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Disease ecology, health and the environment: a framework to account for ecological and socio-economic drivers in the control of neglected tropical diseases

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Reducing the burden of neglected tropical diseases (NTDs) is one of the key strategic targets advanced by the Sustainable Development Goals. Despite the unprecedented effort deployed for NTD elimination in the past decade, their control, mainly through drug administration, remains particularly challenging: persistent poverty and repeated exposure to pathogens embedded in the environment limit the efficacy of strategies focused exclusively on human treatment or medical care. Here, we present a simple modelling framework to illustrate the relative role of ecological and socio-economic drivers of environmentally transmitted parasites and pathogens. Through the analysis of system dynamics, we show that periodic drug treatments that lead to the elimination of directly transmitted diseases may fail to do so in the case of human pathogens with an environmental reservoir. Control of environmentally transmitted diseases can be more effective when human treatment is complemented with interventions targeting the environmental reservoir of the pathogen. We present mechanisms through which the environment can influence the dynamics of poverty via disease feedbacks. For illustration, we present the case studies of Buruli ulcer and schistosomiasis, two devastating waterborne NTDs for which control is particularly challenging.

This article is part of the themed issue 'Conservation, biodiversity and infectious disease: scientific evidence and policy implications'.

1. Introduction

More than one billion people, most of them in tropical and subtropical countries, still live in extreme poverty and suffer a disproportionate burden of neglected tropical diseases (NTDs) [1,2]. NTDs consist of parasitic, viral and bacterial infections (table 1) that are responsible for chronic disabling conditions that account for more than 500 000 deaths and nearly 57 million disability-adjusted life-years

Table 1. NTDs, causes, global burden and current priorities for disease control (based on WHO 2015 [3]).

NTD Group	pathogens responsible	transmission route	disease burden	control target and main strategies ^a
<i>Parasitic infections</i>				
soil-transmitted helminthiasis	<i>Ascaris</i> spp., <i>Trichuris</i> spp., <i>Necator</i> spp. and <i>Ancylostoma</i> spp.	<i>Environmental</i> . Ingestion of eggs and larvae or penetration of the skin	875 million children at risk	<i>Morbidity control</i> . MDA of PC (albendazole or mebendazole)
schistosomiasis	<i>Schistosoma</i> spp. (mostly <i>S. haematobium</i> and <i>S. japonicum</i>)	<i>Waterborne</i> . Direct entry into skin after asexual multiplication in snail hosts	240 million infected	<i>Elimination</i> (some regions). MDA of PC (praziquantel)
lymphatic filariasis	<i>Wuchereria bancrofti</i> , <i>Brugia malayi</i> and <i>B. timori</i>	<i>Vector-borne</i> . Mosquito bites (<i>Anopheles</i> spp., <i>Culex</i> spp. and <i>Aedes</i> spp.)	more than 120 million infected	<i>Elimination</i> . MDA of PC (albendazole + diethylcarbamazine or ivermectin)
onchocerciasis	<i>Onchocerca volvulus</i>	<i>Vector-borne</i> . Bites of blackflies (<i>Simulium</i> spp.)	37 million infected	<i>Elimination</i> . MDA of PC (ivermectin)
dracunculiasis (Guinea worm)	<i>Dracunculus medinensis</i>	<i>Waterborne</i> . Drinking water with infected copepods	100–200 cases/year	<i>Eradication</i> . Active surveillance, water filters, and improved drinking water sources
foodborne trematodiasis	<i>Clonorchis</i> spp., <i>Opisthorchis</i> spp., <i>Fasciola</i> spp. and <i>Paragonimus</i> spp.	<i>Foodborne</i> . Consumption of contaminated fish, seafood or vegetables	unclear; more than 56 million in 2005	<i>Morbidity control</i> . MDA of PC (praziquantel or triclabendazole)
taeniasis/cysticercosis	<i>Taenia solium</i>	<i>Foodborne or faecal–oral</i> . Consumption of pork or ingestion of parasite eggs	unclear; responsible for 1.5 million epilepsy cases	<i>Reduce transmission</i> . Veterinary controls of pork meat; treatment of infected pigs and humans
echinococcosis	<i>Echinococcus granulosus</i> and <i>E. multilocularis</i>	<i>Foodborne</i> . Accidental ingestion of items contaminated with eggs	unclear; 30/100 000 person-year, endemic areas	<i>Reduce transmission</i> . Veterinary controls of livestock; dog vaccination and regular treatment
<i>Protozoan infections</i>				
human African trypanosomiasis	<i>Trypanosoma brucei gambiense</i> and <i>T. brucei rhodesiense</i>	<i>Vector-borne</i> . Bites of tse-tse flies (<i>Glossina</i> spp.)	20 000 cases/year	<i>Elimination</i> . Early diagnosis and treatment (pentamidine, suramin, melarsoprol or eflornithine)
Chagas disease	<i>Trypanosoma cruzi</i>	<i>Vector-borne</i> . Faeces of haematophagous triatomines	seven million infected	<i>Reduce transmission</i> . Sustained vector control; early diagnosis and treatment (benznidazole or nifurtimox)
leishmaniasis	<i>Leishmania</i> spp.	<i>Vector-borne</i> . Bites of sandflies (<i>Phlebotomus</i> spp.)	1.3 million cases/year	<i>Elimination</i> (some regions). Surveillance, early diagnosis and treatment (liposomal amphotericin B)
<i>Bacterial infections</i>				
Buruli ulcer	<i>Mycobacterium ulcerans</i>	<i>Waterborne</i> . Vector-borne and/or direct environmental transmission (unclear)	5000 cases/year	<i>Prevention of disability</i> . Early diagnosis and treatment (streptomycin + rifampin)
leprosy	<i>Mycobacterium leprae</i>	<i>Person-to-person</i> . Skin contact or respiratory route (unclear)	200 000 cases/year	<i>Elimination</i> . Early diagnosis and treatment (multidrug therapy)

(Continued.)

Table 1. (Continued.)

NTD Group	pathogens responsible	transmission route	disease burden	control target and main strategies ^a
trachoma	<i>Chlamydia trachomatis</i>	Person-to-person. Direct personal contact, shared towels and cloths, or flies	more than 21 million with active trachoma	Elimination. MDA of PC (azithromycin)
yaws (endemic treponematoses)	<i>Treponema pallidum</i>	Person-to-person. Skin contact	50 000–80 000 cases/year	Eradication. MDA of PC (azithromycin); early diagnosis and treatment
<i>Viral infections</i>				
dengue and chikungunya	Arbovirus (dengue virus and chikungunya virus)	Vector-borne. Mosquito bites (<i>Aedes</i> spp.)	390 million dengue infections/year	Reduce mortality, morbidity and transmission. Integrated surveillance and vector control
rabies	rabies virus	Zoonosis. Bite from infected dogs	60 000 deaths/year	Elimination. Mass vaccination of domestic dogs

^aMDA of PC, mass drug administration of preventive chemotherapy.

