of Nigeria; Namawala is near the flood plain of the Kilombero River in Tanzania^{36–38} and Butelgut is in the forested foothills of Madang Province in Papua New Guinea.^{39–41} For simplicity, our model estimates EIR for the predominant vector species at each site, which are *Anopheles arabiensis* at Kankiya, *Anopheles gambiae* sensu lato, at Kaduna and Namawala, and *Anopheles punctulatus* at Butelgut.¹ In order to calculate the impact of the combined effects of the intervention on the human biting rate (H_{bt}) it is calculated from the proposed reductions in vector emergence rate (E) and mean number of human bites per lifetime (b_h) similar to equation 12 in the original model for calculating EIR:¹

$$H_{bt} = b_h E / N_h \quad (2)$$

Modeling the impact of integrated transmission control on EIR

We propose the following set of feasible transmission control measures and model their combined impact on EIR by modeling their effects on κ (human reservoir infectiousness), Q (human feeding preference of the vector), P_f (survival per feeding cycle by the vector), and E/N_h based on literature estimates of their potential impact. All assumptions and input values for baseline conditions (no transmission control) are as described in Tables 2 and 3 of the accompanying paper.¹ For each parameter that a given control measure could be expected to affect, a reasonable proportional impact was chosen, based on published values and reasonable guesses. These values were chosen and fixed *a priori* so that the results of the model would not be biased to fit predicted values for outcome variables to those reported in the literature. Initially, we predicted the impact of a program based on only insecticide-treated bed nets. Then the effects of a more integrated program incorporating a malaria vaccine and larval control were predicted.

Modeling the impact of insecticide-treated bed nets

Cost-effective tools that reduce Q or P_f or that increase f, the length of the feeding cycle, include indoor spraying^{33,42,43} and bed-net impregnation^{43–47} with insecticides. Bed nets have been reported to reduce vector survival per feeding cycle by between 0% and 40% for the vector species present at our four modeled sites.^{43–47} Bed nets can also lengthen the feeding cycle of anthropophilic mosquitoes, even when they shift substantially to biting non-human animals,⁴⁴ but this shift has never been quantified. Therefore, in the interest of making a conservative

estimate, we incorporate an impact of 15% extra mortality per feeding cycle and no impact on feeding cycle length.

Insecticides can also have the effect of repelling mosquitoes from dwellings^{46,48} and can discourage them from feeding upon humans.^{44,49} For example, at a site where the reported impact of impregnated bed nets on the survivorship of the vectors is very low, and indeed statistically insignificant, a shift to feeding on animals (a 30% drop in the human blood index, a measure of Q) and longer feeding cycles were demonstrated.⁴⁴ Thus it seems that bed nets sometimes act by enhancing or enabling zoophylaxis. A strong relationship between the human blood index of *An. gambiae* and the availability of cattle as alternative hosts has been demonstrated in The Gambia⁵⁰ and at one of our Nigerian sites, Kaduna.⁵¹ In The Gambia, a cattle to human ratio of one or more reduced the human blood index of *An. gambiae* s.l. (mostly *An. gambiae* sensu stricto) to below 50%.⁵⁰ Human blood indices of less than 50% have also been observed for both *An. arabiensis* and *An. gambiae* s.s. in a number of Kenyan communities which keep large numbers of cattle within their village compounds.^{52–55} Thus, although the presence of livestock has been known to increase malaria transmission intensity and clear guidelines are not available for most vectors,⁵⁶ it appears that zoophylaxis against malaria vectors is achievable in both sub-Saharan Africa and the southwest Pacific and may passively contribute to the effectiveness of bed nets. In order to incorporate the effects of livestock as decoys into our simulation of a bed-net program, we will include a moderate reduction to Q of 20%.

Modeling the impact of a malaria vaccine

Previous modeling studies have determined that regardless of the parasite stage targeted, the impact of any malaria vaccine on malaria transmission is equivalent to its level of efficacious coverage.⁵⁷ Thus the impact of malaria vaccines can be envisaged as a direct function of their efficacious coverage, which we set at 50%, a reasonable target for such vaccines and considerably better than that achieved by existing candidates.⁵⁸

