Climate Change Drives the Transmission and Spread of Vector-Borne Diseases: An Ecological Perspective

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Simple Summary: Vector-borne diseases (VBDs) are a major threat to human health. Climate change has a significant impact on VBDs. To clarify the complex effects of climate change on VBDs, we concluded the effects of climate on the transmission and spread of VBDs from an ecological perspective and summarized VBD changes in response to climate change, specifically including: the nonlinear effects of local climate (temperature, precipitation and wind) on VBD transmission, especially temperature showing n-shape effects; regional climate (the El Niño–Southern Oscillation and North Atlantic Oscillation) has time-lag effects on VBD transmission through indirect impact on local climate; and the u-shaped effect of extreme climates can lead to the geographical spread of VBDs. In terms of non-climatic factors, land use and human mobility through the interactions with climatic factors, will affect transmission and spread of VBD. We further explored the uncertainty of the impact of climate change on VBDs under the COVID-19 pandemic. A systematic understanding of the impact of climate change on the transmission and spread of VBD can provide insights and suggestions for future research on VBD prevention and control.

Abstract: Climate change affects ecosystems and human health in multiple dimensions. With the acceleration of climate change, climate-sensitive vector-borne diseases (VBDs) pose an increasing threat to public health. This paper summarizes 10 publications on the impacts of climate change on ecosystems and human health; then it synthesizes the other existing literature to more broadly explain how climate change drives the transmission and spread of VBDs through an ecological perspective. We highlight the multi-dimensional nature of climate change, its interaction with other factors, and the impact of the COVID-19 pandemic on transmission and spread of VBDs, specifically including: (1) the generally nonlinear relationship of local climate (temperature, precipitation and wind) and VBD transmission, with temperature especially exhibiting an n-shape relation; (2) the time-lagged effect of regional climate phenomena (the El Niño–Southern Oscillation and North Atlantic Oscillation) on VBD transmission; (3) the u-shaped effect of extreme climate (heat waves, cold waves, floods, and droughts) on VBD spread; (4) how interactions between non-climatic (land use and human mobility) and climatic factors increase VBD transmission and spread; and (5) that the impact of the COVID-19 pandemic on climate change is debatable, and its impact on VBDs remains uncertain. By exploring the influence of climate change and non-climatic factors on VBD transmission and spread, this paper provides scientific understanding and guidance for their effective prevention and control.

Keywords: climate change; vector-borne diseases; transmission; spread; interaction; COVID-19 pandemic
1. Introduction

Climate change, which has affected the world since the last century, has caused a general rise in temperatures over the period of 1906–2005 [1]. According to the Sixth Assessment Report (AR 6) of the Intergovernmental Panel on Climate Change (IPCC), the global average surface temperature will reach or exceed 1.5 °C in the next two decades, accompanied by increasing precipitation, melting glaciers and rising sea levels [2]. With the acceleration of climate change, extreme weather conditions will be frequent [3] and pose serious threats to human life and health. At present, about 30% of the global population is exposed to extreme weather that exceeds human thermoregulatory capacity for at least 20 days a year [4]. Moreover, global warming and extreme precipitation can contribute to the prevalence and expansion of diseases, leading to at least 150,000 deaths per year worldwide [5].

The ten publications in this Special Issue illuminate the impacts of climate change on ecosystems and human health from different perspectives in diverse disciplines, including phytochemistry, biology, epidemiology, pathology, and molecular biology. In phytochemistry and biology, the geographical ranges of plants and animals have been shown to be affected by climate change [6–8]. In epidemiology and pathology, the future geographic expansion of vector species that carry vector-borne diseases (VBDs) has been evaluated [9–12], and proved the invasive and evolutionary adaptation of vectors to different ecological and environmental conditions [13]. The impact of non-climatic factors on VBDs has also been assessed [14]. On the basis of these ten articles, we present a further discussion on climate change and VBDs.

As a climate-sensitive type of disease, VBDs are assessed on a global scale with the aim of shedding light on possible future trends, particularly given the increased likelihood of climate change [15]. The impact of climate change on VBDs has become an indisputable fact, and is creating new challenges for public health. As a category, VBDs include rodent-borne (plague, hemorrhagic fever, hemorrhagic fever with renal syndrome, leptospirosis, cutaneous leishmaniasis, and Puumala hantavirus), mosquito-borne (malaria, dengue, Zika, chikungunya, West Nile virus, Ross River virus, and Japanese encephalitis), tick-borne (tick-borne encephalitis, Lyme disease, etc.), and other arthropod-borne diseases [16,17]. Over 700,000 people die from VBDs each year, and more than 80% of the global population lives in high-risk areas threatened by one or more types of VBDs [18]. Of the 250 countries around the world, 86% (218 countries) are suitable for arboviral disease survival and reproduction [19]. Accordingly, a large number of scientists have devoted considerable effort to studying the impact of climate change on VBDs, with 2133 related studies having been published as of August 2022 (Figure 1).

![Figure 1. Annual publications on climate change and VBDs. Articles identified by searching the Web of Science with the combination of the following key words: “climate change” and “vector-borne disease”; counts as of 11 August 2022.](image-url)
The influence of climate factors on the transmission and spread of VBDs can be considered at the levels of local climate, regional climate, and extreme climate. Local climate, represented by temperature, rainfall, and wind, mainly affects the transmission of VBDs by affecting their vectors [20]; regional climate, represented by the El Niño–Southern Oscillation (ENSO) [21], North Atlantic Oscillation (NAO) [22], Pacific Decadal Oscillation (PDO) [23], and Indian Ocean Dipole (IOD) [24], mainly has indirect impacts on VBDs through affecting local climate [25]. Meanwhile, extreme climate events such as heat waves, cold waves, floods, and droughts increase the risk of VBD spillover [26]. We additionally discuss the interaction between non-climatic factors (e.g., land use and human mobility) and climate factors as relates to VBD transmission and spread [27–29]. Moreover, we also discuss the impact of coronavirus disease 2019 (COVID-19) on climate change and the effects of the COVID-19 pandemic on the outbreak risk and incidence of VBDs [30]. All told, our paper aims to comprehensively assess the impacts of climate change on the transmission and spread of VBDs so as to support the precise prevention and control of and comprehensive intervention in VBDs.

