REVIEW

Climate Change and Cascading Risks from Infectious Disease

Jan C. Semenza · Joacim Rocklöv · Kristie L. Ebi

Received: December 21, 2021 / Accepted: April 20, 2022 / Published online: May 19, 2022 @ The Author(s) 2022

ABSTRACT

Climate change is adversely affecting the burden of infectious disease throughout the world, which is a health security threat. Climate-sensitive infectious disease includes vector-borne diseases such as malaria, whose transmission potential is expected to increase because of enhanced climatic suitability for the mosquito vector in Asia, sub-Saharan Africa, and South America. Climatic suitability for the mosquitoes that can carry dengue, Zika, and chikungunya is

Supplementary Information The online version contains supplementary material available at https://doi.org/10.1007/s40121-022-00647-3.

J. C. Semenza (🖂)

Heidelberg Institute of Global Health, University of Heidelberg, 69120 Heidelberg, Germany e-mail: Jan.Semenza@lateralPHC.com

J. Rocklöv

Section of Sustainable Health, Department of Public Health and Clinical Medicine, Umeå University, 901 87 Umeå, Sweden

J. Rocklöv

Heidelberg Institute of Global Health (HIGH), Interdisciplinary Centre for Scientific Computing (IWR), Heidelberg University, Im Neuenheimer Feld 205, 69120 Heidelberg, Germany

K. L. Ebi

Center for Health and the Global Environment (CHanGE), University of Washington, Seattle, WA 98195, USA also likely to increase, facilitating further increases in the geographic range and longer transmission seasons, and raising concern for expansion of these diseases into temperate zones, particularly under higher greenhouse gas emission scenarios. Early spring temperatures in 2018 seem to have contributed to the early onset and extensive West Nile virus outbreak in Europe, a pathogen expected to expand further beyond its current distribution, due to a warming climate. As for tick-borne diseases, climate change is projected to continue to contribute to the spread of Lyme disease and tick-borne encephalitis, particularly in North America and Europe. Schistosomiasis is a waterborne disease and public health concern in Africa, Latin America, the Middle East, and Southeast Asia; climate change is anticipated to change its distribution, with both expansions and contractions expected. Other water-borne diseases that cause diarrheal diseases have declined significantly over the last decades owing to socioeconomic development and public health measures but changes in climate can reverse some of these positive developments. Weather and climate events, population movement, land use changes, urbanization, global trade, and other drivers can catalyze a succession of secondary events that can lead to a range of health impacts, including infectious disease outbreaks. These cascading risk pathways of causally connected events can result in large-scale outbreaks and affect society at large.



We review climatic and other cascading drivers of infectious disease with projections under different climate change scenarios.

Supplementary file1 (MP4 328467 KB)

Keywords: Climate change; Infectious diseases; Cascading risks; Hazard; Vulnerability; Exposure; Malaria; Dengue; Chikungunya; Lyme disease

Key Summary Points

Why carry out this study?

Climate change is considered to be one of the greatest threats to human health in the twenty-first century, with significant increases in temperature extremes, heavy precipitation, and severe droughts.

These climate hazards can activate cascading risk pathways with a sequence of secondary, causally connected events that can disrupt critical infrastructure, vital for a functional society.

What was learned from the study?

This study examines cascading risk pathways from climate change for vector-, water-, food-, and air-borne infectious diseases in a global context.

Cascading effects from climate hazards include also stagnant water that serve as breeding ground for mosquitoes after a flood; contamination of drinking water after a storm surge; breakdown of vector control programs after a hurricane; cholera outbreak after a drought.

A narrow, siloed, and linear assessment of these risks will misinform decision- and policymakers of the magnitude and pattern of future risks, and of the opportunities to modify policies to reduce inherent vulnerabilities and enhance infectious disease control programs.

Elucidating cascading risk pathways from infectious diseases is a first step towards tacking infectious disease threats from climate change.

DIGITAL FEATURES

This article is published with digital features, including a Talking Head video to facilitate understanding of the article. To view digital features for this article go to https://doi.org/10. 6084/m9.figshare.19621077.

INTRODUCTION

Global surface temperature increased at an unprecedented rate over the last 50 years and will continue to do so, until at least the midcentury under all greenhouse emissions scenarios [1]. Climate futures include increases in the frequency and intensity of hot extremes, heavy precipitation, and agricultural and ecological droughts. Pathogens, vectors, hosts, and disease transmission can be sensitive to these changing conditions [2]. Specifically, pathogens develop only within a narrow temperature envelope, with development ceasing at lower or higher temperatures. Temperature influences the reproduction and extrinsic incubation period of pathogens within a vector, with higher temperatures accelerating pathogen maturation. Mosquito biting rate is also a function of temperature, which can affect disease transmission. Moreover, ambient temperature affects the spatial-temporal distribution of disease vectors that carry and transmit pathogen to humans. Disease transmission can in turn be influenced by weather patterns, albeit indirectly, by altered contact rates between human and pathogen, human and vector, or human and host.

However, infectious disease transmission is also a function of underlying vulnerabilities in society (Table 1). Vulnerability is defined as the propensity or predisposition to be adversely affected by infectious disease or a susceptibility to harm and lack of capacity to cope and adapt (Fig. 1). There are multiple vulnerability factors for health impacts of climate change that can be grouped into biomedical (e.g., immunocompromised, malnourished), demographic (e.g., age, sex), geographic (e.g., land use, flood zones), socioeconomic (e.g., civil strife, political Table 1 Combination matrix of climate hazards, vulnerabilities, cascading risks, and climate-sensitive infectious disease (ID) impacts