(DALYs) annually [4,5]. The availability of affordable and safe drug treatments has led to declines in the prevalence of many NTDs, with plans for the elimination of lymphatic filariasis, onchocerciasis and trachoma by 2030 (table 1) [3,6]. Nevertheless, there is growing recognition of the limitations of interventions focused exclusively on treating the human host [3,7]. Most pathogens responsible for NTDs spend part of their life cycle outside the human host and are transmitted through environmental pathways (table 1), such that people may be repeatedly exposed and infected [7]. Moreover, coverage targets for mass drug administration (MDA) are hard to meet in the context of weak healthcare-delivery systems and extreme poverty, and drug resistance can threaten the long-term success of control programmes [8]. Sustainably reducing the burden of many NTDs requires complementary strategies to disrupt disease transmission that are informed by a broader understanding of interactions taking place in coupled natural–human systems [9].

Here, we present a conceptual framework based on a simple model to account for ecological and socio-economic drivers of NTDs for designing context-specific control strategies. As examples, we present the cases of Buruli ulcer and schistosomiasis, two NTDs associated with land-use change in hydrological ecosystems, which represent different ends of the spectrum of diseases for which transmission is highly dependent on the environment (referred to hereafter as ‘environmentally transmitted’). Buruli ulcer is caused by *Mycobacterium ulcerans* which, like other ‘sapronotic’ pathogens [10], can be considered ubiquitous and free-living within the environment (i.e. not dependent on the level of infection within humans or any other known host) in certain regions. It afflicts only a few thousand people per year, but each case of infection can cause permanent disability with catastrophic socio-economic effects on households. Schistosomiasis, which is caused by *Schistosoma* spp., is representative of a larger class of parasitic worms and vector-borne pathogens whose presence in the environment depends upon the local prevalence of infected people. Schistosomiasis infects hundreds of millions of people each year and is responsible for a high burden of chronic morbidity globally. Integration of ecological knowledge and socio-economic feedbacks into a common framework can thus inform the most effective action targets and lead to better interventions for control of NTDs and other infectious diseases of poverty.

2. A framework for coupled ecological and socio-economic determinants of disease

Economic systems of the rural poor are often coupled with natural systems through subsistence-resource exploitation (e.g. agriculture, pastoralism and fishing) and exposure to ambient pathogens (figure 1) [11,12]. This is particularly relevant in the tropics, where climatic conditions are favourable for many pathogens and their vectors (e.g. flies, mosquitoes and ticks), and the diversity of mammal and bird species hosts a greater diversity of pathogens in this region [13–16]. Land-use change for the purpose of economic development (e.g. expansion of road networks, dams and agricultural land conversion) can further affect environmentally transmitted parasites and pathogens through impacts on physicochemical conditions of water and soil [15], ecological community composition (e.g. loss of biodiversity), or inter- and intra-species interactions (e.g. predation and migration) [17,18].

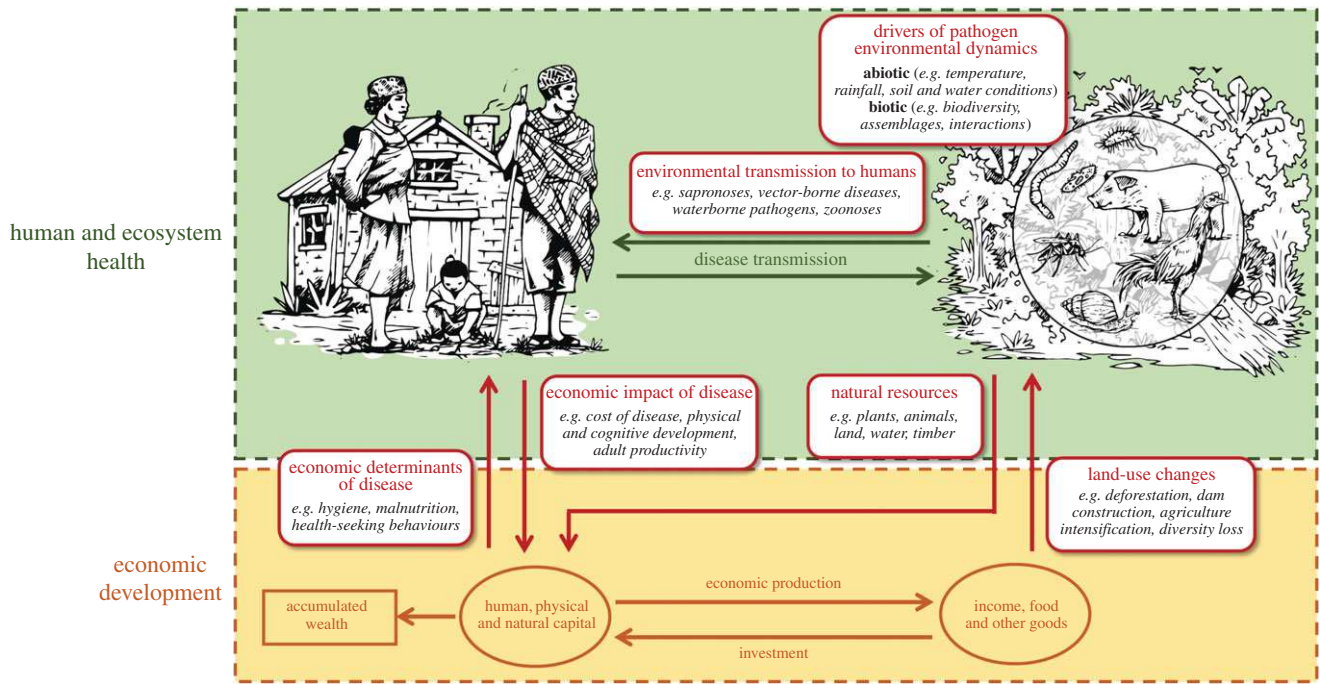


Figure 1. Coupled natural and human systems: feedbacks between ecosystems, infectious diseases and economic development. Many human pathogens in tropical regions, including those that cause NTDs, spend part of their lifecycle outside the human host, where environmental conditions drive their ecology and transmission (upper panel). Ecosystem and disease dynamics influence key forms of capital necessary for economic development of at-risk human communities (lower panel). Simultaneously, economic activities and resources affect those same dynamics by altering vulnerability to disease and introducing land-use change to ecosystems.