2. Non-Linear Effects of Local Climate on VBD Transmission

The effects of local climate factors (mainly considering temperature, precipitation, wind) on VBD transmission are generally nonlinear. These factors can affect the distribution range, population dynamics, and virus transmission ability of vectors [29], and hence the developmental response of pathogens [31].

The non-linear effect of temperature on VBD transmission generally follows an n-shape (Figure 2). Under suitable temperature conditions, the climate adaptability of VBD transmission will be relatively high. When the temperature does not reach suitable conditions, the risk of VBD transmission increases with the increase of temperature, and when the temperature exceeds the peak of the suitable temperature, the risk of VBD transmission decreases with the increase of temperature. Under unsuitable temperature conditions, vector survival may be reduced [32], thereby reducing the transmission capacity of VBDs [33], which also directly affects the development of vector-dependent pathogens [34]. For example, the survival and reproduction range of rodents is generally 10.0–30.0 °C, while 20.0–30.0 °C is the suitable temperature range for rodent-borne disease transmission [35]. The predicted epidemic growth of plague outbreaks is positive between 11.7 °C and 21.5 °C, with a maximum around 17.3 °C [36]. With regard to mosquito-borne diseases, temperature can affect the development and survival of mosquitoes, and there is a thermal optimum which will be suppressed at either heat or cold [37]; including malaria, dengue and Zika, temperature and climate change are reported to have strong nonlinear effects on ectothermic vectors and parasites. The temperature range for the transmission of mosquito-borne diseases is generally 9.0–38.0 °C, with the most suitable range being 23.0–29.0 °C [38].

![Figure 2. The effect of local temperature on VBD transmission shows a clear n-shape. Suitable temperature promotes the development and transmission ability of VBD vectors, while unsuitable temperature will affect reproduction and mobility and even cause death of vectors.](image-url)
Precipitation has also demonstrated a nonlinear effect on VBD transmission in many studies. A suitable level of precipitation may be beneficial for the formation of vector breeding habitats [39], especially in deserts, which can provide rich food sources for rodents [40]. Precipitation is also generally beneficial to mosquito oviposition and reproduction, while the relationship between rainfall and incidence of malaria first increases and then decreases with increasing precipitation, reaching its peak at 120 mm [41]. With regard to ticks, a span of more than 28 precipitation days leads the number of ticks to increase significantly [42]; however, large amounts and long periods of precipitation can wash away ticks, their eggs, and their larvae, thus reducing the population [43].

High wind speeds hamper mosquitos in their flight, can decrease the density of mosquitoes, and make them less likely to stand on and bite their hosts. An example to support this view is the finding that high wind impeded the rate of West Nile virus transmission; conversely, there is no obvious negative trend in the effect of low wind speed on mosquitoes. [44,45]. This also represents a non-linear effect on VBD transmission.

Climate change has an important impact on the transmission of vector-borne diseases, which in general will expand the climate-adaptive transmission zone of vector-borne diseases. In Europe, climate change is likely to expand ticks into higher latitudes and altitudes, thereby increasing the incidence of tick-borne diseases [46]. In South Africa, however, rising temperatures could decrease habitat suitability for some tick species (Acari: Ixodidae), which will decrease the occurrence of the related diseases [47]. Under climate scenarios from the IPCC, the climatic suitability of chikungunya transmission will increase in western and central parts of Europe, but will not generally be suitable in Southern Europe [48]. Due to climate change, the suitability of rodents in certain high-altitude areas has increased by 40% [49]. For mosquitoes and ticks, warming climate generally increases the risk of associated disease transmission at high-latitude and -altitude areas, while the risk of transmission may generally decrease in tropical regions. For rodent-borne diseases such as plague, rodents and fleas both influence pathogen transmission; there is uncertainty about the effect of high temperatures on the inhibition of fleas (vectors) and flea-mediated transmission of pathogenic bacteria [50].

3. Time-Lag Effect of Local and Regional Climate Impacts on VBD Transmission

Regional climate mainly exerts its influence on VBDs through local climate factors, which in turn affect the ecological habitat, distribution, and population dynamics of the vectors [51], and hence the transmission rate of the pathogens [24,52], thereby impacting outbreaks of VBDs [53–55]. The complexity of these indirect effects can create additional time-lag effects (Figure 3).

Numerous studies have concluded that local climate has a time-lag effect on VBD transmission in the short term. For rodent-borne diseases, many studies have also demonstrated a time-lag effect of regional climate, such as on renal hemorrhagic fever [56,57], leptospirosis [58], and cutaneous leishmaniasis [59]. The time-lag is 1–6 months or even one year due to the complex biological characteristics of rodent-borne diseases [35]; for example, temperature affects the human plague in Arizona and New Mexico with a 2–3 month lag effect, while precipitation has a 1–2 year lag effect [60]. For mosquito-borne diseases, a large number of studies have investigated time-lag effects, including on dengue fever [61,62], malaria [63–65], chikungunya [66], Ross River virus [67–69], and Japanese encephalitis [70]. The time-lag for these diseases is usually considered to be about 0–2 months due to indirect effects on the life history and density of mosquitoes [71].