reeding d populations ulations h h burden h chronic or physical using using s	Cascading risk pathways from extreme precipitation and flooding Standing water serves as breeding ground for mosquitoes Flood water damages critical health care infrastructure Rain runoff from agricultural land and pastures results in turbid water runoff, which impedes water disinfection Contamination of oyster farms from storm runoff that mobilizes and transports pathogens Water treatment efficiency is overwhelmed or damaged by storm Water treatment efficiency is overwhelmed or damaged by storm Water treatment efficiency is overwhelmed or damaged by storm Damage to critical water supply and sanitation infrastructure due to floods Contaminoding risk pathways from drought Crop failure, undernutrition, susceptibility to infectious diseases Concentration of pathogens in drinking water sources Water rollection at alternative (contaminated) sources due to water scarcity Intensified demand and sharing (e.g., with livestock) of limited water resources decreases water availability and quality	Borrelia (Lyme) Dengue virus Tick-borne encephalitis Vibrio spp. Leishmania Campylobacter Campylobacter Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
Ine flood Ine actreeding women Immunocompromised populations women Immunocompromised populations Undernourished populations with high chronic Undernourished populations with high chronic Beople with mental or physical disabilities is a vater or exister or a fee Topollation movement the flood disabilities of the origination of the construction of the flood disabilities of the origination of the construction of the origination of the construction of the flood disabilities of the origination of the construction of the construc	vater serves as breeding ground for mosquitoes er damages critical health care infrastructure of from agricultural land and pastures results in turbid water runoff, which impedes water n ation of oyster farms from storm runoff that mobilizes and transports pathogens atment efficiency is overwhelmed or damaged by storm er and floods overwhelmed or damaged by storm er and floods overwhelme containment systems and discharge untreated waste water of inplacement and inadequate sanitation infrastructure due to floods or fictal water supply and sanitation infrastructure due to floods or fictal water supply and sanitation infrastructure due to floods or fictal water supply and sanitation infrastructure due to floods or fictal water supply and sanitation infrastructure due to floods for of pathways from drought e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water resources decreases water and quality.	Dengue virus Tick-bome encephalitis Vibrio spp. Leishmania Campylobacter Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salimonella Shigella
lood (riverine (riverine) (rive	er damages critical health care infrastructure from agricultural land and pastures results in turbid water runoff, which impedes water from agricultural land and pastures results in turbid water runoff, which impedes water ation of oyster farms from storm runoff that mobilizes and transports pathogens attement efficiency is overwhelmed or damaged by storm er and floods overwhelm containment systems and discharge untreated waste water or and floods overwhelm containment systems and discharge untreated waste water are and floods overwhelm containment systems and discharge untreated waste water or critical water supply and sanitation infrastructure due to floods of pathways from drought e. undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water resources decreases water and quality.	Tick-bome encephalitis Vibrio spp. Leishmania Campylobacter Chikungunya virus Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
(riverine Undernourished populations surge/fides surge/fides Populations with high chronic disease burden Populations with high chronic disease burden Populations with high chronic disease burden e water People with mental or physical is Demographic is abilities Sex e Population movement it flood Sex e Population movement it Geographic it Coastal stores ave Flood risk zones Mater-stressed zones Vater-stressed zones ig rain Food-insecure zones	f from agricultural land and pastures results in turbid water runoff, which impedes water n attion of oyster farms from storm runoff that mobilizes and transports pathogens attion of oyster farms from storm runoff that mobilizes and transports pathogens attion of oyster farms from storm runoff that mobilizes and transports pathogens attion of solverwhelmed or damaged by storm er and floods overwhelm containment systems and discharge untreated waste water or critical water supply and sanitation infrastructure due to inundation of isplacement and inadequate sanitation infrastructure due to floods g risk pathways from drought e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water scarcity demand and sharing (e.g., with livestock) of limited water resources decreases water and quality.	Vibrio spp. Leishmania Campylobacter Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salimonella Shigella
Burge/tides surge/tides nelt flood nelt flood Populations with high CD burden nelt flood Passes burden Pepolations with high CD burden Pepolations with high chronic disabilities ake disabilities disabilities disabilities Demographic disabilities over Sex Sex Geographic unplaned urban housing ave Flood risk zones Drought risk zones boar frost) Nater-stressed zones Water-stressed zones food-insecure zones	ation of oyster farms from storm runoff that mobilizes and transports pathogens atment efficiency is overwhelmed or damaged by storm er and floods overwhelm containment systems and discharge untreated waste water oritical water supply and sanitation infrastructure due to inundation displacement and inadequate sanitation infrastructure due to floods grisk pathways from drought e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources dection at alternative (contaminated) sources due to water resources decreases water and quality and quality ettions of water lines with sewer lines due to water shortages results in water contamination	Leishmania Campylobacter Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
od disease burden People with mental or physical displayint mental or physical displayint movement Age Sex Population movement Population movement Flood risk zones Coastal storms and cyclone risk cones Food-insecure zones Food-insecure zones	stment efficiency is overwhelmed or damaged by storm er and floods overwhelm containment systems and discharge untreated waste water o critical water supply and sanitation infrastructure due to floods of tibplacement and inadequate sanitation infrastructure due to floods ng risk pathways from drought e. undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources tion of pathogens in drinking water sources tion at alternative (contaminated) sources due to water resources decreases water and quality and quality	Campylobacter Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
d or Age Sex People with mental or physical disabilities Demographic Age Sex Population movement Geographic Unplanned urban housing Flood risk zones Drought risk zones Conest Sex Vater-stressed zones Flood-insecure zones Food-insecure zones	er and floods overwhelm containment systems and discharge untreated waste water or critical water supply and sanitation infrastructure due to inundation displacement and inadequate sanitation infrastructure due to floods g risk pathways from drought e. undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water scarcity demand and sharing (e.g., with livestock) of limited water resources decreases water and quality nections of water lines with sewer lines due to water shortages results in water contamination	Chikungunya virus Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
d or Age Sex Age Ceographic Fopulation movement Foundation movement Flood risk zones Coastal storms and cyclone risk zones Flood-insecure zones Food-insecure zones Fo	o critical water supply and sanitation infrastructure due to inundation displacement and inadequate sanitation infrastructure due to floods ag risk pathways from drought e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water resources decreases water demand and sharing (e.g., with livestock) of limited water resources decreases water and quality	Cryptosporidium Giardia Hantavirus Rift Valley Fever Salmonella Shigella
de Age or Sex Sex Population movement Population movement Innale urban housing Unplaned urban housing Innale urban housing Flood risk zones Coastal storms and cyclone risk cones Zones Flood-insecure zones Mater-stressed zones	n displacement and inadequate sanitation infrastructure due to floods rg risk pathways from drought e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources tion at alternative (contaminated) sources due to water scarcity to demand and sharing (e.g., with livestock) of limited water resources decreases water and quality and quality ections of water lines with sewer lines due to water shortages results in water contamination	Giardia Hantavirus Rift Valley Fever Salmonella Shigella
or Population movement Geographic Unplaned urban housing Flood risk zones Drought risk zones Castal storms and cyclone risk zones Flood-insecure zones Flood-insecure zones	In the set water from drought and the set of	Hantavirus Rift Valley Fever Salmonella Shigella
Population movement Geographic Unplanned urban housing Unplanned urban housing Flood risk zones Drought risk zones Coastal storms and cyclone risk zones Mater-stressed zones Food-insecure zones	e, undernutrition, susceptibility to infectious diseases tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water scarcity I demand and sharing (e.g., with livestock) of limited water resources decreases water <i>i</i> and quality nections of water lines with sewer lines due to water shortages results in water contamination	Rift Valley Fever Salmonella Shigella
Geographic Unplanned urban housing Unplanned urban housing Flood risk zones Drought risk zones Coastal storms and cyclone risk zones Varter-stressed zones Food-insecure zones	tion of pathogens in drinking water sources ection at alternative (contaminated) sources due to water scarcity I demand and sharing (e.g., with livestock) of limited water resources decreases water and quality	Salmonella Shigella
Unplanned urban housing Flood risk zones Drought risk zones Caastal storms and cyclone risk zones Water-stressed zones Food-insecure zones	ection at alternative (contaminated) sources due to water scarcity I demand and sharing (e.g., with livestock) of limited water resources decreases water r and quality r ections of water lines with sewer lines due to water shortages results in water contamination	Shigella
Flood risk zones Drought risk zones Caastal storms and cyclone risk zones Water-stressed zones Food-insecure zones	I demand and sharing (e.g., with livestock) of limited water resources decreases water r and quality actions of water lines with sewer lines due to water shortages results in water contamination	
Drought risk zones Coastal storms and cyclone risk zones Water-stressed zones Food-insecure zones	nections of water lines with sewer lines due to water shortages results in water contamination	Vero toxin- producing
ost) Coastal storms and cyclone risk 200es Water-stressed zones Food-insecure zones		Escherichia coli West Nile virus
ost) Water-stressed zones Food-insecure zones	Vector breeding in household water containers	Cholera
Food-insecure zones	Poor water quality, hygiene and sanitation (WASH) due to water shortages	Leaionella
	Open defecation due to lack of WASH results in human exposure to pathogens	Rickettsia
Ground frost Socio-economic Low water availa	Low water availability augments travel distance to alternative (contaminated) sources	Hepatatis A
	Cascading risk pathways from elevated temperature	Leptospira
Thaw Gender norms, roles, and relations Longer transmiss	Longer transmission season for vector-borne diseases	Tularemia
e Unsafe, informal occupation	Proliferation of marine bacteria at recreational beaches	Yellow fever virus
Reduced access to health care	Increased prevalence of pathogens in poultry flocks	Yersinosis
Reduced access to education	Extended pathogen survival and replication outside of host	Anthrax
Cho Unsafe water and sanitation	Runoff from areas affected by wildhres during heat waves compromises water quality	Botulism
Gale Inadequate shelter Keduced crop yie Coniare Auronal Conia	Reduced crop yields at higher ambient temperature Fond sonilage due to behavior change (e.g. harbecue)	Listeria
Subtropical storm Political instability Cascading ris Tronical storm Existence of complex emergencies Injundation of dri	Cascading risk pathways from sea-level rise Inundation of drinking water supply and sanitation infrastructure	Loxiella (U Fever) Taxablasma
or conflict		
Tornado Lack of freedom of speech and Saline intrusion i information	Saline intrusion into coastal aquifers and decline in soil and water quality	Schistosoma
I rights and civil ements	Damage of critical infrastructure (e.g., health care, first responders, electricity generation, transportation (e.g., fuel supply, streets, rail, airports, harbors), telecommunication, shelter, fuel extraction(e.g., natural gas, oil), water supply, agriculture, financial services)	Nairovirus (Crimean-Congo haemorrhagic

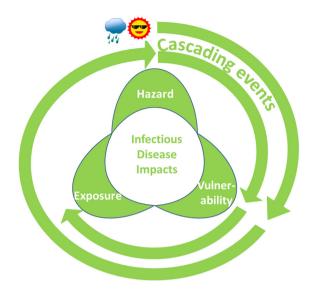


Fig. 1 Cascading risks from infectious disease, due to a nexus of hazard, vulnerability, and exposure. Climatic hazards (e.g., extreme rain event or heat; outer spiral), amplified by societal vulnerabilities can trigger new hazards, such as floodwater contaminated with pathogens or high mosquito densities. Cascading events (inner spiral) caused by these infectious disease hazards and amplified by newly attained vulnerabilities can result in population exposure and give rise to water-borne or mosquito-borne disease outbreaks, respectively

instability) (Table 1) [3]. From an infectious disease standpoint, vulnerability is also determined by the lack of safeguards such as door/ window screens for vectors or flood barriers for storm surges [4].

Furthermore, infectious disease transmission depends on the exposure pattern in human populations, which is defined as the state of people, livelihoods, species, property, (eco-) systems, or other elements present in hazard zones that thereby could be adversely affected. Individuals or communities can be exposed to contaminated drinking water, vectors, or pathogens. The nexus of these three elements hazard, vulnerability, and exposure—determines the current infectious disease impacts or future risks due to climate change (Fig. 1).