(a) Model of environmental transmission of disease

Building on traditional systems in disease ecology and epidemiology [19–21], we first present a general infectious disease model that accounts for direct and environmental pathways of disease transmission. The model is meant to represent the dynamics of NTDs and other tropical diseases of poverty across a wide spectrum of transmission strategies, spanning from human-to-human contagious diseases (e.g. leprosy and trachoma) to parasitic worms (e.g. schistosomiasis), waterborne bacteria (e.g. Buruli ulcer), foodborne pathogens (e.g. echinococcosis), vector-borne diseases (e.g. dengue and malaria), or zoonotic and multi-host pathogens (e.g. rabies and Nipah virus):

$$\frac{dI}{dt} = [\beta_E W + \beta_D I] S - \gamma I \quad (2.1a)$$

and

$$\frac{dW}{dt} = \Omega + V\sigma\lambda I - \delta W. \quad (2.1b)$$

The state variables I and S represent the infected and susceptible proportions of the human population, respectively; W is the environmental reservoir of the pathogen; β_D is the direct (human-to-human) transmission rate; β_E is the transmission rate through the environmental reservoir; γ is the rate of recovery (either into an immune class in an SIR model or back to the susceptible class in an SIS model without immunity); λ is the rate or production of infectious propagules released in the environment (either as free-living stages, such as for soil-transmitted helminths, or pathogen abundance in vectors or intermediate hosts); σ is the fraction of the propagules that reach the environment; V is a parameter proportional to the abundance of vectors (in the case of dengue, chikungunya, etc.), to the abundance of intermediate hosts

(in the case of trematode infections, such as schistosomiasis, fascioliasis, etc.), or to habitat suitability for pathogen environmental persistence (in the case of soil-borne or waterborne diseases, such as intestinal nematode infections and cholera, respectively); Ω is the recruitment rate of infective propagules associated with exogenous processes not directly related to disease prevalence in the population; δ is the mortality of infectious stages in the environment. From this general framework, we define two parameters (or ‘meta-parameters’) that characterize the role of the environment on transmission:

$$\varepsilon = \frac{\beta_E W}{\beta_E W + \beta_D I} \quad (2.2a)$$

and

$$\varphi = \frac{\Omega}{\Omega + V\sigma\lambda I}. \quad (2.2b)$$

Parameter ε is an index of environmental transmission that ranges from zero for direct transmission (i.e. person-to-person) to unity for complete environmental transmission (i.e. pathogens with obligatory non-human hosts, reservoirs or vectors). Parameter φ represents the degree of exogenous versus focal transmission among the environmentally transmitted diseases (ETDs). It ranges from zero when the abundance of infectious propagules in the environmental reservoir is solely a function of the local number/fraction of infected people, with no exogenous recruitment ($\Omega = 0$ and $\lambda > 0$), to unity when the pathogen’s environmental load depends exclusively on its own free-living replication in the environment or on other environmental factors ($\Omega > 0$ and $\lambda = 0$). ‘Sapronotic’ pathogens—those that can replicate within the environment free of any known host (a characteristic of almost all fungal pathogens, about one-third of bacterial pathogens and several parasites that infect man [10])—have

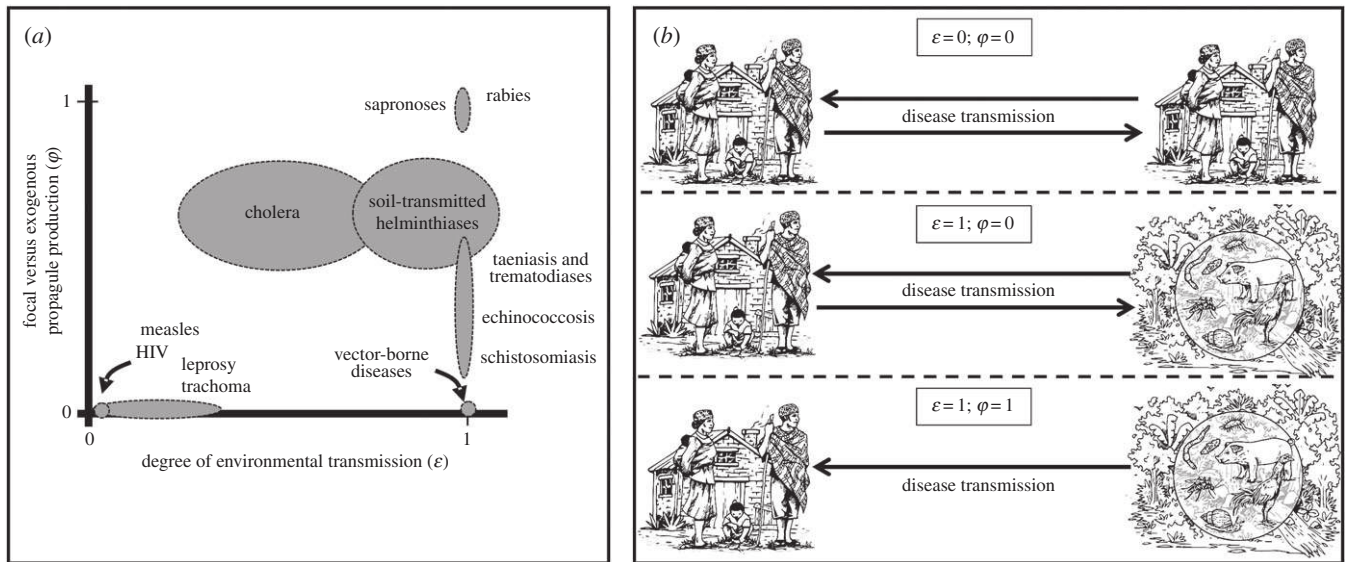


Figure 2. A conceptual framework to account for multiple pathways of disease transmission in coupled economic–epidemiological systems. (a) Qualitative classification of relevant infectious and parasitic diseases in developing countries according to two key parameters that regulate environmental transmission in this conceptual framework (ε and φ). Ellipsoids represent educated guesses about the parameter space occupied by each disease, including major NTD groups. (b) Graphical representation of the three main groups of diseases according to parameters ε and φ : directly transmitted ($\varepsilon = 0$; $\varphi = 0$), environmentally transmitted, with focal transmission dependent on human prevalence ($\varepsilon = 1$; $\varphi = 0$) and environmentally transmitted, driven purely by exogenous environmental factors ($\varepsilon = 1$; $\varphi = 1$).

by definition $\varphi = 1$, while directly transmitted diseases have, by the nature of the close contact required for contagion, $\varphi \rightarrow 0$. For other parasites and pathogens, for example, those that experience intermediate levels of focal to exogenous transmission, the value of φ depends on the scale of analysis.