Time-lag effects of regional climate have wider ranges and longer timespans than local climate effects, and hence are more relevant to making predictions for disease prevention in advance. ENSO is the most significant example of quasi-periodic climate variability on an interannual scale that can affect weather all over the world [21]. The pattern of global climate variability associated with ENSO has been shown to impact a number of infectious diseases, including rodent-borne diseases [24], mosquito-borne diseases [72,73], and tick-borne diseases [59]. For example, increases in the rate of human plague in China were
well-associated with ENSO over short periods (2–3 years), medium periods (6–7 years), and long periods (11–12 years, 30–40 years) [74]. ENSO-driven dengue cases in India between 2010–2017 were likewise positively associated with a 3–6 month time-lag [62], which would help us to predict human outbreaks in advance. However, the ENSO index does not seem to be an accurate index of climate variability in Europe; instead, the NAO has been found to impact outbreaks of 13 infectious diseases [22]. Moreover, multi-decadal temperature changes have been shown to influence the NAO–plague correlation, with 15–22 years lagged impact in different European regions [75]. All told, these regional climates clearly affect the occurrence of VBDs and human health by influencing precipitation and temperature [76–78], and could be used as early signals for disease control and prevention.

![Regional Climate: ENSO, NAO](image)

**Regional climate:**
- ENSO, NAO

**Local climate:**
- temperature, precipitation

**Ecological habitat**

**Vector distribution, population dynamics**

**Pathogen transmission rates**

**VBD transmission**

*Figure 3.* The indirect time-lagged effect of regional climate on VBD transmission in a bottom-up ecosystem. For example, temperature and precipitation have 0–2 month lagged effects on dengue transmission, while the impact of ENSO lags by 3–6 months.

### 4. Impact of Extreme Climate Distribution Expansion on VBD Spread

With the frequency of climate extremes increasing as climate change accelerates, it is increasingly important to understand the impact of climate range edges and limitations on VBDs (Figure 4A). Most extreme climate conditions have a u-shaped effect on the spread of VBDs. Under adaptive climatic conditions, the lowest risk of VBD spread is usually observed, whereas under extreme climates which lower or increase the conditions of adaptive climate, the risk of VBD spread will increase. This means that extreme climate events may increase the risk of disease transmission and the spillover of VBDs, whereas under adaptive climatic conditions, the expansion of VBDs is lower (Figure 4B). Extreme climate has been found to be one of the main causes of disease outbreaks and is a cause of alarm in the global community. The impact of climate change on VBDs is more significant in the fringes of different climatic areas, which strongly influences the geographical distribution of vectors [20].
Different types of extreme climate events (heat waves, cold waves, floods, and droughts) have different effects on the distribution expansion of VBDs. The impact of heat waves on mosquitoes depends on the onset time and duration; such events usually promote mosquito population growth in early developmental stages, but often suppress the entire life cycle [79]. Thus, under short-term heat waves, it is advised to guard against the spreading of mosquito-borne VBDs caused by rapid growth of mosquitoes. On the other hand, an experimental study in Kenya found that cold waves during summer months were more favorable for mosquito growth on account of the extremely warm year-round temperature; hence, cold waves in Kenya keep summer cooler and are conducive to VBD spread [80]. Meanwhile, floods wash away the aquatic stage of mosquitoes and their eggs from their breeding sites [81], while the stagnant water left after flood recession provides a suitable habitat for mosquitoes [82–84]. When wetlands experience occasional droughts, mosquito populations suddenly explode as their predators and competitors are eliminated [85]. Such increases in mosquito populations would also lead to high-risk spillover of mosquito-borne infectious diseases. Climatic trends also impact VBDs; for example, in northern China (arid climate), rodents are expected to respond positively to high precipitation, whereas in southern China (humid climate), excessive precipitation would destroy rodent nests [86], which impacts human plague intensity due to its positive correlation with rodent density [87].

Thus, in the context of climate change, climate extremes are increasingly expected to create additional risks and possibilities for the spread of VBDs [88,89]. When there is extreme heat in winter, the lack of snow cover makes contact between bank voles and humans easier, such as that which occurred to produce the Puumala hantavirus (PUUV) epidemic of 2006–2007 [90]. An Ecuador study found that under extreme climate, Aedes aegypti can expand its distribution in mountainous areas by up to 4215 km², which would put over 12,000 people at risk of disease [37]. In India, an increase of heat wave events has made chikungunya and dengue diseases more prevalent in coastal districts, and Japanese encephalitis and malaria more prevalent in interior districts [91]. Extreme heat, drought, and flooding all have a negative impact on tick distribution, which may disrupt the habitat of Ixodes ticks in Europe, especially Northern and Central Europe; however, extreme weather is expected to expand the distribution of Ixodes ticks in Europe by 3.8% during 2040–2060, and tick-borne encephalitis (TBE) is expected to spread to high altitudes and latitudes [92]. In New York State in America, the annual number of Lyme disease cases increase 4–10% under mild winter temperatures, and increase 2% under extended spring and summer days [93].
5. Interaction between Non-Climate and Climate Factors Alters VBD Spread

Beyond climate factors alone, the interactions of non-climatic factors (land use and human mobility) and climate factors are important to consider for their impacts on VBD spread [27,28,94].

The interaction of land use and climate change will provide opportunities for pathogen exchange among geographically isolated wildlife, and thus in some cases will promote disease spillover [95]. Projections under climate change and land use in 2070 have indicated that in Asia and Africa, species will converge into new communities at high altitudes, biodiversity hotspots, and areas of high population density, resulting in approximately 4000 times greater cross-species transmission of their associated viruses [95].

Human mobility is also a major factor in the global spread of VBDs [96,97]. With climate change making some areas uninhabitable (as with the severe drought in sub-Saharan Africa), the interaction of climate change and human mobility will manifest as viruses traveling along with mass migrants [27].

6. The COVID-19 Pandemic Introduces a New Situation for VBD Epidemics

The impact of the COVID-19 pandemic on climate change is debatable in the short term [98–101]; however, COVID-19 as a background may result in new circumstances that impact the occurrence of VBD epidemics. On the one hand, human activities and air pollution have been reduced during lockdowns in the COVID-19 pandemic, and climate change has been mitigated [98]. On the other hand, a similar reduction in global SO\textsubscript{2} emissions was found to weaken the aerosol cooling effect, which can lead to warming [102]. These possible climate changes may have new effects on the transmission and spread of climate-sensitive VBDs.