In addition to these impacts of weather and climate on pathogens, vectors, hosts, and disease transmission, a weather or climate event can activate cascading risk pathways with a sequence of secondary, causally connected events. Table 1 presents a combination matrix of how climate hazards can be combined with societal vulnerabilities that give rise to cascading risk pathways resulting in climate-sensitive infectious disease outbreaks. The dynamic interactions between climate hazard, exposure, and vulnerability set the stage for cascading risks (Fig. 1; outer spiral). For example, a hurricane can result in flooding or disrupt vector control programs as a result of infrastructure vulnerabilities. Contamination of floodwater with pathogens or high mosquito populations can cause population exposure to pathogens and then trigger cascading events (Fig. 1; inner spiral) such as water-borne or mosquito-borne outbreaks. These resulting impacts of a sequential chain reaction within natural and human systems can be significantly larger than the initial hazard and can cause additional physical, natural, social, or economic disruption (Fig. 1; inner spiral) [5]. If these secondary effects such as flooding disrupt critical infrastructure that is vital for a functional society, the consequences can be substantial. Such events can have a ripple effect across society and generate direct losses through immediate impacts or more secondary losses through consequential impacts. The combination matrix in Table 1 illustrates the predicament of climate change adaptation where virtually any climate hazard can be combined with specific vulnerabilities resulting in cascading risk pathways and infectious disease impacts. Examples of such cascading effects from climate hazards that have been described in the peer-reviewed literature include stagnant water after a flood that serves as a breeding ground for mosquitoes; contamination of drinking water after a storm; injuries from landslides or storm surges with risk of tetanus infections in populations with low vaccination coverage; or a cholera outbreak after a drought (Table 2).

Protecting and promoting health requires understanding and preparing for these causal interdependencies that permeate many aspects of society. A climate hazard can adversely affect one sector in society by surpassing a threshold that causes system failure (Fig. 1). As a result of

Climate hazard	Vulnerability and cascading events	Infectious disease outcome	References
Hurricane	Lack of WASH in mega-shelter after hurricane Katrina	Widespread outbreak of norovirus gastroenteritis among evacuees	Yee EL, Palacio H, Atmar RL, et al. Widespread outbreak of norovirus gastroenteritis among evacuees of Hurricane Katrina residing in a large "megashelter" in Houston, Texas: lessons learned for prevention. Clin Infect Dis. 2007;44:1032–9
Typhoon	Serious flooding in Metro Manila	Outbreak of Leptospirosis	Amilasan A-shereT, Ujiie M, Suzuki M, et al. Outbreak of leptospirosis after flood, the Philippines, 2009. Emerg Infect Dis. 2012;18:91–4
Cyclones: Idai and Kenneth in Mozambique	Lack of access to safe water, poor sanitation, contact with stagnant floodwater, overcrowding in the camps for displaced people	Diarrheal diseases, malaria	Mugabe VA, Gudo ES, Inlamea OF, et al. Natural disasters, population displacement and health emergencies: multiple public health threats in Mozambique. BMJ Global Health. 2021;6:e006778. doi:10.1136/bmjgh-2021- 006778
Heavy rainfall and elevated temperature	Contamination of surface water by compromised WASH systems	Cholera outbreaks in Yemen	Camacho A, Bouhenia M, Alyusfi R, et al. Cholera epidemic in Yemen, 2016–18: an analysis of surveillance data. Lancet Glob Health. 2018 Jun;6(6):e680–e690. https:// doi.org/10.1016/S2214- 109X(18)30230-4
Monsoon: heavy rain	Record-breaking deluge and floods	Acute diarrhea, skin and eye infections, leptospirosis, malaria epidemic, leishmaniasis, respiratory infections, hepatitis	Baqir M, Sobani ZA, Bhamani A, et al. Infectious diseases in the aftermath of monsoon flooding in Pakistan. Asian Pac J Trop Biomed. 2012;2:76–9

Table 2 Selected examples of climate hazards, societal vulnerabilities, and cascading events resulting in infectious disease outbreaks

Climate hazard	Vulnerability and cascading events	Infectious disease outcome	References
Floods	Health care access	Inadequate access to health care after the disaster	Jacquet GA, Kirsch T, Durrani A, Sauer L, Doocy S. Health care access and utilization after the 2010 Pakistan floods. Prehosp Disaster Med. 2016 Oct;31(5):485–91
Floods, storms, droughts	Displacement	Infectious disease outbreaks including measles, cholera, cutaneous leishmaniasis, dengue	Desai AN, Ramatowski JW, Marano N, Madoff LC, Lassmann B. Infectious disease outbreaks among forcibly displaced persons: an analysis of ProMED reports 1996–2016. Confl Health. 2020 Dec;14(1):1–0
Heavy rain	Overwhelmed water treatment and distribution system	Water-borne disease outbreaks, e.g., cryptosporidium	Semenza JC, Nichols G. Cryptosporidiosis surveillance and water-borne outbreaks in Europe. Euro Surveill. 2007 May;12(5):E13–14
Extreme temperatures, droughts	Crop failures, undernutrition	Vulnerability to infectious diseases, especially diarrhea, pneumonia, and measles	Gwela A, Mupere E, Berkley JA, Lancioni C. Undernutrition, host immunity and vulnerability to infection among young children. Pediatr Infect Dis J. 2019 Aug 1;38(8):e175–7

Table 2 continued

Climate hazard	Vulnerability and cascading events	Infectious disease outcome	References
Droughts	Water scarcity, hygiene	Cholera outbreaks; infectious disease outbreaks	Charnley GE, Kelman I, Green N, Hinsley W, Gaythorpe KA, Murray KA. Exploring relationships between drought and epidemic cholera in Africa using generalised linear models. BMC Infect Dis. 2021 Dec;21(1):1–2
			Jofre J, Blanch AR, Lucena F. Water-borne infectious disease outbreaks associated with water scarcity and rainfall events. In: Sabater S, Barceló D, editors. Water scarcity in the Mediterranean; 2009: pp. 147–59. Berlin, Heidelberg: Springer

 Table 2
 continued

the interconnected nature of modern society, this can have implications for other sectors [6]. The potential for cross-scale failures need to be defined in relation to the initial climate trigger. By doing so, the complexity of interactions within the public health network can be modeled mathematically. That way, cascading risk pathways can be simulated and analyzed. Understanding the weaknesses in the system can help advance adaptive capacity and intervention measures. Only then can cascading failures be anticipated and intercepted quickly with targeted measures that can prevent a public health impact. Here we critically review interlinked drivers of infectious disease transmission and elucidate the cascading risks associated with climate variability and change. We thoroughly examine cascading risk pathways from climate change for vector-, water-, food-, and air-borne infectious diseases in a global context, as opposed to a focused assessment of one infectious disease category [7] or one geographic area [4], which, to our knowledge, has not been attempted before. Such а

comprehensive assessment is critical in order to elucidate cascading risk pathways from infectious diseases, in the context of the complex branching configuration of a globalized, dendritic society; indeed, it is a first step towards tackling infectious disease threats from climate change.

METHODS

Peer-reviewed research articles were retrieved from PubMed using the following search terms: infectious diseases, vector-, water-, food-, airborne diseases, climate change, climate variability, global warming, temperature, heat wave, precipitation, flooding. Disease-specific searches by pathogen name were also conducted. Keywords of the concepts and MeSH terms (when available) were used in the search strategies. A special focus was given to English language publications from the last 5 years. Of particular interest were publications that examined the specific aims of this study,

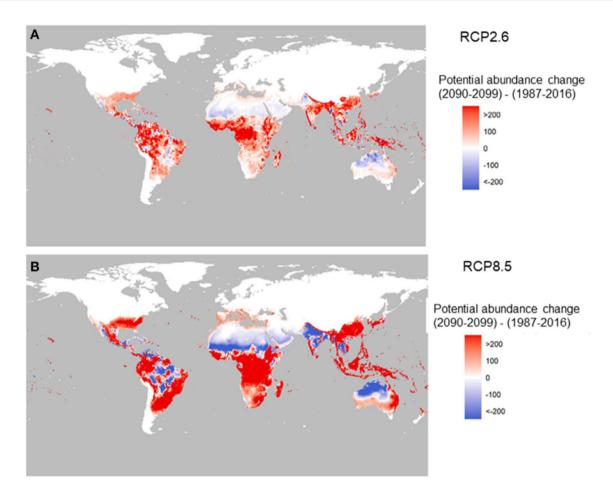


Fig. 2 Change in the potential abundance of *A. aegypti* (per larval site) over the twenty-first century (2090–2099 relative to 1987–2016). The two panels correspond to two

climate change and cascading risks from infectious disease, and assessed the association between climate change and disease transmission. Research reports from international organizations and gray literature were also included in our analysis. This article is based on previously conducted studies and does not contain any new studies with human participants or animals performed by any of the authors.

VECTOR-BORNE DISEASES

Climatic conditions indirectly affect vectorborne diseases such as mosquito- and tick-borne diseases. Alterations in environmental conditions can have secondary effects on vector carbon emission scenarios: RCP2.6 (a) and RCP8.5 (b). Source: Reference [27]

populations, replication rates of pathogens, and vector-host interactions. Further, climatic events can result in cascading secondary effects that can alter the transmission pathway for vector-borne diseases discussed in this section.