Figure 2 provides a simple qualitative description of common infective agents in tropical regions according to their transmission strategies based on parameters ε and φ . Many pathogens responsible for mortality in developing countries have low ε and φ , such as childhood diseases (measles and whooping cough), respiratory infections (tuberculosis) or sexually transmitted diseases (HIV and syphilis). By contrast, for most pathogens responsible for NTDs, environmental transmission plays an important role ($\varepsilon \rightarrow 1$). The general model described above, although difficult to parametrize in its simplest form [19,22], can be used to explore conceptual differences in dynamics and control strategies for a wide variety of pathogens and transmission strategies. A heuristic illustration of such insights is provided in figure 3, where we contrast the impact of common strategies of disease control for directly transmitted diseases ($\varepsilon = 0$, $\varphi = 0$) and ETDs ($\varepsilon = 1$, $\varphi \rightarrow 1$).

We show that for direct and ETDs with comparable basic reproduction numbers R_0 (i.e. number of secondary infections caused by an infected individual introduced in a susceptible population), control can be effectively achieved for directly transmitted disease through periodic medical treatment. All else being equal (i.e. in terms of the R_0 of the disease and the frequency and intensity of intervention), the same drug-focused strategies can have a reduced efficacy for diseases where the environmental reservoir plays an important role (figure 3*a,b*). In the latter case, more effective control can be achieved when classic treatment strategies are complemented with interventions that act on the environmental reservoir of the pathogen or reduce exposure (figure 3*c*). A detailed analysis of this system of equations, the parametrization for the presented simulations, and further explanations are provided in the electronic supplementary material, Appendix S1.

(b) Feedbacks between poverty and disease in coupled environment–human systems

Global patterns of poverty and disease have a strong geographic signature, indicative of underlying environmental determinants (figure 4) [26,27]. Building on recent advances in the modelling of coupled economic–epidemiological systems [11,28], we explore how the framework presented in §2*a* can be integrated with existing models of disease-driven poverty traps to understand the potential role of environmental drivers. This integrated framework is based on the intersection of the following empirical facts: (i) economic and environmental conditions are co-determinants of NTDs, and (ii) NTDs have well-documented negative effects on economic productivity, primarily through child development, educational attainment and labour productivity [1,27,29–31].

The effect of economic drivers on disease dynamics in a particular system can be captured by transforming some parameters in equations (2.1*a,b*) to be functions of *per capita* income or wealth: i.e. $\beta_D = \beta_D(k)$, $\beta_E = \beta_E(k)$, $\Omega = \Omega(k)$, $\lambda = \lambda(k)$; where k is capital, which we define broadly as a stock of economic resources used to produce income. For instance, transmission rates (β_D , β_E) and environmental recruitment of propagules (σ and Ω) are generally decreasing functions of capital k , because capital is necessary for prevention measures that reduce disease transmission, such as systems for water and sanitation, improved housing conditions or protective clothing [4,5,7]. In addition, the recovery rate $\gamma = \gamma(k)$ is generally an increasing function of capital because capital (or income generated from capital) is used for better nutrition and accessing healthcare [32,33]. The relationship of vectors $V = V(k)$ and the mortality rate of infective propagules $\delta = \delta(k)$ with capital (k , or economic development) are more context- and disease-specific [28].

The effects of disease on income can be captured by adding capital as a third state variable, in the tradition of economic

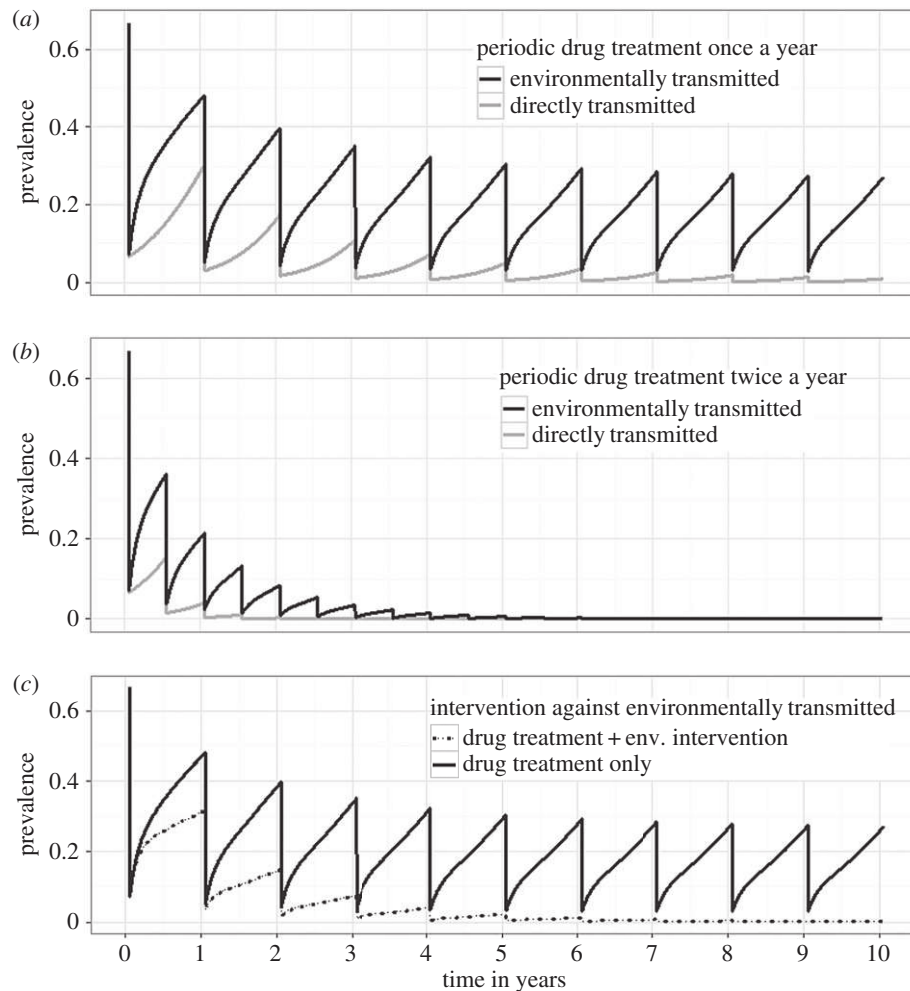


Figure 3. Effectiveness of periodic drug treatment and environmental interventions for the control of diseases with different transmission pathways. Population prevalence dynamics over 10 years for a directly transmitted disease (grey lines) and an ETD (black lines) with human drug treatment dispensed once a year (a) or twice a year (b). We considered a 90% drop in prevalence after each treatment and $R_0 = 3$ for both the directly transmitted and the ETD (see the electronic supplementary material, Appendix S1 for further parameter values). In (c), only the dynamics of an ETD are displayed, with model parameters set as in (a) and two interventions considered: only drug treatment administered once a year (solid line), and the same drug administration complemented with an environmental intervention (e.g. water sanitation, use of insecticide or biological control) that reduces life expectancy of the pathogen environmental stages by one-third (dashed line).

growth theory, to the infectious disease model [34,35]:

$$\frac{dk}{dt} = r(I)y(I,k) - dk, \quad (2.3)$$

where k is *per capita* physical or human capital; d is capital depreciation rate; r is capital accumulation or savings rate, which is negatively affected by disease I , and $y = y(I,k)$ is *per capita* income, an increasing function of capital and a decreasing function of disease prevalence in the population.