Besides climate change, lack of vector testing and control activities [103–106] and insufficient financial support for VBD surveillance [107,108] during the COVID-19 pandemic have led to increased prevalence of VBDs. Routine vector testing and control activities required by the Department of Prevention and Control, such as regular household surveys, have been suspended during COVID-19 quarantine [103–106]. Many countries have temporarily suspended adult surveillance and larval control measures for Aedes aegypti, resulting in an increased risk of dengue transmission [109]. At the same time, due to the economic pressure caused by COVID-19, financial support for VBD surveillance is insufficient [107,108]. In addition, some VBDs with similar symptoms have been marginalized and underdiagnosed during the COVID-19 pandemic, resulting in VBDs being ignored rather than eliminated [30,110].

However, lockdown policies have greatly reduced imported cases and blocked sources of VBD transmission [110–112]. Statistics indicate that the number of vector-borne cases declined dramatically during the COVID-19 pandemic in many countries [111–114]. There are two important reasons for the decrease in the number of VBDs. One reason is that entry–exit control in different countries have greatly reduced imported cases and blocked the source of disease transmission [110]. The other is that the decreasing of human outdoor activities and physical distancing interventions reduced the bite chance of mosquitoes, and consequently reduced the risk of mosquito-borne disease transmission [111,112]. Therefore, prevention and control targeting COVID-19 transmission also has a preventive effect on VBDs.

7. Conclusions

In this paper, we summarized the different impacts of multiple climatic factors on the transmission and spread of VBDs in the context of climate change. Local climate exerts non-linear direct effects, resulting in rapid transmission in suitable conditions and decline in an unsuitable environment, with local temperature in particular showing a clear n-shape. Regional climate has an indirect impact on VBDs, affecting transmission through its effects on local climate, which by necessity produces a certain time-lag for the effect on disease transmission. Extreme climate events can increase the spread of disease, leading to the
expansion of VBD distributions. Moreover, land use and human mobility have an important interaction effect on VBD spread, increasing the possibilities for spread and spillover. The impact of the COVID-19 pandemic on how climate change affects VBD transmission and spread is yet uncertain.

Quarantine policies during the COVID-19 pandemic successfully blocked the import of VBD cases; this effective prevention and control policy is worth adopting and applying in the field of VBDs. Meanwhile, the impact of COVID-19 on climate change is controversial, and its potential effect on VBDs may gradually become clear in the future. Accordingly, it remains necessary to further explore the potential of COVID-associated climate change to drive effects on VBDs. Meanwhile, the improvement of surveillance systems in relation to the COVID-19 pandemic and the construction of a surveillance network are also worthy of application in VBD surveillance. With the continuous improvement of monitoring systems, it also becomes necessary to adopt methods from the fields of machine learning and artificial intelligence to handle large databases with complex algorithms in the future.

There is still a lot of research worth undertaking with regard to climate change and VBDs. One important research direction is to integrate multidisciplinary factors to analyze the impact of climate change on VBDs, especially with reference to the fields of computer science, zoology, entomology, ecology, and epidemiology. Through the integration of multiple disciplines, we can not only better understand the impacts of climate change on VBDs, but also develop a deeper understanding of the occurrence and development mechanisms of these infectious diseases. Such findings could contribute to achieving a better understanding of how climate change drives effects on VBD risk and spread, thereby improving the prevention and control of VBDs and so improving human health.

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References
1. IPCC. Climate Change 2007: The Physical Science Basis. Contribution of Working Group I to the Fourth Assessment Report of the Intergovernmental Panel on Climate Change; Cambridge University Press: Cambridge, UK; New York, NY, USA, 2007.
2. IPCC. Climate Change 2021: The Physical Science Basis. Contribution of Working Group I to the Sixth Assessment Report of the Intergovernmental Panel on Climate Change; Cambridge University Press: Cambridge, UK; New York, NY, USA, 2021; pp. 33–144. [CrossRef]
3. Mora, C.; Frazier, A.G.; Longman, R.J.; Dacks, R.S.; Walton, M.M.; Tong, E.J.; Sanchez, J.J.; Kaiser, L.R.; Stender, Y.O.; Anderson, J.M.; et al. The Projected Timing of Climate Departure from Recent Variability. Nature 2013, 502, 183–187. [CrossRef] [PubMed]
4. Mora, C.; Doussset, B.; Caldwell, I.; Powell, F.; Geronimo, R.; Bielecki, C.; Counsell, C.; Dietrich, B.; Johnston, E.; Louis, L.; et al. Global Risk of Deadly Heat. Nat. Clim. Chang. 2017, 7, 501–506. [CrossRef]
5. Patz, J.A.; Campbell-Lendrum, D.; Holloway, T.; Foley, J.A. Impact of Regional Climate Change on Human Health. Nature 2005, 438, 310–317. [CrossRef] [PubMed]
6. Yuan, X.; Yang, L.; Li, H.; Wang, L. Spatiotemporal Variations of Plague Risk in the Tibetan Plateau from 1954-2016. Biology 2022, 11, 304. [CrossRef]
7. Yin, Y.; He, Q.; Pan, X.; Liu, Q.; Wu, Y.; Li, X. Predicting Current Potential Distribution and the Range Dynamics of Pomacea Canaliculata in China under Global Climate Change. Biology 2022, 11, 110. [CrossRef]
8. Kumari, P.; Wani, I.A.; Khan, S.; Verma, S.; Mushtaq, S.; Gulnaz, A.; Paray, B.A. Modeling of Valeriana Wallichii Habitat Suitability and Niche Dynamics in the Himalayan Region under Anticipated Climate Change. Biology 2022, 11, 498. [CrossRef]
11. Ma, D.; Lun, X.; Li, C.; Zhou, R.; Zhao, Z.; Wang, J.; Zhang, Q.; Liu, Q. Predicting the Potential Global Distribution of Amblyomma americanum (Acari: Ixodidae) under Near Current and Future Climatic Conditions, Using the Maximum Entropy Model. *Biology* 2021, 10, 1057. [CrossRef]