Mosquito-Borne Diseases

Malaria

Malaria is caused by five species of plasmodium parasites that are transmitted by *Anopheles* mosquitoes. The disability-adjusted life years (DALYs) rate of malaria declined by almost 40% globally between 2007 and 2017. The largest burden of disease occurs in Africa where more than 90% of all malaria-related deaths occur [8, 9]. The RTS,S/AS01 malaria vaccine for the

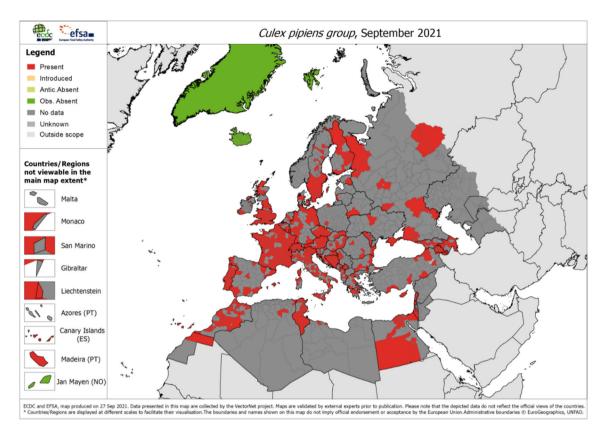


Fig. 3 The vector for West Nile virus: *C. pipiens* mosquito distribution in Europe as of September 2021. Source: ECDC, https://www.ecdc.europa.eu/en/publications-data/culex-pipiens-group-current-known-distribution-september-2021

prevention of *Plasmodium falciparum* malaria in children living in regions with moderate to high transmission in combination with chemoprevention should further decrease the disease burden [10, 11]. While there have been recurrent, local outbreaks of malaria in Europe [12], the risk for widespread transmission is relatively low.

Malaria has expanded its geographic range into higher altitudes during warmer years in the highland areas of Columbia and Ethiopia [13, 14]. Thus, without interventions, it is possible that declining trends in the number of malaria DALYs will be offset by additional climate change. Cascading risk pathways from a number of hurricanes disrupted anti-malaria vector control programs and resulted in a resurgence of *P. falciparum* malaria in Haiti, Guatemala, and Nicaragua in the 1980s and 1990s [15, 16]. In the Amazon region, the dry season is getting longer and the rainy season that used to start at the end of October now starts at the beginning of December; this exacerbates and accelerates the burning of the rainforest. Heat stress and fires in the Amazon rainforest along with deforestation, road density, and selective logging are associated with malaria risk in the Amazon [17–19]. Smoke exposure from the Amazon fires is associated with increased respiratory illnesses such as pneumonia, acute bronchitis, and asthma in indigenous populations [20].

The *Anopheles* vector is projected to expand its geographic range under different climate change scenarios, in part due to an increase in temperature and an expansion of the rainy season in the tropical areas of Africa [21, 22]. Similarly, an expansion is projected in South Africa and China but a contraction in India and Southeast Asia as a result of a reduction in climatic suitability [23, 24]. In China, Plasmodium vivax and P. falciparum malaria distributions are projected to increase under higher emission scenarios, such as representative concentration pathway (RCP) 4.5 and RCP8.5 [25]. Under a multi-scenario climate change framework, these are also the areas projected to experience a lengthening of the transmission season by 1.6 additional months in the tropical highlands in the African region, the Americas, and the Eastern Mediterranean region. The improvement in climatic suitability is expected to be greater in rural areas than in urban areas, and the epidemic belt would expand towards temperate regions [26]. Similarly, a general rise in months suitable for transmission is projected to increase in India; in other areas, the transmission season is projected to contract, when the climatic conditions will be too extreme for the vector species [27].

Arboviral Diseases

Dengue Aedes aegypti, the principal mosquito vector of many arboviral diseases, such as dengue, chikungunya, yellow fever, and Zika, has experienced a global expansion, threatening almost half of the world's population [28, 29]. This expansion is attributed in part to the global temperature increase [30] but also globalized population movement through air traffic and urbanization, and insufficient vector control measures [31, 32]. Dengue is responsible for an estimated 10.000 deaths and 100 million symptomatic infections per year in over 125 countries [33, 34]. Dengue incidence is positively associated with temperature, precipitation, and relative humidity in several settings worldwide, including the Americas [35], India [36], and Philippines [37]. Sea surface temperature, rain, and variation in wind associated with the El Niño Southern Oscillation over the Pacific Ocean has also been used as a predictor of dengue incidence [38]. Cascading risks due to a breakdown of vector control measures in countries of Central America after hurricane Mitch in 1998 resulted in almost 40,000 cases of dengue and dengue hemorrhagic fever [16].

The change in potential for global abundance in the future of the dengue vector *A. aegypti* is shown in Fig. 2, with big increased potential in Southeast Asia, China, Japan, East Australia, and Africa [39]. More favorable temperatures and increased rainfall by 2050 from climate change could increase the suitability for dengue in southern and western Africa, southeastern USA, central Mexico, northern Argentina, and inland areas of Australia. In addition, coastal cities in eastern China and Japan are projected to become more suitable by 2050 [29]. The potential transmission season will lengthen by 4 months because of an increase in climatic suitability, particularly in lowlands in the Western Pacific region and the Eastern Mediterranean region [26].

Chikungunya The chikungunya virus was first identified in Tanzania in 1952, where it caused a localized outbreak in Africa and parts of Asia, and then spread to countries around the Indian Ocean. Travel and trade have contributed to a continuous geographic spread to temperate areas [40]. The virus has also been repeatedly imported into Europe where conducive climatic conditions contributed to two large outbreaks in Italy in 2007 and 2017 [41, 42]. Projecting the chikungunya risk under RCP4.5 and RCP8.5 indicates an expansion of the transmissionsuitable areas in China, sub-Saharan Africa, South America, the USA, and continental Europe, although also some contraction of the transmission risk along parts of the Adriatic coast of Europe as a result of unfavorable climatic conditions for example [43].

Zika The Zika virus has also expanded globally, causing large outbreaks in South America in 2016 following a period of record high temperatures and severe drought conditions in 2015 [44]. Storing drinking water in open containers at home as a result of the drought might have created ideal vector breeding and exposure conditions that contributed to the outbreak. Zika could expand north with longer seasons as temperatures move towards the predicted thermal optimum (29 °C) [45].

West Nile Fever Europe experienced uncharacteristically high spring temperatures in 2018 followed by an exceptionally early and intense West Nile virus (WNV) transmission season,

with 2083 cases [46, 47]. These weather anomalies could have activated the mosquito breeding season early and reduced the extrinsic incubation period, which would explain the high prevalence of WNV in mosquito vectors (Culex pipiens) (Fig. 3) and avian hosts, compared to previous years. In fact, birds infected with WNV were discovered before the virus was detected in humans, in both the Netherlands and Germany [48, 49]. In Europe, progressive expansion of WNV is projected along the edges of the current transmission areas, which can result in a cascading risk to the safety of blood banks [50]. WNV-infected, but asymptomatic donors can inadvertently contaminate the blood supply which can arrest blood transfusion services. In order to prevent cascading risk pathways affecting the blood supply, a number of steps need to be taken, such as screening, deferral and pathogen reduction strategies [51].

TICK-BORNE DISEASES

Lyme Disease

Vector surveillance in Canada has documented a geographic range expansion of the black-legged tick Ixodes scapularis, the main vector of Borrelia burgdorferi, the agent of Lyme disease [52]. This expansion is associated with elevated temperatures, the emergence of tick populations, increases in their range and recent geographic spread, as well as with a rapid increase in human Lyme disease cases [53-56]. In Europe, transmission by Ixodes ricinus ticks is also determined by factors besides temperature, such as host populations and habitats [57]. A higher transmission risk is projected under all RCP scenarios for I. scapularis in some areas of Canada [58]. The season of Lyme disease is projected to expand in the USA under a 2 °C warming scenario with a 20% increase of cases over the coming decades and lead to an earlier onset and longer length of the annual Lyme disease season [58, 59]. In Europe, the risk of transmission could even be reduced under the low-emission greenhouse gas scenario [60].

Tick-Borne Encephalitis

Tick-borne encephalitis (TBE) is an important zoonotic infection, with an increasing disease burden and expanding geographic range across Europe, Russia, and parts of Asia. Among a number of contributing factors, milder winters and warmer springs due to climate change have been implicated in this range expansion [61-63]. TBE is a potentially serious disease, with 3411 TBE cases reported in EU/EEA countries in 2019 of which 20 died (case fatality, 0.7%). TBE cases generally display a seasonal peak in the months of July and August. Conducive climatic conditions can result in a cascading chain of events where goats, sheep, or cows are infected with the TBE virus that can result in alimentary infection of humans after consumption of unpasteurized milk and cheese from domestic ruminants [64-66].

Rocky Mountain Spotted Fever

Rocky Mountain spotted fever is the most common fatal tick-borne disease in the USA, with 21 fatalities between 2003 and 2016, caused by *Rickettsia rickettsia* [67]. Warmer wetter climates have led to an expansion of tick habitat range and distribution [68, 69]. For example, the range expansion of the lone star tick is correlated with an increased incidence of spotted fever group rickettsiosis in the USA [70].