In coupled disease–economic systems [11,36,37], an increase in prevalence of either directly transmitted (electronic supplementary material, Appendix S1; $\varepsilon = 0$) or ETD (figure 5a–c; $\varepsilon = 1$) can trigger a sequence of cascading effects that reinforce conditions of poverty. Lower capital in turn fosters disease transmission (high β) and reduces recovery (low γ), ultimately creating a negative reinforcing cycle of poverty and disease. A substantial reduction in disease prevalence can trigger the opposite sequence of cascading effects, leading to a reinforcing cycle of better health, increased labour productivity and capital accumulation. Therefore, disease–economic systems can, under some circumstances, be characterized by bistability [11,36,37], where one locally stable equilibrium is associated with low income and high disease

prevalence (i.e. a ‘poverty trap’) and the other equilibrium is characterized by high income and little or no disease (figure 5). The equilibrium outcome of either a poverty trap or a virtuous cycle towards health and productivity, is contingent on the history of the system and on specific ‘shocks’ in economic and/or healthcare investments that might push the disease–economic system from one basin of attraction to the other [36]. In the case of ETDs, such as most NTDs, disease–economic dynamics are highly influenced by the dynamics of the pathogen in the environment (figure 5a–c). As a result, the same strategies that could help break a disease-driven poverty trap for directly transmitted diseases, might not be sufficient for ETDs (figure 5d–f). Indeed, failure to act on the environmental reservoir, despite administration of periodic drug treatment, may prevent a sustained prevalence reduction under the critical threshold that switches the basin of attraction for ETDs, therefore perpetuating the poverty trap (figure 5d–f).

These models, which have not previously incorporated an environmental component of disease transmission, have been extended to include within-human population dynamics, where bistability can occur on coupled epidemiological/economic networks [34]. Even if diseases have low prevalence, and no bistability, disease–economic feedbacks can lead to

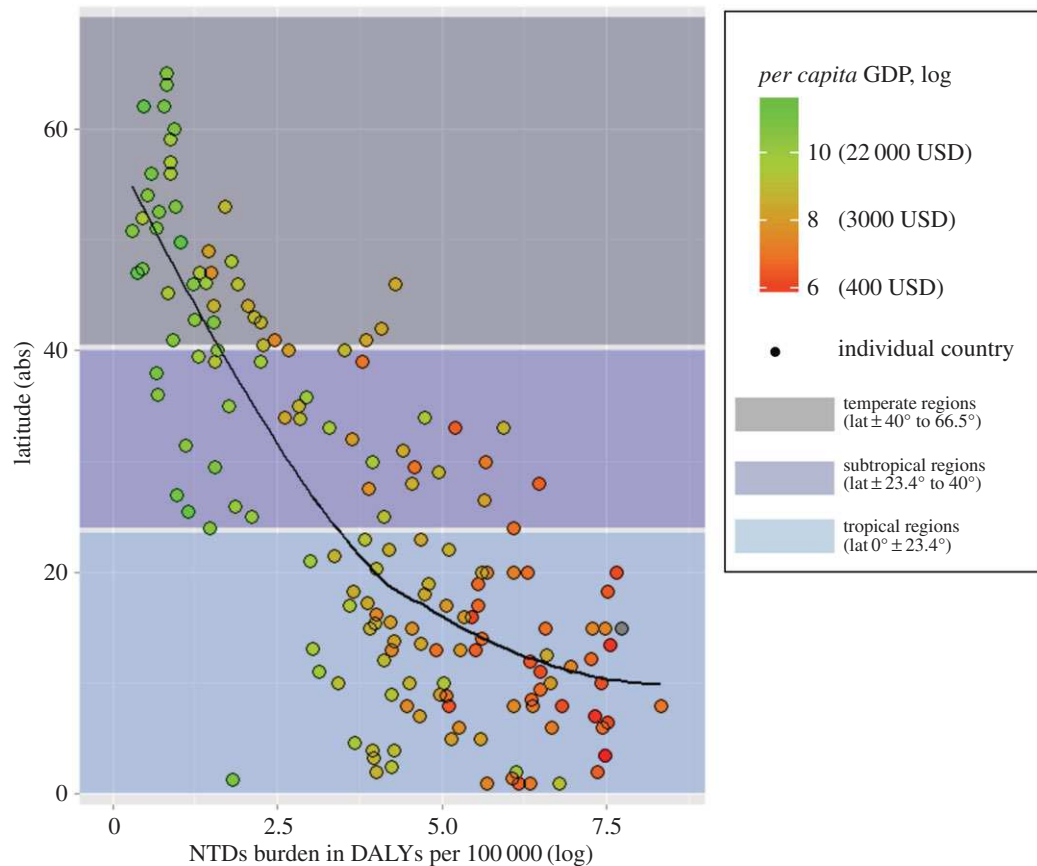


Figure 4. Latitudinal gradient in NTDs burden and poverty. Tropical and subtropical regions (blue and light blue shade) bear an overwhelming burden of NTDs, represented in the x-axis as DALY per 100 000 population (logarithmic scale). These regions accumulate 88.0% and 11.8% of the global number of NTD-related DALYs, respectively, which together represents more than 99% of the world burden. Concurrently, tropical and subtropical regions present some of the lowest levels of *per capita* GDP in the world (dot colour gradient from red for lowest GDP to green for highest GDP). The black solid line in the graph shows the relationship between latitude and NTD-related DALYs (loess nonlinear smoother). Figure was created with R statistical software (package ggplot2) with country data obtained from [23–25].

inequalities in cases of NTDs that cause lifelong disability [38]. Equations ((2.1)–(2.3)) therefore represent a simple modelling framework capable of accounting for a continuum of alternative infectious disease transmission pathways and socio-economic drivers that can be extended to account for system-specific contexts. To illustrate the concepts introduced in our framework for specific systems, we present in §§3 and 4, as case studies, recent evidence on socio-economic impacts, environmental drivers and ecological solutions for two NTDs with environmental transmission ($\varepsilon = 1$) at opposite extremes of the local-to-exogenous transmission (φ) axis in figure 2*a*. Buruli ulcer is a debilitating disease caused by a waterborne mycobacterium whose transmission to people depends on exogenous environmental reservoirs ($\varphi = 1$). Schistosomiasis is caused by a blood fluke, whose life cycle depends strongly on a transmission loop from infected people to the local aquatic environment (snails) and back to people ($\varphi = 0$ or close to zero).