12. Li, C.; Gao, Y.; Chang, N.; Ma, D.; Zhou, R.; Zhao, Z.; Wang, J.; Zhang, Q.; Liu, Q. Risk Assessment of Anopheles Philippinensis and Anopheles Nivipes (Diptera: Culicidae) Invading China under Climate Change. *Biology* 2021, 10, 998. [CrossRef]

13. Abubakar, M.; Sami, H.; Mahdi, I.; Altahir, O.; Abdelbagi, H.; Mohamed, N.S.; Ahmed, A. The Phylogeography and Spread of the Invasive Asian Malaria Vectors, Anopheles Stephensi, in Sudan. *Biology* 2022, 11, 409. [CrossRef] [PubMed]

14. Li, Z.; Gurgel, H.; Xu, L.; Yang, L.; Dong, J. Improving Dengue Forecasts by Using Geospatial Big Data Analysis in Google Earth Engine and the Historical Dengue Information-Aided Long Short Term Memory Modeling. *Biology* 2022, 11, 169. [CrossRef] [PubMed]

15. Githeko, A.K.; Lindsay, S.W.; Confalonieri, U.E.; Patz, J.A. Climate Change and Vector-Borne Diseases: A Regional Analysis. *Bull. World Health Organ.* 2000, 78, 1136–1147.

16. Kilpatrick, A.M.; Randolph, S.E. Drivers, Dynamics, and Control of Emerging Vector-Borne Zoonotic Diseases. *Lancet* 2012, 380, 1946–1955. [CrossRef]

17. Wilder-Smith, A.; Gubler, D.J.; Weaver, S.C.; Monath, T.P.; Heymann, D.L.; Scott, T.W. Epidemic Arboviral Diseases: Priorities for Research and Public Health. *Lancet Infect. Dis.* 2017, 17, e101–e106. [CrossRef]

18. World Health Organization & UNICEF/UNDP/World Bank. WHO Special Programme for Research and Training in Research and Training in Vector-Borne Diseases, Global Vector Control Response 2017–2030; World Health Organization: Geneva, Switzerland, 2017; ISBN 978-92-4-151297-8.

19. Leta, S.; Beyene, T.J.; Clercç, E.M.D.; Amenu, K.; Kraemer, M.U.G.; Revie, C.W. Global Risk Mapping for Major Diseases Transmitted by Aedes Aegypti and Aedes Albopictus. *J. Infect. Dis.* 2018, 67, 25–35. [CrossRef]

20. Fouque, F.; Reeder, J.C. Impact of Past and On-Going Changes on Climate and Weather on Vector-Borne Diseases Transmission: A Look at the Evidence. *Infect. Dis. Res.* 2014, 8, 51. [CrossRef]

21. Kovats, R.S. El Niño and human health. *Bull. World Health Organ.* 2000, 78, 1127–1135.

22. Morand, S.; Owers, K.A.; Varetn-Szukta, A.; McIntyre, K.M.; Baylis, M. Climate Variability and Outbreaks of Infectious Diseases in Europe. *Sci. Rep.* 2013, 3, 1774. [CrossRef]

23. Ben Ari, T.; Gershunov, A.; Gage, K.L.; Snäll, T.; Ettestad, P.; Kausrud, K.L.; Stenseth, N.C. Human Plague in the USA: The Importance of Regional and Local Climate. *Biol. Lett.* 2008, 4, 737–740. [CrossRef]

24. Kreppel, K.S.; Caminade, C.; Telfer, S.; Owers, K.; McIntyre, K.; Baylis, M. A Non-Stationary Relationship between Global Climate Phenomena and Human Plague Incidence in Madagascar. *PLoS Negl. Trop. Dis.* 2014, 8, e3155. [CrossRef] [PubMed]

25. Fisman, D.N.; Tuite, A.R.; Brown, K.A. Impact of El Niño Southern Oscillation on Infectious Disease Hospitalization Risk in the United States. *Proc. Natl. Acad. Sci. USA* 2016, 113, 14589–14594. [CrossRef]

26. El-Sayed, A.; Kamel, M. Climatic Changes and Their Role in Emergence and Re-Emergence of Diseases. *Environ. Sci. Pollut. Res.* 2020, 27, 22336–22352. [CrossRef]

27. Rosenthal, J. Climate Change and the Geographic Distribution of Infectious Diseases. *EcoHealth* 2009, 6, 489–495. [CrossRef]

28. Franklínos, L.H.V.; Jones, K.E.; Redding, D.W.; Abubakar, I. The Effect of Global Change on Mosquito-Borne Disease. *Lancet Infect. Dis.* 2019, 19, e302–e312. [CrossRef]

29. Medlock, J.M.; Leach, S.A. Effect of Climate Change on Human Health Related Vector-Borne Diseases Risk in the UK. *Lancet Infect. Dis.* 2015, 15, 721–730. [CrossRef]

30. Singh, N.S.; Singh, D.P. Impact of Climate Change on Human Health Related Vector-Borne Diseases in the Present Scenario of COVID-19 in India. *Environ. Res. 2020, 4*, 631–638. [CrossRef]

31. Sternberg, E.D.; Thomas, M.B. Local Adaptation to Temperature and the Implications for Vector-Borne Diseases. *Trends Parasitol.* 2014, 30, 115–122. [CrossRef]

32. Christiansen-Jucht, C.; Parham, P.E.; Saddler, A.; Koella, J.C.; Basañez, M.-G. Temperature during Larval Development and Adult Maintenance Influences the Survival of Anopheles Gambiense s.s. *Parasites Vectors* 2014, 7, 489. [CrossRef]