Leishmaniasis

The protozoa Leishmania infantum, the main causative agent of zoonotic visceral leishmaniasis and cutaneous leishmaniasis, are transmitted bv infected Phlebotomine sandflies. Leishmaniasis is found in southern Europe and northern Africa around the Mediterranean as well as parts of Asia [71]. The primary reservoirs of human infections are domestic and stray dogs that have experienced a progressive increase of Leishmania seroprevalence rates at higher latitudes and altitudes, more so than might be expected in Mediterranean countries [72]. There has been a climate change-related expansion of sandflies into more northern

Infect Dis Ther (2022) 11:1371-1390

latitudes and higher altitudes in Italy [73], in the Pyrenees [74], and Germany [75]. The convergence of vector dispersion and the scattering of infected dogs (e.g., through adoption services) can compound this public health issue. The climate in Central Europe is projected to become increasingly more suitable for sandflies in the future, under climate change scenarios [76].

FOOD-BORNE DISEASES

Salmonella is climate sensitive and grows in a narrow temperature envelope with a strong seasonality. An increase in ambient temperature is associated with an upsurge in Salmonella incidence in a number of settings, indicating a direct impact on replication rates [77-79]. The situation for Campylobacter is different, because the pathogen cannot replicate outside of the host. Thus, warm weather conditions may not directly affect Campylobacter replication rates but rather reflect human behavioral issues such as riskier patterns of food production/consumption or other seasonal factors [80-82]. Nevertheless, with an increase in ambient temperature and extreme weather events, the food safety risk is anticipated to increase as a result of risks from existing and emerging food-borne pathogens along the food chain [83–85].

The transmission pathway of food-borne diseases through the food chain is complex and susceptible to several climatic drivers [86]. For example, in 2017, hurricane Irma contaminated many commercial fruit and vegetable fields in Florida with pathogens and parasites, which led the US Food and Drug Administration to warn against the consumption of fresh produce that had been in contact with floodwater [87]. Hurricanes can also disrupt food processing, preparation, transport and spoil foodstuff.

WATER-BORNE DISEASES

Water-borne diseases causing diarrheal diseases have been declining globally since the 1990s, owing to improvements in water, sanitation, and hygiene (WASH), reductions in poverty, and vaccination programs [88]. Despite this, the disease burden is considerable in low- and highincome countries [7] and is usually caused by microbial contamination of the drinking water supply [89–91]. Bacteria, protozoa, viruses, or parasites have been implicated in water-borne outbreaks due to an inability of the water treatment system to clear pathogens from the water supply [89, 91].

Water-borne outbreaks can occur because of climate variability and change followed by secondary events that are causally connected. Such cascading risk pathways can lead to a succession of system failures and damage critical infrastructure. For instance, extreme precipitation can mobilize pathogens from pastures and fields and overwhelm water treatment and distribution systems resulting in drinking water contamination [92]. In fact, empirical studies have documented cascading risks from heavy rain that give rise to water-borne outbreaks [93, 94]. Similarly, cascading risk pathways of floods can contaminate drinking water wells, treatment and distribution systems, and produce waterborne outbreaks [95, 96]. Conversely, water shortages and drought can also give rise to cascading risks and cause diarrheal diseases [97, 98], although this association has been documented inconsistently [99].

Cholera

Natural disasters can trigger a sequence of cascading events that can compromise WASH. For example, poor sanitation after a hurricane can result in cholera outbreaks [100]. Cholera, is an acute diarrheal disease, caused by the bacterium Vibrio cholerae that can result in severe morbidity and mortality; it has been associated with several climatic parameters, in situations with poor WASH and where cholera has already been seeded in the population [101–103]. For example, elevated ambient temperature is a key parameter for cholera incidence [104, 105], as well as lower and higher precipitation [106, 107]. The projected risk for non-cholera Vibrio cases, including gastroenteritis, wound infections, and septicemia, is projected to increase in the Baltic Sea region with higher sea surface temperatures [108].

Leptospirosis

Leptospira bacteria can contaminate water, soil, or food through the urine of infected animals, and cause leptospirosis, a bacterial disease that affects humans and animals. For example, recreational water use during hot weather increases the risk of exposure to water contaminated with Leptospira [109]. As part of a time series from 2006 to 2016, human leptospirosis notification was significantly associated with rainfall and land surface temperature in highrisk counties in China [110]. Floodwater or drinking water contaminated with Leptospira can cause outbreaks of leptospirosis [111, 112], vividly illustrating the nexus of climate hazard, societal vulnerability, and population exposure in creating cascading risk pathways for leptospirosis (Fig. 1).

Schistosomiasis

Infections with parasitic blood flukes of the species Schistosoma mansoni, S. japonicum, S. intercalatum, S. mekongi, and S. guineensis can cause intestinal schistosomiasis, associated with systemic inflammation. Cases have been reported in Africa, Latin America, the Middle East, and Southeast Asia [113–115]. Temperature is a determinant of the geographic distribution of flukes and a climate change assessment indicated increased transmission as well as potential shrinkages in certain areas [116]. For example, a decade-long drought between 2001 and 2009 resulted in the disappearance of significant clustering around historical transmission hot spots in coastal Kenya, due to the disappearance of the flukes from ponds along with urinary schistosomiasis transmission [117]. Conversely, extreme precipitation in June 2017 in Brazil resulted in a large schistosomiasis outbreak after a large flooding period [118].

Hotter areas in Africa are projected to experience reduced snail populations but higher populations are projected in areas with currently lower winter temperature [119, 120]. In nic areas in Sichu

1383

China, currently endemic areas in Sichuan Province are projected to contract, but non-endemic areas in Sichuan and Hunan/Hubei provinces expand [116, 121].

Respiratory Infections

Respiratory infections tend to be highly seasonal, with higher incidence in the winter months, in part due to increased pathogen survival, indoor crowding, and elevated host susceptibility [122]. For example, temperature and humidity determine the incidence of influenza in temperate regions of the world. Low daily temperatures and both low and high relative humidity were associated with an increased risk of influenza incidence in Seoul, Republic of Korea [123]. Similarly, low temperature correlated with peaks of influenza virus activity in Northern Europe [124]. Conversely, warm winters tend to be followed by severe and early-onset influenza incidence the following season [125], partially due to waning population immunity to previous infections.

A respiratory disease outbreak caused by hantavirus in 1999 and 2000 in Santos, Panama was preceded by extreme precipitation [126]. Such unusual rainfall patterns may have led to increases in rodent populations and contact rates with humans [127, 128]. Infected rodents can harbor the hantavirus in their saliva, urine, or feces which can result in human exposure. Another aspect of cascading effects in a warming world is the increased risk for flooding, which can result in respiratory infections; the incidence of respiratory infections has been observed to increase after flooding in a number of settings [129–131].

CONCLUSION

Increasing climate variability is already leading to cascading risks from infectious disease. With projections of increases in multiple modes of climate variability with additional climate change, it is vital for health systems to prepare for more and more extreme cascading risks, taking into account multiple other drivers of outbreaks of infectious diseases. The preparation needs to consider that these changes will vary over time and space, so are inherently difficult to predict. That means health systems need to prepare for uncertainty as much as for climate change. This requires flexibility in planning modifications to vector control programs to ensure they are prepared for a range of possible futures.

Climate effects can have far-reaching implications for public health through inherent societal vulnerabilities that can magnify the impacts of cascading risk pathways (Fig. 1). A narrow, siloed, and linear assessment of these risks will misinform decision- and policymakers of the magnitude and pattern of future risks, and of the opportunities to modify policies to enhance infectious disease control programs, building from current programs. For example, current malaria control programs include treatment, bed nets, and vector control; also incorporating land use (e.g., wetland management, drainage of standing water) and socioedeterminants conomic (e.g., housing, occupational exposure) can indirectly counteract the impacts of increasing climate variability on disease transmission.

A comprehensive understanding is critical of the interconnected nature of public health with social, demographic, and environmental drivers of infectious diseases. To this end, a better collaboration is warranted between public health practitioners, climate scientists, civil engineers, social scientists, network modelers, and decision-makers. Mathematical modelling of climate hazards, vulnerabilities, and exposures can improve projections of cascading events and facilitate transformative adaptation.

Failure to invest in research and health systems would jeopardize the resilience of individuals and communities to cascading impacts from infectious disease, leading to a sicker future.

ACKNOWLEDGEMENTS

Funding. No funding or sponsorship was received for this study or publication of this article.

Authorship. All named authors meet the International Committee of Medical Journal Editors (ICMJE) criteria for authorship for this article, take responsibility for the integrity of the work as a whole, and have given their approval for this version to be published.

Author Contributions. JCS had the idea for the article, performed the literature search and data analysis, and drafted the article. All authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

Disclosures. Jan C. Semenza, Joacim Rocklöv, and Kris Ebi all have nothing to disclose.

Compliance with Ethics Guidelines. This article is based on previously conducted studies and does not contain any new studies with human participants or animals performed by any of the authors.

Open Access. This article is licensed under a Creative Commons Attribution-NonCommercial 4.0 International License, which permits any non-commercial use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/bync/4.0/.