Modeling the impact of larval control

Reducing E, the vector emergence rate, represents a formidable task in most parts of Africa and Papua New Guinea. Nevertheless, methods do exist which can substantially reduce the availability of vector breeding sites in many situations.¹⁹ It has even been possible to eradicate, almost exclusively by larval control, the sub-Saharan African vector *An. gambiae* in both Brazil³¹ and Egypt³² after they were accidentally introduced there. Much of the larval habitat in both Butelgut⁴¹ and Kaduna^{34,51} are associated with rivers and streams, representing readily identifiable targets for larval control undertaken once a week or so. Although the topography of Kankiya was not well described, like Kaduna it is part of the dry savanna region of Nigeria. At Namawala, the striking peak of vector emergence observed annually is clearly associated with rice cultivation during the rainy season.³⁶ Although methods do exist for vector control in such circumstances, they require rigorous and intensive implementation for complete success.^{31,32} The ability of all of the predominant vector species at these sites to breed in small, dispersed, temporary pools makes control even more difficult. Nevertheless, the complete eradication of *An. gambiae* s.l. from seemingly ideal habitats in flooding river valleys of northeast Brazil has proven that even this challenging vector is vulnerable to well-organized and determined larval control programs.³¹ Furthermore, a similar program in Egypt was effectively implemented at the peak of World War II, demonstrating that it can be achieved even with limited resources.³² For the purpose of a larval control campaign simulation which is conservative enough to be realistic but ambitious enough to be useful, we incorporate a two-fold (50%) reduction of emergence rates, to account for the level of larval habitat control that is possible with existing, cost-effective methods.¹⁹

RESULTS

The use of bed nets without controlling the infectiousness of the human parasite reservoir or larval habitats was predicted to reduce EIR by 7- to 13-fold at all four sites. Predicted reductions of the sporozoite rate (S) and the human biting rate (H_{Bt}) with bed nets were 3- to 5-fold and 2- to 3-fold, respectively. Correspondingly, the largest predicted impact of the integrated program on any of the three contributors to EIR is on L , the lifetime malaria transmission potential of individual mosquitoes (Figure 1B). However, the impact on L is amplified by two-fold reductions of E/N_h , the vector emergence rate relative to human population size (Figure 1C), and of κ , the infectiousness of the human reservoir (Figure 1A). This results in combined EIR reductions by the integrated program of between 30- and 50-fold (Figure 1D). Note, however, that the impact on H_{Bt} , the human biting rate, is not as striking, being approximately 5-fold (Figure 1E). This generalized control program was also predicted to have remarkably consistent effects on other relevant entomological parameters (Figure 2). The integrated control program reduced the lifetime biting capacity of individual vectors (b_h) by only 2- to 3-fold but decreased their lifetime *infectious* biting capacity (β) by 15- to 27-fold. The integrated program reduced the sporozoite rates (S) and individual lifetime transmission potentials (L) of the vector populations by 6- to 11-fold and 8- to 13-fold, respectively.

DISCUSSION

This model yields outputs that are not only accurate under baseline conditions,¹ but also predict the impact of domestic insecticide-based vector control remarkably well. The predictions of this model illustrate how a small decrease in vector populations can have a dramatic impact on EIR, and help explain the proven efficacy of bed nets and indoor spraying.^{12,14,20} The model demonstrates how integrated control using a few modestly effective tools can meaningfully reduce EIR but cannot eliminate malaria completely from areas of intense transmission.

The predicted impact of bed nets alone is strikingly consistent with observations made during insecticide-based control programs.^{42–47,59} Indeed, a DDT-spraying campaign at Kankiya reduced sporozoite rates and human biting rates by approximately 4- and 3-fold, respectively.³³ These reductions are almost identical to the predictions of the model at this site, despite the *a priori* impact choices on the input parameters P_f and Q . The application of these predicted effects on EIR to the empirically established relationship between EIR and *P. falciparum* prevalence,² allows us to estimate that 20–25% reductions in parasite prevalence can be expected. This compares well with observed impact in the field which varies from approximately 10% to 40% reductions in prevalence.^{59–62} Such large reductions of transmission intensity at the community level are consistent with the observed benefits of such programs, in terms of morbidity and mortality^{12,14,20} and their observed impact on unprotected people over considerable distances.⁶³ Also, the modest reductions of b_h , the total number of human bites a female mosquito will take, and hence the number of ovipositions she will complete during her lifetime (Figure 2) indicate that limited selective pressure will be exerted on the vector. This is consistent with the observation that pyrethroid resistance does not seem to emerge in vector populations exposed to impregnated bed nets in areas where these same insecticides are not used widely in agriculture.⁶⁴

The ability of the model to explain the efficacy of bed nets further emphasizes the value of this model as a predictive tool for planning vector-control research and implementation.¹ However, accurately verifying the impact of insecticide-based interventions on the EIR may be difficult in practice. For example, pyrethrum spray catches are likely to underestimate biting rates in dwellings which have insecticide-treated bed nets or walls, because repelled mosquitoes will exit after feeding.^{46,48} Conversely, standard human landing catches tend to overestimate the

biting rate in communities with bed nets where residents are protected, but mosquito collectors are not.⁴⁶ It is also important to bear in mind that the level of impact on Q , P_f , or f required to usefully reduce EIR is quite small and hence difficult to confirm statistically. This model demonstrates how the effects of bed nets may be explained by quite small reductions of these parameters and how the reduction of malaria burden can be achieved without decimating the vector population. Given that bed nets can exert their effects through any of the three parameters, accurate assessment of how transmission intensity is reduced in any trial probably requires detailed measurements of all of them.

