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# **Diversity Cascades and Malaria Vectors**

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# Abstract

The interactions between predator diversity and primary consumer abundance can include direct effects and indirect, cascading effects. Understanding these effects on immature *Anopheles* mosquitoes is important in sub-Saharan Africa, where most cases of malaria occur. Aquatic predators and immature mosquitoes were collected from shallow pools of varying age previously excavated by brickmakers in the western highlands of Kenya. Path analysis showed an indirect negative effect of habitat age on *An. gambiae* (Giles, 1902) mediated by effects on predator diversity. Disturbance resets habitats to an earlier successional stage, diminishing predator diversity and increasing *An. gambiae* populations. The increase in vector abundance as a result of reduced predator diversity highlights the public health value in conserving native insect diversity.

#### Keywords

trophic cascades; diversity; Anopheles gambiae; malaria; Kenya

Diversity cascades are the indirect interactions between diversity and other community metrics across trophic levels. The effect of predator diversity on primary consumers (i.e., top-down diversity cascades, sensu Dyer 2008) varies depending on the food web being studied, but it is clear that biodiversity can have significant population-level effects on disease vectors (Ostfeld and Keesing 2000). For example, incidence of Lyme disease is positively correlated with species richness of good disease hosts, such as birds, but the densities of effective arthropod vectors of Lyme disease are negatively correlated with the species richness of small mammals, which vary widely in their ability to carry the disease. Increases in predator diversity can promote herbivore abundance by increasing intraguild predation and omnivory among predators (Hochberg 1996, Denoth et al. 2002), further showing that diversity at one trophic level can act to enhance or constrain populations at other levels. Alternately, if specialist predators are present, increased predator diversity may reduce the abundance of a particular prey species. In such a case, as predator diversity increases within a habitat, the probability of specialist predators being present also increases (Stireman et al. 2004). Several studies of topdown diversity cascades have been published by terrestrial ecologists (Dyer and Letourneau 2003, Dyer and Stireman 2003, Schmitz 2004, Pearson and Dyer 2006, Dyer 2008, Pearson et al. 2008), whereas the effect of predator diversity on the aquatic food webs that are dominated by immature mosquitoes has not been examined in detail, despite the large number of singlepredator studies performed in such systems (Lacey and Orr 1994).

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Enhancement or conservation of aquatic predator species richness may be an effective supplement to existing malaria control programs, particularly in Africa where a substantial majority of the world's estimated 515 million malaria cases occur (Snow et al. 2005). Malaria epidemics in the African highlands have generated a great deal of political and scientific interest, and an ongoing effort to determine why malaria has spread to these previously malariafree locations (Malakooti et al. 1998, Hay et al. 2002) and to establish methods for preventing these epidemics from recurring (Delacollette 2002, Abeku 2007). Surveys of aquatic habitats in the highlands of western Kenya have suggested that altered land use plays an important ecological role in determining the distribution of mosquito larvae (Carlson et al. 2004, Omlin et al. 2007, Howard and Omlin 2008); in particular, in many highland valleys, the excavation of clay for the large-scale production of bricks has generated hundreds of shallow pools that support high densities of immature Anopheles gambiae (Giles, 1902) but very few predator species (Carlson et al. 2004). In this paper, An. gambiae is sensu Giles, 1902, referring to a cryptic species complex, frequently denoted as An. gambiae s.l. for studies in which polymerase chain reaction (PCR) identification of the sibling species was not accomplished. Members of this complex are the primary malaria vectors in sub-Saharan Africa. Similarly, An. funestus is used sensu Giles, 1900; members of this cryptic species complex are secondary vectors of malaria in sub-Saharan Africa.

This study examined simple aquatic food webs in the same system described above: shallow brickmaking pools in the highlands of western Kenya. The following general hypotheses were tested:

- 1. Predator diversity and prey abundance will be positively associated with the length of time since a habitat was last disturbed (hereafter referred to as habitat age) caused by successional changes in habitat quality.
- 2. Habitat age affects *An. gambiae* abundance both directly, through discriminating oviposition behavior, and indirectly, through direct and indirect effects on predator diversity.

#### Materials and Methods

Mosquito larvae and aquatic predators were collected from shallow pools of varying ages, and path analysis was used to test hypothesized causal relationships between habitat age and predator diversity and mosquito abundance.