REFERENCES

- 1. The Intergovernmental Panel on Climate Change (IPCC). Sixth assessment report; climate change 2021: the physical science basis. https://www.ipcc. ch/report/ar6/wg1/#SPM. Accessed 17 Apr 2022.
- Wu X, Lu Y, Zhou S, Chen L, Xu B. Impact of climate change on human infectious diseases: empirical evidence and human adaptation. Environ Int. 2016;1(86):14–23.
- 3. The impacts of climate change on human health in the United States: a scientific assessment. https:// health2016.globalchange.gov/. Accessed 1 Dec 2021.
- 4. Semenza JC, Paz S. Climate change and infectious disease in Europe: impact, projection and adaptation. Lancet Reg Health Eur. 2021;1(9):100230.
- 5. Intergovernmental Panel on Climate Change (IPCC). https://www.ipcc.ch/report/sixthassessment-report-working-group-ii/. Accessed 17 Apr 2022.
- 6. Lawrence J, Blackett P, Cradock-Henry NA. Cascading climate change impacts and implications. Clim Risk Manag. 2020;1(29): 100234.
- 7. Semenza JC. Cascading risks of waterborne diseases from climate change. Nat Immunol. 2020;21(5): 484–7.
- 8. M'Bra RK, Kone B, Soro DP, et al. Impact of climate variability on the transmission risk of malaria in northern Côte d'Ivoire. PLoS ONE. 2018;13(6): e0182304.
- 9. Caminade C, McIntyre KM, Jones AE. Impact of recent and future climate change on vector-borne diseases. Ann N Y Acad Sci. 2019;1436(1):157.
- Bejon P, Lusingu J, Olotu A, et al. Efficacy of RTS, S/AS01E vaccine against malaria in children 5 to 17 months of age. N Engl J Med. 2008;359(24): 2521–32.
- 11. Chandramohan D, Zongo I, Sagara I, et al. Seasonal malaria vaccination with or without seasonal malaria chemoprevention. N Engl J Med. 2021;385(11):1005–17.
- 12. Sudre B, Rossi M, Van Bortel W, et al. Mapping environmental suitability for malaria transmission, Greece. Emerg Infect Dis. 2013;19(5):784.
- 13. Siraj AS, Santos-Vega M, Bouma MJ, Yadeta D, Carrascal DR, Pascual M. Altitudinal changes in malaria incidence in highlands of Ethiopia and Colombia. Science. 2014;343(6175):1154–8.

- 14. Midekisa A, Beyene B, Mihretie A, Bayabil E, Wimberly MC. Seasonal associations of climatic drivers and malaria in the highlands of Ethiopia. Parasit Vectors. 2015;8(1):1–1.
- 15. Bissell RA. Delayed-impact infectious disease after a natural disaster. J Emerg Med. 1983;1(1):59–66.
- 16. PAHO. Impact of Hurricane Mitch on Central America. Epidemiol Bull. 1998;19:1–14.
- 17. Vittor AY, Gilman RH, Tielsch J, et al. The effect of deforestation on the human-biting rate of Anopheles darlingi, the primary vector of falciparum malaria in the Peruvian Amazon. Am J Trop Med Hyg. 2006;74(1):3–11.
- Olson SH, Gangnon R, Silveira GA, Patz JA. Deforestation and malaria in Mancio Lima county, Brazil. Emerg Infect Dis. 2010;16(7):1108.
- Hahn MB, Gangnon RE, Barcellos C, Asner GP, Patz JA. Influence of deforestation, logging, and fire on malaria in the Brazilian Amazon. PLoS ONE. 2014;9(1): e85725.
- 20. Alves L. Amazon fires coincide with increased respiratory illnesses in indigenous populations. Lancet Respir Med. 2020;8(11): e84.
- 21. Akpan GE, Adepoju KA, Oladosu OR, Adelabu SA. Dominant malaria vector species in Nigeria: modelling potential distribution of Anopheles gambiae sensu lato and its siblings with MaxEnt. PLoS ONE. 2018;13(10): e0204233.
- 22. Le PV, Kumar P, Ruiz MO, Mbogo C, Muturi EJ. Predicting the direct and indirect impacts of climate change on malaria in coastal Kenya. PLoS ONE. 2019;14(2): e0211258.
- 23. Khormi HM, Kumar L. Future malaria spatial pattern based on the potential global warming impact in South and Southeast Asia. Geospat Health. 2016;11(3):416.
- 24. Laporta GZ, Linton YM, Wilkerson RC, et al. Malaria vectors in South America: current and future scenarios. Parasit Vectors. 2015;8(1):1–3.
- 25. Hundessa S, Williams G, Li S, et al. Projecting potential spatial and temporal changes in the distribution of *Plasmodium vivax* and *Plasmodium falciparum* malaria in China with climate change. Sci Total Environ. 2018;15(627):1285–93.
- 26. Colón-González FJ, Sewe MO, Tompkins AM, et al. Projecting the risk of mosquito-borne diseases in a warmer and more populated world: a multi-model, multi-scenario intercomparison modelling study. Lancet Planet Health. 2021;5(7):e404–14.

- 27. Sarkar S, Gangare V, Singh P, Dhiman RC. Shift in potential malaria transmission areas in India, using the fuzzy-based climate suitability malaria transmission (FCSMT) model under changing climatic conditions. Int J Environ Res Public Health. 2019;16(18):3474.
- 28. Bhatt S, Gething PW, Brady OJ, et al. The global distribution and burden of dengue. Nature. 2013;496(7446):504–7.
- 29. Messina JP, Brady OJ, Golding N, et al. The current and future global distribution and population at risk of dengue. Nat Microbiol. 2019;4(9):1508–15.
- 30. Liu-Helmersson J, Brännström Å, Sewe MO, Semenza JC, Rocklöv J. Estimating past, present, and future trends in the global distribution and abundance of the arbovirus vector *Aedes aegypti* under climate change scenarios. Front Public Health. 2019;7:148.
- 31. Semenza JC, Sudre B, Miniota J, et al. International dispersal of dengue through air travel: importation risk for Europe. PLoS Negl Trop Dis. 2014;8(12): e3278.
- 32. Kraemer MU, Sinka ME, Duda KA, et al. The global distribution of the arbovirus vectors *Aedes aegypti* and *Ae. albopictus*. Elife. 2015;4:e08347.
- 33. Stanaway JD, Shepard DS, Undurraga EA, et al. The global burden of dengue: an analysis from the Global Burden of Disease Study 2013. Lancet Infect Dis. 2016;16(6):712–23.
- 34. Messina JP, Brady OJ, Pigott DM, Brownstein JS, Hoen AG, Hay SI. A global compendium of human dengue virus occurrence. Sci Data. 2014;1(1):1–6.
- 35. López MS, Müller GV, Sione WF. Analysis of the spatial distribution of scientific publications regarding vector-borne diseases related to climate variability in South America. Spat Spatiotemporal Epidemiol. 2018;1(26):35–93.
- 36. Mutheneni SR, Morse AP, Caminade C, Upadhyayula SM. Dengue burden in India: recent trends and importance of climatic parameters. Emerg Microbes Infect. 2017;6(1):1.
- 37. Carvajal TM, Viacrusis KM, Hernandez LF, Ho HT, Amalin DM, Watanabe K. Machine learning methods reveal the temporal pattern of dengue incidence using meteorological factors in metropolitan Manila, Philippines. BMC Infect Dis. 2018;18(1):1–5.
- Petrova D, Lowe R, Stewart-Ibarra A, Ballester J, Koopman SJ, Rodó X. Sensitivity of large dengue epidemics in Ecuador to long-lead predictions of El Niño. Clim Serv. 2019;1(15):100096.

- 39. Liu-Helmersson J, Brännström Å, Sewe MO, Semenza JC, Rocklöv J. Estimating past, present, and future trends in the global distribution and abundance of the arbovirus vector *Aedes aegypti* under climate change scenarios. Front Public Health. 2019;12(7):148.
- Yactayo S, Staples JE, Millot V, Cibrelus L, Ramon-Pardo P. Epidemiology of Chikungunya in the Americas. J Infect Dis. 2016;214(suppl_5):S441–5.
- 41. Rezza G, Nicoletti L, Angelini R, et al. Infection with chikungunya virus in Italy: an outbreak in a temperate region. Lancet. 2007;370(9602):1840–6.
- 42. Rocklöv J, Tozan Y, Ramadona A, et al. Using big data to monitor the introduction and spread of Chikungunya, Europe, 2017. Emerg Infect Dis. 2019;25(6):1041.
- 43. Tjaden NB, Suk JE, Fischer D, Thomas SM, Beierkuhnlein C, Semenza JC. Modelling the effects of global climate change on chikungunya transmission in the 21st century. Sci Rep. 2017;7(1):1–1.
- 44. Paz S, Semenza JC. El Niño and climate change contributing factors in the dispersal of Zika virus in the Americas? Lancet. 2016;387(10020):745.
- 45. Tesla B, Demakovsky LR, Mordecai EA, et al. Temperature drives Zika virus transmission: evidence from empirical and mathematical models. Proc R Soc B. 1884;2018(285):20180795.
- 46. Marini G, Calzolari M, Angelini P, et al. A quantitative comparison of West Nile virus incidence from 2013 to 2018 in Emilia-Romagna, Italy. PLoS Negl Trop Dis. 2020;14(1): e0007953.
- Farooq Z, Rocklöv J, Wallin J, Abiri N, Odhiambo Sewe M, Sjödin H, Semenza JC. Artificial intelligence to predict West Nile virus outbreaks with ecoclimatic drivers. Lancet Reg Health – Eur 2022;17: 100370. https://doi.org/10.1016/j.lanepe.2022. 100370.
- 48. Ziegler U, Santos PD, Groschup MH, et al. West Nile virus epidemic in Germany triggered by epizootic emergence, 2019. Viruses. 2020;12(4):448.
- 49. Vlaskamp DR, Thijsen SF, Reimerink J, et al. First autochthonous human West Nile virus infections in the Netherlands, July to August 2020. Eurosurveillance. 2020;25(46):2001904.
- 50. Semenza JC, Tran A, Espinosa L, Sudre B, Domanovic D, Paz S. Climate change projections of West Nile virus infections in Europe: implications for blood safety practices. Environ Health. 2016;15(1): 125–36.