3. Case study 1. Buruli ulcer: the need for ecological and socio-economic insights for disease control

Buruli ulcer, a devastating skin disease caused by the environmental pathogen *Mycobacterium ulcerans*, is probably one of the least-studied of all recognized NTDs. Discovered in the 1960s, it has rapidly emerged in tropical and subtropical countries and currently infects 5000 new people every year

[39]. Most of the disease burden concentrates in rural areas of central and western Africa, where it presents a highly focal distribution, even within endemic regions [39]. Although initial stages of the disease (i.e. nodules, plaques and small ulcers) can be treated with an eight-week antibiotic regimen, poor access to healthcare in these areas frequently leads to catastrophic disease progression [40,41]. One in four cases experience functional limitations for life as a consequence of complications associated with the later stages of the disease (i.e. extensive ulcers, osteomyelitis) that require invasive surgery and lengthy hospital stays [40,41]. These pathological problems have socio-economic causes and consequences. Besides the enormous cost of the disease episode itself, estimated at 25% of household yearly income in Cameroon [42], handicap and deformity are associated with significant stigma, social isolation, job loss and school dropout [41,42]. Early detection and treatment of cases can avert much of the unnecessary suffering [3]. Yet, for ETDs like Buruli ulcer whose dynamics are driven by extrinsic environmental factors ($\varepsilon \rightarrow 1$, $\varphi \rightarrow 1$, figure 2), our poor understanding of the environmental mechanisms that drive pathogen ecological dynamics and its transmission to human populations undermines our capacity to prevent the disease.

Buruli ulcer is associated with aquatic ecosystems, and *M. ulcerans* is found in water, mud, aquatic plants and freshwater insects from representative taxa of the whole aquatic community [43]. The ubiquity of this multi-host environmentally persistent pathogen is exacerbated by the lack of safe water in endemic rural areas, where daily activities, such as bathing or washing, repeatedly expose people to infection

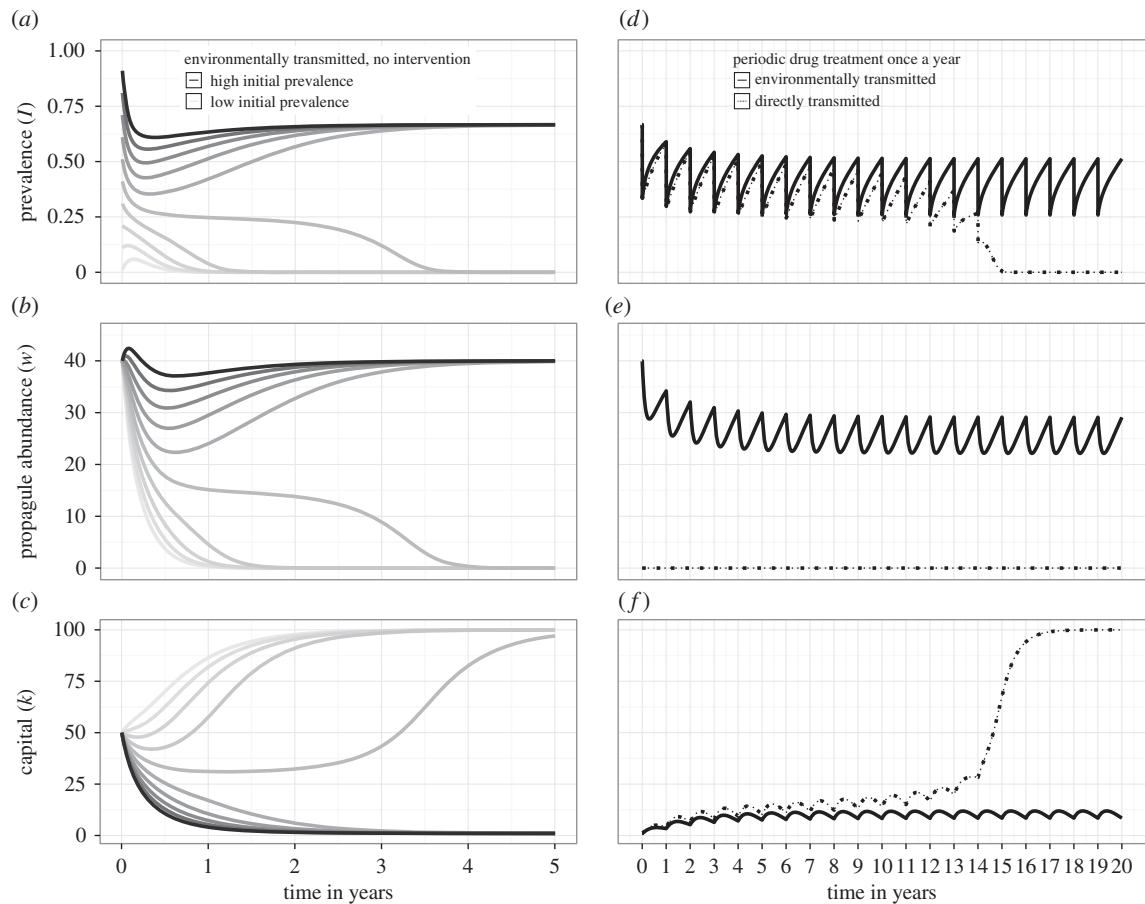


Figure 5. Dynamics of coupled economic–epidemiological systems and impact of periodic drug treatment for directly and environmentally transmitted diseases. Graphs are based on the coupled system described in the main text, and parameter values described in the electronic supplementary material, Appendix S1. (a–c) Dynamics over 5 years in a coupled system for the case of an ETD with $R_0 = 3$. Graphs display dynamics of human prevalence (I), pathogen environmental abundance (W), and *per capita* capital (k), starting from a range of initial conditions for prevalence I , with same k and W . Shades are increasingly dark for increasing values of initial prevalence. Bistability emerges from this system, with two equilibria depending on initial conditions: a high prevalence low capital equilibrium, and a low prevalence high capital equilibrium. (d–f) Impact of human drug treatment dispensed once a year on the dynamics over 20 years for a directly transmitted disease (dotted lines) and an ETD (solid lines) in a coupled economic–epidemiological system. Initial conditions were set as in the high-prevalence–low-capital equilibrium in the left panel, to assess the effectiveness of this intervention to break a theoretical disease-driven poverty trap. $R_0 = 3$ for both types of disease.