#### Study System

This study was conducted in Itumbe Valley, Kisii District, one of many similar valleys in the densely populated highlands of western Kenya. Itumbe Valley is the site of intensive brick production, for which clay is excavated and processed in shallow water-filled pits. These habitats were linked to proliferation of *An. gambiae* after a 2002 malaria epidemic in the region (Carlson et al. 2004).

#### Field Methods

One hundred brick-making pits were evaluated in April 2003. Abundance of immature mosquitoes was obtained by standard dipping method, averaging the yields of five dips per pit (Service 1993). Identification of immature mosquitoes was constrained by field conditions; larvae were identified as *An. gambiae*, *An. funestus*, or belonging to the culicine "competitor" mosquito genera. Macroscopic predators were qualitatively sampled by aquatic sweep net for a duration of ~5 min in each pit. Invertebrate predators were later separated into distinct morpho-species, and voucher specimens were preserved in ethanol for subsequent taxonomic

identification. To establish habitat age, brick-makers were questioned regarding the date when each pit had last been used for clay extraction or processing.

#### **Conceptual Model**

Models incorporating *Anopheles* abundance began soon after the discovery was made that malaria was spread by mosquitoes (Ross 1911). As the complexity of the models has increased, there has been a tendency to examine the variables involved in producing adult mosquitoes separately from the interactions between the adult mosquitoes, humans, and *Plasmodium* parasites. The most comprehensive model available that describes adult *Anopheles* production hypothesizes that the effects of predation and competition are important variables affecting *Anopheles* larvae (Depinay et al. 2004).

In addition to the direct effects of predation and competition on Anopheles larvae, it is likely that there are important interactions within and between populations of predators and competitors. In this regard, the abundance of competitors may have an additional indirect effect on Anopheles larvae by altering the abundance, diversity, or feeding habits of predators (Chesson 1989). Like other dispersing insects, An. gambiae mosquitoes use various combinations of visual (Huang et al. 2005) and olfactory (Sumba et al. 2008) cues to select high-quality aquatic habitats for oviposition. Some of these cues are associated with biological succession of the habitat, such as the extent of microbial growth (Huang et al. 2006) and the presence of top predators (Munga et al. 2006). Thus, successional stage of a habitat is expected to play an important role in determining abundance of An. gambiae larvae directly by affecting oviposition behavior, and indirectly by affecting colonization rates of predators and competitor mosquitoes. The hypothesized relationships among these latent variables- habitat disturbance, predator diversity, competitor abundance, and vector abundance-and the ultimate production of adult vector mosquitoes were delineated in a conceptual model (Fig. 1), guided by previous structural equation models of diversity cascades in agricultural systems and grasslands (Dyer and Stireman 2003, Pearson and Dyer 2006).

#### **Statistical Analyses**

To conform to the prerequisites for multiple regression analysis, all variables were tested and, with the exception of predator diversity, log transformed to meet assumptions of normality and homogeneity of variance. Pathway coefficients were generated using a series of multiple regressions using the CALIS procedure (SAS Institute 1989). These standardized coefficients, representing the relative magnitude of the relationships, are presented on a path diagram (Fig. 2).

## Results

Mosquito dips from the sampled pits yielded an average of  $8.0 \pm 0.95$  *An. gambiae* larvae and  $5.7 \pm 0.72$  culicine larvae. A smaller number of *An. funestus*  $(1.8 \pm 0.32)$  larvae were also collected. Across all pools, there were 28 predator morphospecies collected, representing 21 genera and 12 families (Table 1). The most common predators were Hemiptera (12 morpho spp.), including *Hynesionella* (Gerridae) and *Anisops* (Notonectidae), the only two genera found in >50% of the sampled pools. The next most common were Coleoptera, in particular the nine morphospecies of Dytiscidae, which were present in some combination in 52% of habitats. Mean species richness of predators for all pools was  $5.0 \pm 0.25$ . The average time since last disturbance of pools was  $16.4 \pm 2.8$  mo, ranging from 0 (actively disturbed) to 13 yr since disturbance.