- 51. Semenza JC, Domanović D. Blood supply under threat. Nat Clim Change. 2013;3(5):432–5.
- 52. Clow KM, Leighton PA, Ogden NH, et al. Northward range expansion of Ixodes scapularis evident over a short timescale in Ontario, Canada. PLoS ONE. 2017;12(12): e0189393.
- 53. Cheng A, Chen D, Woodstock K, Ogden NH, Wu X, Wu J. Analyzing the potential risk of climate change on Lyme disease in eastern Ontario, Canada using time series remotely sensed temperature data and tick population modelling. Remote Sens. 2017;9(6): 609.
- 54. Gasmi S, Ogden NH, Lindsay LR, et al. Emerging infections: surveillance for Lyme disease in Canada: 2009–2015. Can Commun Dis Rep. 2017;43(10): 194.
- 55. Ebi KL, Ogden NH, Semenza JC, Woodward A. Detecting and attributing health burdens to climate change. Environ Health Perspect. 2017;125(8): 085004.
- 56. Burrows H, Talbot B, McKay R, et al. A multi-year assessment of blacklegged tick (*Ixodes scapularis*) population establishment and Lyme disease risk areas in Ottawa, Canada, 2017–2019. PLoS ONE. 2021;16(2):e0246484.
- 57. Estrada-Peña A, de la Fuente J. Host distribution does not limit the range of the tick *Ixodes ricinus* but impacts the circulation of transmitted pathogens. Front Cell Infect Microbiol. 2017;11(7):405.
- McPherson M, García-García A, Cuesta-Valero FJ, et al. Expansion of the Lyme disease vector *Ixodes scapularis* in Canada inferred from CMIP5 climate projections. Environ Health Perspect. 2017;125(5): 057008.
- 59. Monaghan AJ, Moore SM, Sampson KM, Beard CB, Eisen RJ. Climate change influences on the annual onset of Lyme disease in the United States. Ticks Tick-Borne Dis. 2015;6(5):615–22.
- 60. Li S, Gilbert L, Vanwambeke SO, Yu J, Purse BV, Harrison PA. Lyme disease risks in Europe under multiple uncertain drivers of change. Environ Health Perspect. 2019;127(6): 067010.
- 61. Lindgren E, Gustafson R. Tick-borne encephalitis in Sweden and climate change. Lancet. 2001;358(9275):16–8.
- 62. Lukan M, Bullova E, Petko B. Climate warming and tick-borne encephalitis, Slovakia. Emerg Infect Dis. 2010;16(3):524.

- 63. Semenza JC, Suk JE. Vector-borne diseases and climate change: a European perspective. FEMS Microbiol Lett. 2018;365(2):fnx244.
- 64. Blomqvist G, Näslund K, Svensson L, Beck C, Valarcher JF. Mapping geographical areas at risk for tick-borne encephalitis (TBE) by analysing bulk tank milk from Swedish dairy cattle herds for the presence of TBE virus-specific antibodies. Acta Vet Scand. 2021;63(1):1–9.
- 65. Wallenhammar A, Lindqvist R, Asghar N, et al. Revealing new tick-borne encephalitis virus foci by screening antibodies in sheep milk. Parasit Vectors. 2020;13(1):1–2.
- 66. Paulsen KM, Stuen S, Das Neves CG, et al. Tickborne encephalitis virus in cows and unpasteurized cow milk from Norway. Zoonoses Public Health. 2019;66(2):216–22.
- 67. https://www.cdc.gov/ticks/tickbornediseases/rmsf. html. Tickborne Diseases of the United States. Accessed 17 Apr 2022.
- 68. Adem PV. Emerging and re-emerging rickettsial infections. Semin Diagn Pathol. 2019;36(3):146–51.
- 69. Karbowiak G. The occurrence of the *Dermacentor reticulatus* tick-its expansion to new areas and possible causes. Ann Parasitol. 2014;60(1):37–47.
- 70. Dahlgren FS, Paddock CD, Springer YP, Eisen RJ, Behravesh CB. Expanding range of *Amblyomma americanum* and simultaneous changes in the epidemiology of spotted fever group rickettsiosis in the United States. Am J Trop Med Hyg. 2016;94(1):35.
- 71. Alten B, Maia C, Afonso MO, et al. Seasonal dynamics of phlebotomine sand fly species proven vectors of Mediterranean leishmaniasis caused by *Leishmania infantum*. PLoS Negl Trop Dis. 2016;10(2):e0004458.
- 72. Antoniou M, Gramiccia M, Molina R, Dvorak V, Volf P. The role of indigenous phlebotomine sandflies and mammals in the spreading of leishmaniasis agents in the Mediterranean region. Eurosurveillance. 2013;18(30):20540.
- 73. Maroli M, Rossi L, Baldelli R, et al. The northward spread of leishmaniasis in Italy: evidence from retrospective and ongoing studies on the canine reservoir and phlebotomine vectors. Trop Med Int Health. 2008;13(2):256–64.
- 74. Ballart C, Barón S, Alcover MM, Portús M, Gállego M. Distribution of phlebotomine sand flies (Diptera: Psychodidae) in Andorra: first finding of *P. perniciosus* and wide distribution of *P. ariasi*. Acta Trop. 2012;122(1):155–9.

1387

- 1388
 - 75. Naucke TJ, Menn B, Massberg D, Lorentz S. Sandflies and leishmaniasis in Germany. Parasitol Res. 2008;103(1):65–8.
 - 76. Fischer D, Moeller P, Thomas SM, Naucke TJ, Beierkuhnlein C. Combining climatic projections and dispersal ability: a method for estimating the responses of sandfly vector species to climate change. PLoS Negl Trop Dis. 2011;5(11): e1407.
 - 77. Wang P, Goggins WB, Chan EY. Associations of Salmonella hospitalizations with ambient temperature, humidity and rainfall in Hong Kong. Environ Int. 2018;1(120):223–30.
 - 78. Lal A, Hales S, Kirk M, Baker MG, French NP. Spatial and temporal variation in the association between temperature and salmonellosis in NZ. Aust N Z J Public Health. 2016;40(2):165–9.
 - 79. Milazzo A, Giles LC, Zhang Y, Koehler AP, Hiller JE, Bi P. Heatwaves differentially affect risk of Salmonella serotypes. J Infect. 2016;73(3):231–40.
 - Lake IR, Colon-Gonzalez FJ, Takkinen J, et al. Exploring campylobacter seasonality across Europe using the European surveillance system (TESSy), 2008 to 2016. Eurosurveillance. 2019;24(13): 1800028.
 - 81. Rosenberg A, Weinberger M, Paz S, Valinsky L, Agmon V, Peretz C. Ambient temperature and agerelated notified Campylobacter infection in Israel: a 12-year time series study. Environ Res. 2018;1(164): 539–45.
 - 82. Djennad A, Iacono GL, Sarran C, et al. Seasonality and the effects of weather on Campylobacter infections. BMC Infect Dis. 2019;19(1):1.
 - 83. Smith BA, Fazil A. Climate change and infectious diseases: the challenges: how will climate change impact microbial foodborne disease in Canada? Can Commun Dis Rep. 2019;45(4):108.
 - 84. Akil L, Ahmad HA, Reddy RS. Effects of climate change on Salmonella infections. Foodborne Pathog Dis. 2014;11(12):974–80.
 - 85. Lake IR. Food-borne disease and climate change in the United Kingdom. Environ Health. 2017;16(1): 53–9.
 - Semenza JC, Herbst S, Rechenburg A, et al. Climate change impact assessment of food-and waterborne diseases. Crit Rev Environ Sci Technol. 2012;42(8): 857–90.
 - Food Safety News. Floodwater pathogens can't be washed off of fresh produce. Food Safety News. 2017. https://www.foodsafetynews.com/2017/09/

floodwater-pathogens-cant-be-washed-off-of-freshproduce/. Accessed 17 Apr 2022.