The predictions of the model incorporating larval control and a malaria vaccine give an approximate estimate of the level of impact which can be achieved with affordable tools which are available now or likely to be available soon. Only one of the interventions included in the simulated transmission control program, an effective malaria vaccine, is unavailable at this time. Removing this intervention from the program still results in 15- to 25-fold reductions in EIR. Note also that these high reductions in EIR are accomplished with modest, 5-fold reductions in the overall human biting rate, H_{Bt} . This is important to bear in mind because H_{Bt} is the factor that communities often more readily perceive and are motivated by.^{62,65,66} Furthermore, in areas of very high transmission, a large impact on EIR may not be as readily appreciated as in areas of low transmission even though the impact, in terms of averted deaths, is equivalent.²⁰ Because the overall disease burden in high transmission areas is much higher, reductions in mortality seem smaller when expressed and perceived as a proportion.²⁰ Alleviation of mortality burden appears to be a consistent outcome, although the impact of transmission control in terms of protective efficacy is lower in areas with high baseline transmission intensity and with better medical facilities.^{12,14,20}

The predicted impact of the combined control measures implies that an integrated approach is preferable for controlling intense malaria transmission and can achieve consistently large reductions of EIR, irrespective of baseline value. This is consistent with the fact that bed nets have roughly the same impact on absolute mortality over a wide range of transmission intensity.²⁰ Clearly, integrated malaria transmission control should be implemented to the fullest extent possible in all endemic areas.^{2,12,14,20} Perhaps the most important question remaining with regard to bed nets and other transmission control measures is how to implement their widespread use rather than whether they are effective.^{12,20,67} We suggest that this model may be useful for deciding how to combine different tools when planning vector-control programs.

Despite their dramatic predicted impact on EIR, the integrated control programs modeled here failed to reduce EIR to less than one at any of the four sites, even with the theoretical inclusion of a malaria vaccine. Endemic malaria with its concomitant disease burden is readily maintained by EIRs considerably below this level.² Thus, although meaningful reductions in malaria burden can be achieved with existing technology, elimination from areas of intense transmission is currently beyond the grasp of many developing nations in Africa and the southwest Pacific. Techniques being explored at present include odor-baited traps, vector population replacement, vaccines, and biological control, but none of these interventions are likely to yield useful tools in the immediate future.^{68,69}

The search for ways to combat malaria transmission should, perhaps, look as much to the past as to the future. In the interest of a conservative estimate, we incorporated a reduction of only 50% to vector emergence rates as being feasible at these sites. However, we suggest that far greater reductions can be achieved with existing tools, which have come to be regarded as old-fashioned. Although members of the *An. gambiae* complex, in particular, are notoriously difficult to control, complete elimination has been accomplished in both Brazil³¹ and Egypt³² in less than three years, even in areas with extensive seasonal flooding. Many of the

habitats cleansed of *An. gambiae* in Brazil, appear to have been as ideal for *An. gambiae* proliferation as they were challenging to the implementation of control programs.³¹ The Egyptian campaign was executed with very limited resources during the peak of World War II, necessitating the use of sand rather than oil as a larvicide diluent.³² Despite these considerable obstacles, complete elimination was achieved in both instances by combining simple larvicides with effective, practical, and determined management of human resources.^{31,32} Although many of the principles of these vector-eradication campaigns are not practicable under the conditions of sustainable malaria transmission control programs, three key contributors are worth highlighting: 1) political support and freedom from bureaucratic obstacles, including full authority to enforce necessary measures, 2) clear chains of command and individualization of responsibilities, and 3) assignment of control and surveillance responsibilities to separate personnel.

Sustainable environmental tools for larval habitat source reduction and biological control are also well established,¹⁹ but remain grossly underused in most endemic areas. Such methods center on managing resources for food, water, and fuel. We suggest that such methods can meaningfully reduce malaria transmission intensity but that their broader application will require greater coordination between those concerned with malaria control and those concerned with agricultural, industrial, and economic development.