Path analysis showed the following direct and indirect interactions: (1) strong positive effects of habitat age on predator diversity and on abundance of culicine competitors; (2) strong

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negative effects of predator diversity on *An. gambiae* abundance; and (3) a strong indirect negative effect of habitat age on *An. gambiae* abundance (Fig. 2). Our data support the hypothesis that habitat age both directly and indirectly increased predator diversity, which in turn depressed *An. gambiae* abundance; the model was supported by the data ( $\chi^2 = 0.88$ , df = 1, P = 0.35) and the path coefficients were significant (P < 0.05).

## Discussion

Changes to predator diversity of any magnitude, even the loss of a single species, can have major impacts on ecosystems and on infectious disease risk (Keesing et al. 2006). This study showed the important role of aquatic predator diversity in suppressing local populations of the predominant malaria vector. Predator diversity was markedly diminished by habitat disturbance and recovered as biological succession of the habitat progressed over time. Part of this effect is attributable to the reduction of competitor mosquito populations, which may serve as an alternate food source for predators. In recently disturbed habitats, the reduction in predator diversity was associated with greater abundance of *An. gambiae*, the primary malaria vector in Africa. Ecological disturbance resulting from altered land use in highland swamps is further implicated as a possible cause for the puzzling increase in highland malaria (Minakawa et al. 2006).

In contrast to insecticidal control, biological control by a diverse predator assemblage is likely to persist over time (Ostfeld and Holt 2004). For example, Shililu et al. (2003) compared the efficacy against *Anopheles* larvae of the organophosphate temephos and two biolarvicides (*Bacillus thuringiensis isrealensis* and *B. sphaericus*) in a variety of aquatic habitats in Eritrea. For all three larvicides, mosquitoes were almost entirely eliminated from standing streambed pools in the days immediately after treatment, but within 14 d after treatment, mosquito densities were equivalent to untreated habitats (mean of 15.36 larvae per pool). Although vector control through use of larvicides is possible in aquatic habitats, it requires weekly evaluation and/or treatment (Fillinger and Lindsay 2006). Because competitor abundance has a positive effect on predator diversity, larvicidal elimination of non-target organisms (including other Dipteran larvae) may indirectly promote a resurgence of *An. gambiae* larval populations even when predators are not harmed by the treatment itself.

In this study, it is clear that diversity of aquatic predators is important for suppressing *An. gambiae* larval abundance and is associated with habitats that have remained undisturbed for an extended period of time. Although larval abundance is only one factor influencing subsequent vector-biting rate and malaria transmission, reductions in malaria cases have been observed after large-scale implementation of larval control (reviewed by Walker and Lynch 2007). This study found statistical support for a strong interaction between predator diversity and *An. gambiae* abundance, indicated by the magnitude of the path coefficient (-0.15, from Fig. 2); in the context of the study design, each additional predator species in a brick-making pit reduced the number of *An. gambiae* by approximately one larva per dip sample.

This study did not incorporate the rapidly diminishing areas of undisturbed wetlands in the Kenyan highlands, wherein the diversity of predators is expected to prohibit establishment of substantial *An. gambiae* populations. The benefits of ecosystem services have been cited as a compelling incentive for the preservation of biodiversity in tropical areas (Daily et al. 1997, Keesing et al. 2006). The local paucity of predator species after human disturbance is linked with the proliferation of malaria-vectoring mosquitoes in the mountains of western Kenya, implicitly showing the value of natural enemy diversity to human health.