- Dicker D, Nguyen G, Abate D, et al. Global, regional, and national age-sex-specific mortality and life expectancy, 1950–2017: a systematic analysis for the Global Burden of Disease Study 2017. Lancet. 2018;392(10159):1684–735.
- 89. Ferreira DC, Graziele I, Marques RC, Gonçalves J. Investment in drinking water and sanitation infrastructure and its impact on waterborne diseases dissemination: the Brazilian case. Sci Total Environ. 2021;20(779): 146279.
- 90. Chen L, Deng Y, Dong S, et al. The occurrence and control of waterborne viruses in drinking water treatment: a review. Chemosphere. 2021;30: 130728.
- 91. Mutono N, Wright J, Mutembei H, et al. The nexus between improved water supply and water-borne diseases in urban areas in Africa: a scoping review protocol. AAS Open Res. 2020;3:12.
- 92. Semenza JC, Nichols G. Cryptosporidiosis surveillance and water-borne outbreaks in Europe. Euro Surveill. 2007;12(5):E13–4.
- 93. Herrador BR, De Blasio BF, MacDonald E, et al. Analytical studies assessing the association between extreme precipitation or temperature and drinking water-related waterborne infections: a review. Environ Health. 2015;14(1):1–2.
- 94. Levy K, Woster AP, Goldstein RS, Carlton EJ. Untangling the impacts of climate change on waterborne diseases: a systematic review of relationships between diarrheal diseases and temperature, rainfall, flooding, and drought. Environ Sci Technol. 2016;50(10):4905–22.
- 95. Suk JE, Vaughan EC, Cook RG, Semenza JC. Natural disasters and infectious disease in Europe: a literature review to identify cascading risk pathways. Eur J Public Health. 2020;30(5):928–35.
- 96. Zhang N, Song D, Zhang J, et al. The impact of the 2016 flood event in Anhui Province, China on infectious diarrhea disease: an interrupted timeseries study. Environ Int. 2019;1(127):801–9.
- 97. Epstein A, Benmarhnia T, Weiser SD. Drought and illness among young children in Uganda, 2009–2012. Am J Trop Med Hyg. 2020;102(3):644.
- 98. Subiros M, Brottet E, Solet JL, LeGuen A, Filleul L. Health monitoring during water scarcity in Mayotte, France, 2017. BMC Public Health. 2019;19(1): 1.

- 99. Asmall T, Abrams A, Röösli M, Cissé G, Carden K, Dalvie MA. The adverse health effects associated with drought in Africa. Sci Total Environ. 2021;1(793): 148500.
- 100. Kouadio IK, Aljunid S, Kamigaki T, Hammad K, Oshitani H. Infectious diseases following natural disasters: prevention and control measures. Expert Rev Anti Infect Ther. 2012;10(1):95–104.
- 101. Jutla A, Khan R, Colwell R. Natural disasters and cholera outbreaks: current understanding and future outlook. Curr Environ Health Rep. 2017;4(1): 99–107.
- 102. Moore S, Dongdem AZ, Opare D, et al. Dynamics of cholera epidemics from Benin to Mauritania. PLoS Negl Trop Dis. 2018;12(4): e0006379.
- 103. Camacho A, Bouhenia M, Alyusfi R, et al. Cholera epidemic in Yemen, 2016–18: an analysis of surveillance data. Lancet Glob Health. 2018;6(6): e680–90.
- 104. Olago D, Marshall M, Wandiga SO, et al. Climatic, socio-economic, and health factors affecting human vulnerability to cholera in the Lake Victoria basin, East Africa. AMBIO J Hum Environ. 2007;36(4):350–8.
- 105. Luque Fernández MÁ, Bauernfeind A, Jiménez JD, Gil CL, Omeiri NE, Guibert DH. Influence of temperature and rainfall on the evolution of cholera epidemics in Lusaka, Zambia, 2003–2006: analysis of a time series. Trans R Soc Trop Med Hyg. 2009;103(2):137–43.
- 106. Ruiz-Moreno D, Pascual M, Bouma M, Dobson A, Cash B. Cholera seasonality in Madras (1901–1940): dual role for rainfall in endemic and epidemic regions. EcoHealth. 2007;4(1):52–62.
- 107. Hashizume M, Faruque AS, Wagatsuma Y, Hayashi T, Armstrong B. Cholera in Bangladesh: climatic components of seasonal variation. Epidemiology. 2010;21(5):706–10.
- Semenza JC, Trinanes J, Lohr W, et al. Environmental suitability of Vibrio infections in a warming climate: an early warning system. Environ Health Perspect. 2017;125(10): 107004.
- 109. Lau CL, Smythe LD, Craig SB, Weinstein P. Climate change, flooding, urbanisation and leptospirosis: fuelling the fire? Trans R Soc Trop Med Hyg. 2010;104(10):631–8.
- 110. Dhewantara PW, Hu W, Zhang W, et al. Climate variability, satellite-derived physical environmental data and human leptospirosis: a retrospective ecological study in China. Environ Res. 2019;1(176): 108523.

- 111. Dechet AM, Parsons M, Rambaran M, et al. Leptospirosis outbreak following severe flooding: a rapid assessment and mass prophylaxis campaign; Guyana, January–February 2005. PLoS ONE. 2012;7(7): e39672.
- 112. Al-shere TA, Ujiie M, Suzuki M, et al. Outbreak of leptospirosis after flood, the Philippines, 2009. Emerg Infect Dis. 2012;18(1):91.
- 113. Cissé G. Food-borne and water-borne diseases under climate change in low-and middle-income countries: further efforts needed for reducing environmental health exposure risks. Acta Trop. 2019;1(194):181–8.
- 114. Mbereko A, Chimbari MJ, Manyangadze T, Mukaratirwa S. Knowledge and perceptions of schistosomiasis, a water-borne disease, in two semi-arid rural areas of South Africa (Ndumo) and Zimbabwe (Ntalale). Food Waterborne Parasitol. 2020;1(21): e00091.
- 115. Archer J, O'Halloran L, Al-Shehri H, et al. Intestinal schistosomiasis and giardiasis co-infection in sub-Saharan Africa: can a one health approach improve control of each waterborne parasite simultaneously? Trop Med Infect Dis. 2020;5(3):137.
- 116. Yang GJ, Bergquist R. Potential impact of climate change on schistosomiasis: a global assessment attempt. Trop Med Infect Dis. 2018;3(4):117.
- 117. Mutuku FM, King CH, Bustinduy AL, Mungai PL, Muchiri EM, Kitron U. Impact of drought on the spatial pattern of transmission of *Schistosoma haematobium* in coastal Kenya. Am J Trop Med Hyg. 2011;85(6):1065–70. https://doi.org/10.4269/ ajtmh.2011.11-0186.
- 118. DM Fujita, de Albuquerque Luna EJ. Natural disasters, ecotourism and adventure travel may contribute to Schistosoma mansoni in Chapada Diamantina/Bahia—Brazil—2017. Travel Med Infect Dis. 2019;31:101341.
- 119. Kalinda C, Chimbari MJ, Mukaratirwa S. Effect of temperature on the *Bulinus globosus—Schistosoma haematobium* system. Infect Dis Poverty. 2017;6(1): 1–7.
- 120. McCreesh N, Nikulin G, Booth M. Predicting the effects of climate change on *Schistosoma mansoni* transmission in eastern Africa. Parasit Vectors. 2015;8(1):1–9.
- 121. Zhu G, Fan J, Peterson AT. *Schistosoma japonicum* transmission risk maps at present and under climate change in mainland China. PLoS Negl Trop Dis. 2017;11(10): e0006021.

- 122. Nichols GL, Gillingham EL, Macintyre HL, et al. Coronavirus seasonality, respiratory infections and weather. BMC Infect Dis. 2021;21(1):1–5.
- 123. Park JE, Son WS, Ryu Y, Choi SB, Kwon O, Ahn I. Effects of temperature, humidity, and diurnal temperature range on influenza incidence in a temperate region. Influenza Other Respir Viruses. 2020;14(1):11–8.
- 124. Ianevski A, Zusinaite E, Shtaida N, et al. Low temperature and low UV indexes correlated with peaks of influenza virus activity in Northern Europe during 2010–2018. Viruses. 2019;11(3):207.
- 125. Towers S, Chowell G, Hameed R, et al. Climate change and influenza: the likelihood of early and severe influenza seasons following warmer than average winters. PLoS Curr. 2013;28:5.
- 126. Bayard V, Kitsutani PT, Barria EO, et al. Outbreak of hantavirus pulmonary syndrome, Los Santos, Panama, 1999–2000. Emerg Infect Dis. 2004;10(9): 1635.
- 127. Williams RJ, Bryan RT, Mills JN, et al. An outbreak of hantavirus pulmonary syndrome in western Paraguay. Am J Trop Med Hyg. 1997;57(3):274–82.

- 128. Engelthaler DM, Mosley DG, Cheek JE, et al. Climatic and environmental patterns associated with hantavirus pulmonary syndrome, Four Corners region, United States. Emerg Infect Dis. 1999;5(1): 87.
- 129. Kondo H, Seo N, Yasuda T, et al. Post-flood—infectious diseases in Mozambique. Prehosp Disaster Med. 2002;17(3):126–33.
- 130. Biswas R, Pal D, Mukhopadhyay SP. A communitybased study on health impact of flood in a vulnerable district of West Bengal. Indian J Public Health. 1999;43(2):89–90.
- 131. Yusof A, Siddique AK, Baqui AH, Eusof A, Zaman K. 1988 floods in Bangladesh: pattern of illness and causes of death. J Diarrhoeal Dis Res. 1991;1:310–4.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.