33. Westbrook, C.J.; Reiskind, M.H.; Pesko, K.N.; Greene, K.E.; Lounibos, L.P. Larval Environmental Temperature and the Susceptibility of Aedes Albopictus Skuse (Diptera: Culicidae) to Chikungunya Virus. *Vector Borne Zoonotic Dis.* 2010, 10, 241–247. [CrossRef]

34. Perkins, T.A.; Metcalf, C.J.E.; Grenfell, B.T.; Tatem, A.J. Estimating Drivers of Autochthonous Transmission of Chikungunya Virus in Its Invasion of the Americas. *PloS Curr.* 2015, *7*, ecurrents.outbreaks.a47b6ac10e0420b1788e9767946d1fe. [CrossRef] [PubMed]

35. Liu, Q. Impact of Climate Change on Vector-Borne Diseases and Related Response Strategies in China: Major Research Findings and Recommendations for Future Research. *Chin. J. Vector Biol Control* 2021, 32, 1–11. [CrossRef]

36. Krauer, F.; Viljugrein, H.; Dean, K.R. The Influence of Temperature on the Seasonality of Historical Plague Outbreaks. *Proc. R. Soc. B-Biol. Sci.* 2021, 288, 20202725. [CrossRef]
37. Reinhold, J.M.; Lazzari, C.R.; Lahondere, C. Effects of the Environmental Temperature on Aedes Aegypti and Aedes Albopictus Mosquitoes: A Review. *Insects* **2018**, *9*, 158. [CrossRef] [PubMed]

38. Mordecii, E.A.; Caldwell, J.M.; Grossman, M.K.; Lippi, C.A.; Johnson, L.R.; Neira, M.; Rohr, J.R.; Ryan, S.J.; Savage, V.; Shocket, M.S.; et al. Thermal Biology of Mosquito-Borne Disease. *Ecol. Lett.* **2019**, *22*, 1690–1708. [CrossRef] [PubMed]

39. Orland, M.; Kelt, D. Responses of a Heteromyid Rodent Community to Large and Small-Scale Resource Pulses: Diversity, Abundance, and Home-Range Dynamics. *J. Mammal.* **2007**, *88*, 1280–1287. [CrossRef]

40. Matsushita, N.; Kim, Y.; Ng, C.F.S.; Moriyama, M.; Igarashi, T.; Yamamoto, K.; Otiennio, W.; Minakawa, N.; Hashizume, M. Differences of Rainfall-Malaria Associations in Lowland and Highland in Western Kenya. *Int. J. Environ. Res. Public Health* **2019**, *16*, 3693. [CrossRef]

41. Wan, X.; Holyoak, M.; Yan, C.; Le Maho, Y.; Dirzo, R.; Krebs, C.J.; Stenseth, N.C.; Zhang, Z. Broad-Scale Climate Variation Drives Host-Seeking Behavior and Its Associated Climatic Variables in Aedes albopictus. *Parasites Vectors* **2019**, *12*, 474. [CrossRef] [PubMed]

42. Fischer, D.; Thomas, S.M.; Suk, J.E.; Sudre, B.; Hess, A.; Tjaden, N.B.; Beierkuhnlein, C.; Semenza, J.C. Climate Change Effects on the Dynamics of Animal Populations: A Global Multi-Taxa Analysis. *Biol. Rev.* **2009**, *84*, 595–615. [CrossRef] [PubMed]

43. Strathnairn, M.J.; Stryers, V.; Petroysqliakis, T.; Part, P.; Gewehr, S.; Kalaitzopoulou, S.; Mourelatos, S.; Baka, A.; Pervanidou, D.; Vontas, J.; et al. Identification of Climatic Factors Affecting the Epidemiology of Human West Nile Virus Infections in Northern Greece. *PLoS ONE* **2016**, *11*, e0161510. [CrossRef] [PubMed]

44. Estrella-Peña, A. Climate Change Decreases Habitat Suitability for Some Tick Species (Acari: Ixodidae) in South Africa. *PLoS ONE* **2019**, *12*, 247. [CrossRef] [PubMed]

45. Yin, Q.; Li, L.; Guo, X.; Wu, R.; Shi, B.; Wang, Y.; Liu, Y.; Wu, S.; Pan, Y.; Wang, Q.; et al. A Field-Based Modeling Study on Ecological Characterization of Hourly Host-Seeking Behavior and Its Associated Climatic Variables in Aedes albopictus. *Parasites Vectors* **2019**, *12*, 474. [CrossRef] [PubMed]

46. Canzani, J.; Gock, G.; Kochmann, J.; Klimpel, S. Ticks on the Move—Climate Change-Induced Range Shifts of Three Tick Species in Europe: Current and Future Habitat Suitability for *Ixodes ricinus* in Comparison with *Dermacentor reticulatus* and *Dermacentor marginatus*. *Parasitol. Res.* **2022**, *121*, 2241–2252. [CrossRef] [PubMed]

47. Estrada-Peña, A. Climate Change Decreases Habitat Suitability for Some Tick Species (Acari: Ixodidae) in South Africa. *Onderstepoort J. Vet. Res.* **2003**, *70*, 79–93. [PubMed]

48. Fischer, D.; Thomas, S.M.; Suk, J.E.; Sudre, B.; Hess, A.; Tjaden, N.B.; Beierkuhnlein, C.; Semenza, J.C. Climate Change Effects on Chikungunya Transmission in Europe: Geospatial Analysis of Vector’s Climatic Suitability and Virus’ Temperature Requirements. *Int. J. Health Geograph.* **2013**, *12*, 51. [CrossRef]

49. Carlson, C.J.; Bevins, S.N.; Schmid, B.V. Plague Risk in the Western United States over Seven Decades of Environmental Change. *Glob. Chang. Biol.* **2022**, *28*, 753–769. [CrossRef]