[44,45]. Prevalence of the disease is concentrated near swamps, flooded areas and along the basin of slow-flowing rivers at low elevation. Land-use change due to agricultural practices, deforestation, or the construction of dams and roads have been correlated with the emergence of Buruli ulcer endemic areas [43]. Recent advances in disease ecology are shedding light on the conditions that favour environmental persistence and growth of *M. ulcerans*, and drive its spatio-temporal dynamics (W in equation (2.1b)). *M. ulcerans* thrives in periods of intensive rainfall, and certain physicochemical conditions of lentic ecosystems (i.e. low water flow, low oxygen, mildly acidic pH, high temperature) may allow persistence as free-living stages [46,47] (figure 6a). Under unfavourable water conditions, trophic transmission between aquatic hosts seems to ensure the presence of *M. ulcerans* in the environment, independently of the infections in people, which may explain the ubiquity of the pathogen in endemic regions [46,47].

An ecological perspective can also provide new insights on Buruli ulcer transmission. It is unclear whether direct entry into the skin [43] or vector transmission through the bites of water bugs [49] are major contributors to transmission. Ecological studies recently provided the first field-based evidence that environmental transmission to humans could play a greater role than vector transmission (figure 6b) and suggested that

M. ulcerans' environmental load could predict the temporal and spatial patterns of Buruli ulcer incidence [48]. Integration of this ecological understanding with known socio-economic features of the disease in coupled economic–epidemiological models (as shown in equations (2.1)–(2.3)) is helping to clarify the economic consequences of land-use change through their effects on Buruli ulcer [38]. These models show that in contexts of high environmental risk, Buruli ulcer can cause economic inequalities at the population level (figure 6c), with disproportionate impacts on the poorest socio-economic groups due to disparities in vulnerability and healthcare access [38]. Thus, the negative consequences of land-use change could fall disproportionately on the poor. These studies represent an initial step towards a more comprehensive understanding of Buruli ulcer environmental risk, transmission and economics that could lead to the implementation of novel control strategies.

4. Case study 2. Schistosomiasis: an ecological solution for disease control and economic development

Schistosomiasis is a debilitating disease caused by freshwater trematodes of the genus *Schistosoma* and one of the most

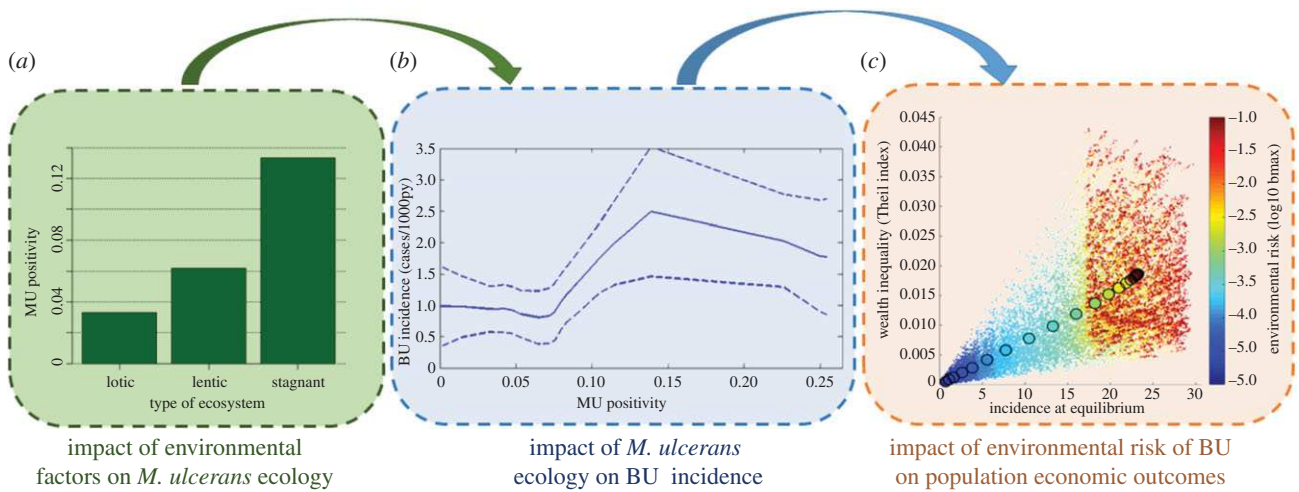


Figure 6. Understanding feedbacks between environment, Buruli ulcer and economic development. A recent research initiative in Cameroon sought to characterize the environmental dynamics of *M. ulcerans* through frequent sampling of multiple aquatic ecosystems and molecular analyses, in order to gain new insights on the ecology and epidemiology of the disease. (a) The prevalence of the bacteria in aquatic sites (MU positivity) appeared to be favoured by specific conditions in stagnant and slow-flowing (lentic) ecosystems [46]. (b) Comparisons of *M. ulcerans* ecological dynamics with patterns of Buruli ulcer incidence from the same areas in mathematical and statistical models, showed that spatial and temporal patterns of Buruli ulcer (y-axis) were influenced by the environmental prevalence of *M. ulcerans* (x-axis) [48]. (c) Individual-based model simulations of a coupled economic–epidemiological system adapted to Buruli ulcer revealed that the disease can be responsible of economic inequalities in endemic populations, with increases in Theil index (y-axis) as a function of BU incidence (x-axis) and *M. ulcerans* environmental load (colour gradient) [38]. Figures are adapted from Garchitorena *et al.* [38,46,48].

important NTDs based on number of people at risk (almost 800 million) and number infected (approximately 250 million) [3,4,50,51]. More than 90% of infections occur in sub-Saharan Africa, the vast majority of which occur in school-aged children between 5 and 15 years old [5]. Infection occurs through exposure to natural freshwater ecosystems in tropical areas that are inhabited by obligate intermediate hosts of the parasites—snails of the genus *Bulinus* and *Biomphalaria*. The snails become infected when people (or, in some case other vertebrate hosts such as rodents or cattle) shed parasite eggs into the environment through urine or faeces ($\varepsilon \rightarrow 1$, $\varphi \rightarrow 0$, figure 2). People (or other vertebrate hosts) are infected when they come into contact with waters containing infectious larvae shed from snails. As with Buruli ulcer, exposure commonly occurs through bathing, washing dishes, fishing or other water-based daily activities. Infection with *Schistosoma* spp. can trigger acute and chronic inflammatory processes such as haematuria, scarring, calcification, squamous cell carcinoma and occasional embolic egg granulomas in brain or spinal cord (*S. japonicum*) as well as fever, hepatic egg granulomas, fibrosis and portal hypertension (*S. mansoni*) [52]. Inflammatory processes lead to debilitation, stunted growth and cognitive impairment in children, increased chances of acquiring HIV in the case of urogenital schistosomiasis in women, and may eventually lead to liver, lung or kidney failure. The drug of choice for schistosomiasis, praziquantel, is cheap (US \$0.32 to treat a child [53]) and effective in removing parasites from infected people [52,53]. Yet, despite more than US \$1.4 billion spent in the past two decades on MDA programmes in developing countries, prevalence and intensity of infection have changed little compared with the pre-praziquantel era [54]. This is because, after treatment, rural villagers have few alternatives but to return to the parasite-contaminated waters (figure 3), and parasitic loads in people return to pre-treatment levels in as little as six months [55,56].