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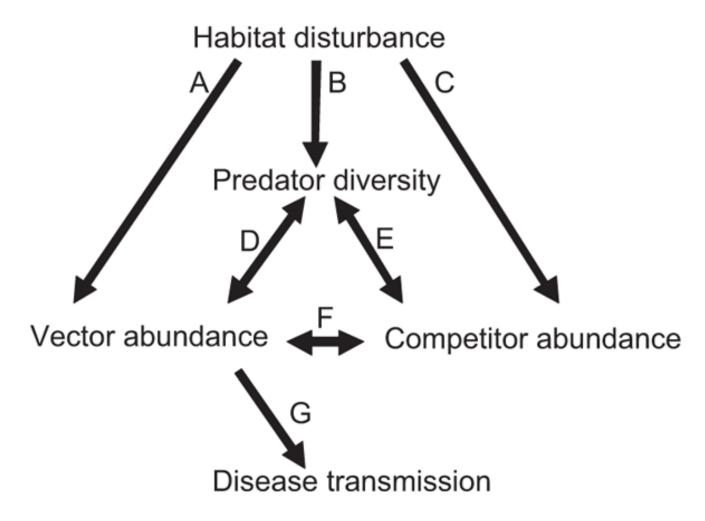
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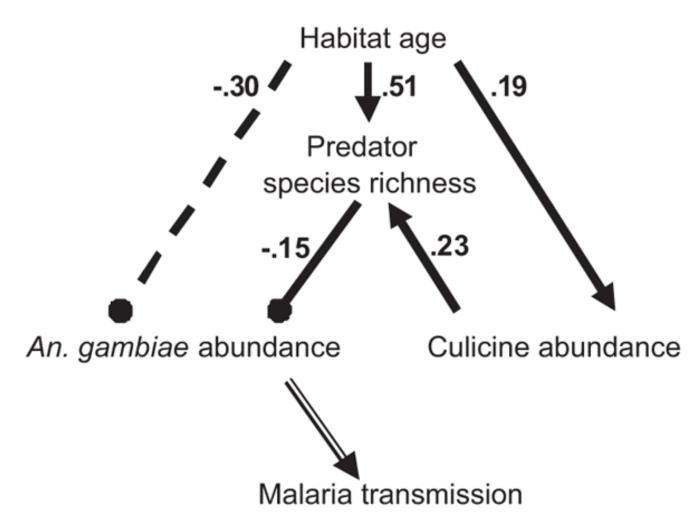
#### Fig. 1.

Conceptual model showing a hypothesized set of interactions in aquatic ecosystems that could combine to influence malaria transmission. Habitat condition affects immature vector abundance directly (A) by influencing oviposition behavior of gravid female *Anopheles* mosquitoes. Habitat similarly affects colonization by predators (B) and competitors (C), although the magnitude and sign of the effect may be inconsistent. The interactions between predator diversity and vector abundance (D) may be mitigated or enhanced by additional interactions between predators (F). Direct and indirect effects on vector abundance ultimately influence the more well-established relationship between vector abundance and malaria transmission (G).

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#### Fig. 2.

This causal pathway diagram is statistically supported by field data (i.e., no significant differences between the correlational matrix and that predicted by the causal model;  $\chi^2 = 0.88$ , df = 1, P = 0.35). Lines ending in arrowheads indicate a positive effect; lines ending in circles indicate a negative effect. Solid lines represent direct effects, whereas the dashed line shows an indirect effect. Only the relationships with statistically significant path coefficients ( $\alpha = 0.05$ ) are shown in the diagram, with the exception of the relationship between *An. gambiae* abundance and malaria transmission (indicated with a double lined arrow), but was not measured in this study. Habitat age had a direct positive effect on both predator diversity and abundance of culicine mosquito larvae but no detectable direct effect on *Anopheles* abundance. Habitat age also indirectly promoted predator diversity thorugh a direct enhancement of culicine larvae (alternative prey), which in turn acted to enhance predator colonization. The direct and indirect increases in predator diversity caused by habitat age resulted in an overall reduction of *An. gambiae* larvae, overwhelming any potential positive effect of habitat age on vector abundance.

# Table 1 100 brickmaking pits in Itumbe Valley, K

# Insect predators collected from 100 brickmaking pits in Itumbe Valley, Kenya

Order	Family	Genus	% habitats colonized
Hemiptera	Gerridae	Hynesionella	66
		Limnogonus	5
	Hydrometridae	Hydrometra	12
	Veliidae	Rhagovelia	40
	Notonectidae	Anisops	68
		Enithares	5
	Pleidae	Plea	12
	Naucoridae	Laccocoris	3
	Nepidae	Ranatra	8
		Laccotrephes sp 1	23
		Laccotrephes sp 2	18
		Nepa	2
Odonata	Lestidae	Lestes	32
	Coenagrionidae	Enallagma	25
	Libellulidae	Palpopleura	20
		Orthetrum	16
Coleoptera	Hydrophilidae	Hydrochara	3
	Dytiscidae	Laccophilini sp. 1	32
		Laccophilini sp. 2	12
		Laccophilini sp. 3	9
		Laccophilini sp. 4	2
		Laccophilini sp. 5	2
		Laccophilini sp. 6	1
		Copelatus	5
		Cybister	7
		Hydaticus	3