50. Snäll, T.; Benestad, R.E.; Stenseth, N.C. Expected Future Plague Levels in a Wildlife Host under Different Scenarios of Climate Change. *Glob. Chang. Biol.* **2009**, *15*, 500–507. [CrossRef]

51. Wang, X.; Holyoak, M.; Yan, C.; Le Maho, Y.; Dirzo, R.; Krebs, C.J.; Stenseth, N.C.; Zhang, Z. Broad-Scale Climate Variation Drives the Dynamics of Animal Abundance and Weather Conditions Potentially Drive Hemorrhagic Fever with Renal Syndrome Outbreaks in Xian, China, 2002–2015. *PLoS Neglect. Trop. Dis.* **2015**, *9*, e0003530. [CrossRef] [PubMed]

52. Joshi, Y.P.; Kim, E.-H.; Cheong, H.-K. The Influence of Climatic Factors on the Development of Hemorrhagic Fever with Renal Syndrome and Leptospirosis during the Peak Season in Korea: An Ecologic Study. *Proc. Natl. Acad. Sci. USA* **2017**, *114*, 119–124. [CrossRef]

53. Hiko, A.; Malicha, G. Climate Change and Animal Health Risk. In *Climate Change and the 2030 Corporate Agenda for Sustainable Development*; Emerald group Publishing Limited: Howard House, Bingley, UK, 2017; Volume 19, pp. 77–111, ISBN 978-1-78635-818-9.

54. Ciota, A.T.; Keyel, A.C. The Role of Temperature in Transmission of Zoonotic Arboviruses. *Viruses* **2019**, *11*, 1013. [CrossRef]

55. Rocklov, J.; Dubrow, R. Climate Change: An Enduring Challenge for Vector-Borne Disease Prevention and Control. *Nat. Immunol.* **2020**, *21*, 479–483. [CrossRef]

56. Tian, H.-Y.; Yu, P.-B.; Luis, A.D.; Bi, P.; Cazelles, B.; Laine, M.; Huang, S.-Q.; Ma, C.-F.; Zhou, S.; Wei, J.; et al. Changes in Rodent Abundance and Weather Conditions Potentially Drive Hemorrhagic Fever with Renal Syndrome Outbreaks in Xian, China, 2002–2015. *PLoS Neglect. Trop. Dis.* **2015**, *9*, e0003530. [CrossRef] [PubMed]

57. Joshi, Y.P.; Kim, E.-H.; Cheong, H.-K. The Influence of Climatic Factors on the Development of Hemorrhagic Fever with Renal Syndrome and Leptospirosis during the Peak Season in Korea: An Ecologic Study. *BMJ Infect. Dis.* **2017**, *17*, 406. [CrossRef] [PubMed]

58. Hubalek, Z. North Atlantic Weather Oscillation and Human Infectious Diseases in the Czech Republic, 1951–2003. *Eur. J. Epidemiol.* **2005**, *20*, 263–270. [CrossRef]

59. Fernando Chaves, L.; Calzada, J.E.; Valderrama, A.; Saldana, A. Cutaneous Leishmaniasis and Sand Fly Fluctuations Are Associated with El Niño in Panama. *PLoS Neglect. Trop. Dis.* **2014**, *8*, e3210. [CrossRef]

60. Enscore, R.E.; Biggerstaff, B.J.; Brown, T.L.; Fulgham, R.E.; Reynolds, P.J.; Engelthaler, D.M.; Levy, C.E.; Parmenter, R.R.; Montenier, J.A.; Cheek, J.E.; et al. Modeling Relationships between Climate and the Frequency of Human Plague Cases in the Southwestern United States, 1960–1997. *Am. J. Trop. Med. Hyg.* **2002**, *66*, 186–196. [CrossRef]

61. Hii, Y.L.; Rocklov, J.; Wall, S.; Ng, L.C.; Tang, C.S.; Ng, N. Optimal Lead Time for Dengue Forecast. *PLoS Neglect. Trop. Dis.* **2012**, *6*, e1848. [CrossRef] [PubMed]
Biology 2022, 11, 1628

88. Cai, W.; Zhang, C.; Zhang, S.; Ai, S.; Bai, Y.; Bao, J.; Chen, B.; Chang, N.; Chen, H.; Cheng, L.; et al. The 2021 China Report of the Lancet Countdown on Health and Climate Change: Seizing the Window of Opportunity. Lancet Public Health 2021, 6, e932-e947. [CrossRef]

89. Cai, W.; Zhang, C.; Suen, H.P.; Ai, S.; Bai, Y.; Bao, J.; Chen, B.; Cheng, L.; Cui, X.; Dai, H.; et al. The 2020 China Report of the Lancet Countdown on Health and Climate Change. Lancet Public Health 2021, 6, e64-e81. [CrossRef]

90. Evander, M.; Ahlm, C. Milder Winters in Northern Scandinavia May Contribute to Larger Outbreaks of Haemorrhagic Fever Virus. Glob. Health Action 2009, 2, 2020. [CrossRef]

91. Karmakar, M.; Pradhan, M.M. Climate Change and Public Health: A Study of Vector-Borne Diseases in Odisha, India. Nat. Hazards 2020, 102, 659-671. [CrossRef]

92. Lin, S.; Shrestha, S.; Prusinski, M.A.; White, J.L.; Lukacik, G.; Smith, M.; Lu, J.; Backenson, B. The Effects of Multiyear and Seasonal Weather Factors on Incidence of Lyme Disease and Its Vector in New York State. Sci. Total Environ. 2019, 665, 1182-1188. [CrossRef]

93. Boeckmann, M.; Joyner, T.A. Old Health Risks in New Places? An Ecological Niche Model for I. Ricinus Tick Distribution in Europe under a Changing Climate. Health Place 2014, 30, 70–77. [CrossRef]