In Africa, the development of irrigation systems and the construction of dams often lead to a further expansion of the habitat of snails (an increase in parameter V in

equation (2.1b)), and hence new potential transmission sites for schistosomiasis [57,58]. The construction of the Diama Dam, built at the mouth of the Senegal River in 1986, is a textbook case. Urinary schistosomiasis was endemic at very low levels before the construction of the dam and intestinal schistosomiasis was absent. The first new cases of both intestinal and urinary schistosomiasis were reported about 1 year after the completion of the Diama Dam [59]. Following two decades of exponential growth in the number of infected people, the lower basin of the Senegal River has now become a hyper-endemic region for schistosomiasis [59]. The construction of the Diama Dam not only led to an expansion of good snail habitat—i.e. a year-around stable water body at low salinity—but also decimated populations of the migratory prawn, *Macrobrachium vollohovenii*, a voracious predator of snails [58,60,61]. The combination of increased habitat for the snail population with the removal of a very effective predator provided ideal environmental conditions for *Bulinus* and *Biomphalaria* snails to thrive and amplify schistosomiasis transmission.

A pilot study carried out in 2011–2014 provided supportive evidence that the reintroduction of the freshwater prawn can reduce infected snails and significantly lower reinfection rates in the human population [56]. Computer simulations showed that the combination of MDA and prawn reintroduction can achieve results that neither of these approaches can attain independently [56]. As prawns are a delicious food commodity with market value, the development of prawn production through aquaculture could also alleviate hunger and provide a source of revenue for local villagers, so that prawn restoration could become an actionable ecological solution leading to a mutually beneficial outcome for biodiversity, schistosomiasis reduction, food production and income [56]. An extended analysis of the geographical distribution of dams and 24 marketable prawn species around the world in areas where schistosomiasis is endemic [58] shows that the removal of natural predators of schistosomiasis-bearing snails might have occurred in many other places in addition

to Senegal. The reintroduction of these predators, either through ecological restoration or aquaculture, might protect people from the risk of contracting schistosomiasis and could complement drug-distribution programmes [58].

5. Discussion and Conclusion

While poverty, infectious disease, and environmental change have been the subject of much independent investigation, they are interlinked in complex ways that require integrated study in order to meet key targets in the Sustainable Development Goals [9]. Poverty and weak health systems arguably pose the most important risk factors for acquiring and succumbing to infectious diseases. Less recognized is the fact that pathogens and parasites responsible for most tropical diseases of poverty are predominantly transmitted through environmental pathways and highly influenced by ecological factors. Here, we describe a simple framework for studying the complex relationships between poverty, infectious diseases, and the environment. The aim of this framework is to help investigate environmental drivers of joint poverty–disease dynamics, with the ultimate goal of identifying sustainable solutions for human and planetary health. Preliminary analyses illustrate the limited effectiveness of human drug treatment for the control of ETDs, when compared with directly transmitted diseases. Although the model cannot be realistically parametrized for specific diseases in its simplest form, this key conceptual difference highlights opportunities for environmental interventions for disease control. By integrating ecological dynamics into coupled disease–economic systems, we show that the presence of persistent environmental reservoirs of pathogens may affect both NTD prevalence and poverty. Future research is needed to investigate how interrupting environmental transmission cycles [3,7] can complement current efforts to expand preventive and curative services for the poorest groups [62,63] to break cycles of poverty and disease.

Building on current ecological knowledge, cost-effective interventions for disease prevention can be designed for many infectious diseases of poverty [64–66]. Mosquito control strategies are among the most widely recognized and extensively deployed of these interventions [67]. In the absence of an effective vaccine for malaria and caution against drug resistance, the recent reduction in disease burden in sub-Saharan Africa has been largely driven by the massive scale-up of indoor residual spraying and long-lasting insecticidal nets [67]. Other examples can be found for zoonotic diseases such as Nipah virus, where the use of bamboo skirts provides a physical barrier that prevents bats (the natural reservoir of NiV) from accessing date palm sap, subsequently reducing transmission to human populations [68,69]. The integration of drug administration with basic infrastructure investments in water, sanitation and hygiene (WASH) has been recognized as a priority in the prevention and care for multiple NTDs, along with improved living conditions, integrated vector control, health education and stronger health systems in endemic areas [3,70]. Yet, the immediate investment costs of infrastructure development hinder the implementation of WASH and similar interventions, leaving drug treatment as the main control strategy for most NTDs. For sapronoses, whose presence in the environment is independent of human infections, such as fungal and waterborne bacterial infections (e.g. Buruli ulcer),

more research on environmental drivers will help identify transmission hot spots, contribute to early detection of cases, and inform prevention strategies [46,48]. For ETDs with well-known ecological drivers, such as trematode and vector-borne infections (e.g. schistosomiasis), control strategies based on natural enemies of pathogens or their wildlife hosts/vectors could represent sustainable ecological solutions. A recent analysis of a century of schistosomiasis control programmes [54] showed that, before the availability of the anti-parasitic drug praziquantel, successful elimination was achieved via interventions that directly targeted the snail intermediate host [54,71]. Supportive evidence [56,58,60] suggests that freshwater prawns may reduce disease transmission through predation on schistosomiasis-bearing snails. More research is needed to identify what other natural enemies of parasites or of their wildlife hosts could be successfully harnessed to achieve a better control of disease transmission [72,73]. While broadly used for integrated pest control in agriculture, such ecological strategies have received little attention in the literature of human infectious diseases. Despite showing promising results in the fight against malaria and other vector-borne diseases, environmental strategies to complement medical approaches still remain poorly developed and under recognized [74].

The many dimensions of the Sustainable Development Goals and an increased attention to the interdependence of human wellbeing and the Earth's provisioning of ecosystem services, have recently expanded the focus of the global health community into the new scientific movement of Planetary Health [9]. While this movement reflects a wide consensus on the threats and challenges currently faced by human populations and natural systems, we still lack actionable solutions that cost-effectively leverage our understanding of the ecology of disease transmission to break the reinforcing cycles of poverty, environmental change and disease. To make the Planetary Health concept actionable, research efforts can help identify specific keystone systems for developing and deploying ecological interventions that influence both poverty and disease.

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