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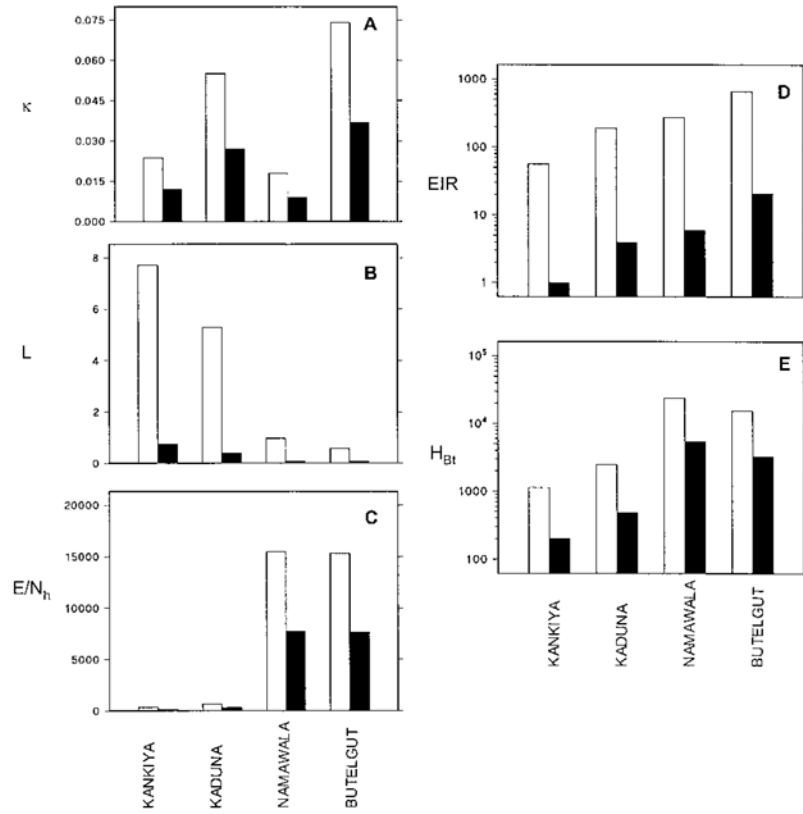


Figure 1.

A: Simulated between-site variations in the size of the infectiousness of the human reservoir κ . B: The lifetime malaria transmission capacity of individual vectors (L); C: Mosquito emergence rate relative to the size of the human population E/N_h ; D: The entomologic inoculation rate (EIR); and E: The human biting rate (H_{Bt}). Baseline conditions are represented by empty bars and those of an integrated transmission control program by solid bars. For further details of definitions and units see Killeen and others.¹

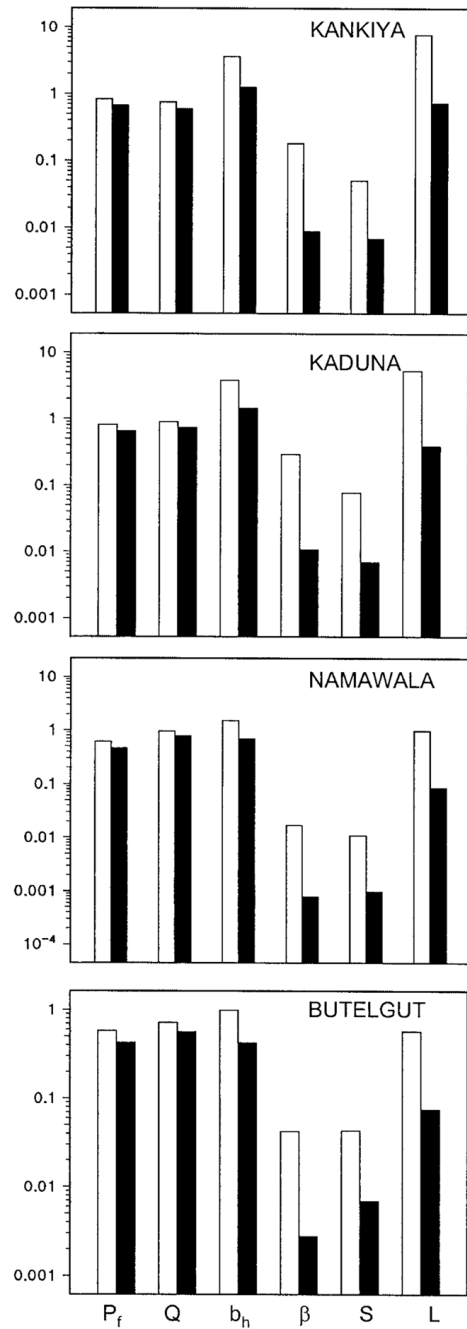


Figure 2.

Predicted between-site variations in vector survival per feeding cycle (P_f), blood-feeding preferences (Q), number of human bites per lifetime (b_h), number of infectious human bites per lifetime (β), sporozoite prevalence (S), and infectious bites per lifetime per bite on an infective human host (L) under baseline conditions (empty bars) and those of an integrated transmission control program (solid bars). See Killeen and others¹ for further details of definitions and units.