94. Ali, S.; Gugliemini, O.; Harber, S.; Harrison, A.; Houle, L.; Ivory, J.; Kersten, S.; Khan, R.; LeBoa, C.; et al. Environmental Changes Increase Cross-Species Viral Transmission Risk. Nature 2022, 607, 555–562. [CrossRef]

95. Wilson, M. Infectious Diseases: An Ecological Perspective. BMJ 1995, 311, 1681-1684. [CrossRef]

96. Ratnam, I.; Leder, K.; Black, J.; Torresi, J. Dengue Fever and International Travel. J. Travel Med. 2013, 20, 384–393. [CrossRef]

97. Muhammad, S.; Long, X.; Salman, M. COVID-19 Pandemic and Environmental Pollution: A Blessing in Disguise? Sci. Total Environ. 2020, 728, 138820. [CrossRef] [PubMed]

98. Rosenbloom, D.; Markard, J. A COVID-19 Recovery for Climate. Science 2020, 368, 447. [CrossRef] [PubMed] [PubMed]

99. Yang, Y.; Ren, L.; Li, H.; Wang, H.; Wang, P.; Chen, L.; Yue, X.; Liao, H. Fast Climate Responses to Aerosol Emission Reductions During the COVID-19 Pandemic. Geophys. Res. Lett. 2020, 47, e2020GL089788. [CrossRef]

100. Dutheil, F.; Baker, J.; Navel, V. COVID-19 as a Factor Influencing Air Pollution? Environ. Pollut. 2020, 263, 114466. [CrossRef] [PubMed]

101. Forster, P.M.; Forster, H.I.; Evans, M.J.; Gidden, M.J.; Jones, C.D.; Keller, C.A.; Lamboll, R.D.; Quere, C.L.; Rogelj, J.; Rosen, D.; et al. Current and Future Global Climate Impacts Resulting from COVID-19. Nat. Clim. Chang. 2020, 10, 913–919. [CrossRef]

102. Castaneda-Gomez, J.; Gonzalez-Acosta, C.; Jaime-Rodriguez, J.L.; Villegas-Trejo, A.; Moreno-Garcia, M. COVID-19 Pandemic and Environmental Pollution: A Blessing in Disguise? Gac. Med. Mex. 2021, 157, 194–200. [CrossRef]

103. Olive, M.-M.; Baldet, T.; Devillers, J.; Fite, J.; Paty, M.-C.; Paupy, C.; Quenel, P.; Quillery, E.; Raude, J.; Stahl, J.-P.; et al. The COVID-19 Pandemic Should Not Derailed Global Vector Control Efforts. PLoS Negl. Trop. Dis. 2020, 14, e0008716. [CrossRef]

104. Seelig, F.; Bezerra, H.; Cameron, M.; Hii, J.; Hiscox, A.; Irish, S.; Jones, R.T.; Lang, T.; Lindsay, S.W.; Lowe, R.; et al. The COVID-19 Pandemic Should Not Derailed Global Vector Control Efforts. PLoS Negl. Trop. Dis. 2020, 14, e0008606. [CrossRef]

105. Webb, C.E. Reflections on a Highly Unusual Summer: Bushfires, COVID-19 and Mosquito-Borne Disease in NSW, Australia. Public Health Res. Pract. 2020, 30, e0342027. [CrossRef]

106. Wilder-Smith, A.; Tissera, H.; Ooi, E.E.; Coloma, J.; Scott, T.W.; Gabul, D.J. Preventing Dengue Epidemics during the COVID-19 Pandemic. Am. J. Trop. Med. Hyg. 2020, 103, 570–571. [CrossRef] [PubMed]

107. Khan, S.A.; Webb, C.E.; Abu Kassim, N.F. Prioritizing Mosquito-Borne Diseases during and after the COVID-19 Pandemic. West. Pac. Surveill. Response J. 2021, 12, 40–41. [CrossRef] [PubMed]

108. Reegan, A.D.; Gandhi, M.R.; Asharaja, A.C.; Devi, C.; Shanthakumar, S.P. COVID-19 Lockdown: Impact Assessment on Aedes Larval Indices, Breeding Habitats, Effects on Vector Control Programme and Prevention of Dengue Outbreaks. Heliyon 2020, 6, e05181. [CrossRef] [PubMed]

109. Zui, M.; Rigatelli, G.; Roncon, L. Reduction of West Nile Virus Infections in Italy during 2020 Early Summer: A Secondary “COVID-19” Effect? Pathog. Glob. Health 2020, 114, 345–346. [CrossRef]

110. Lim, J.T.; Dickens, B.S.L.; Chew, L.Z.X.; Choo, E.L.W.; Koo, J.R.; Aik, J.; Ng, L.C.; Cook, A.R. Impact of Sars-Cov-2 Interventions on Dengue Transmission. PLoS Negl. Trop. Dis. 2020, 14, e0008719. [CrossRef]

111. McCormick, D.W.; Kugeler, K.J.; Marx, G.E.; Jayanthi, P.; Dietz, S.; Mead, P.; Hinckley, A.F. Effects of COVID-19 Pandemic on Reported Lyme Disease, United States, 2020. Emerg. Infect. Dis. 2021, 27, 2715–2717. [CrossRef]

112. Ullrich, A.; Schranz, M.; Rexroth, U.; Hamouda, O.; Schaade, L.; Diercke, M.; Boender, T.S. Impact of the COVID-19 Pandemic and Associated Non-Pharmaceutical Interventions on Other Notifiable Infectious Diseases in Germany: An Analysis of National Surveillance Data during Week 1-2016—Week 32-2020. Lancet Reg. Health-Eur. 2021, 6, 100103. [CrossRef]

113. Lai, C.-C.; Chen, S.-Y.; Yen, M.-Y.; Lee, P.-I.; Ko, W.-C.; Hsueh, P.-R. The Impact of the Coronavirus Disease 2019 Epidemic on Notifiable Infectious Diseases in Taiwan: A Database Analysis. Travel Med. Infect. Dis. 2021, 40, 101997. [CrossRef]