

Climate factors, climate change, and their health impacts

Ania Kawiecki Peralta (ania.kawiecki@bsc.es)

Carles Milà Garcia (carles.milagarcia@bsc.es)

Global Health Resilience group

Advanced Webinar Series on Spatiotemporal Modeling of Climate-Sensitive Diseases
21st of January 2026

About us



**Barcelona
Supercomputing
Center**
Centro Nacional de Supercomputación

Ania Kawiecki Peralta



I am a postdoc at the Global Health Resilience group at the BSC. My background is in Veterinary Medicine, and I have a PhD in Epidemiology studying dengue virus vector surveillance and control.

Currently I am working on developing R packages to facilitate disease risk modeling and prediction using Bayesian spatio-temporal models in INLA.

Carles Milà



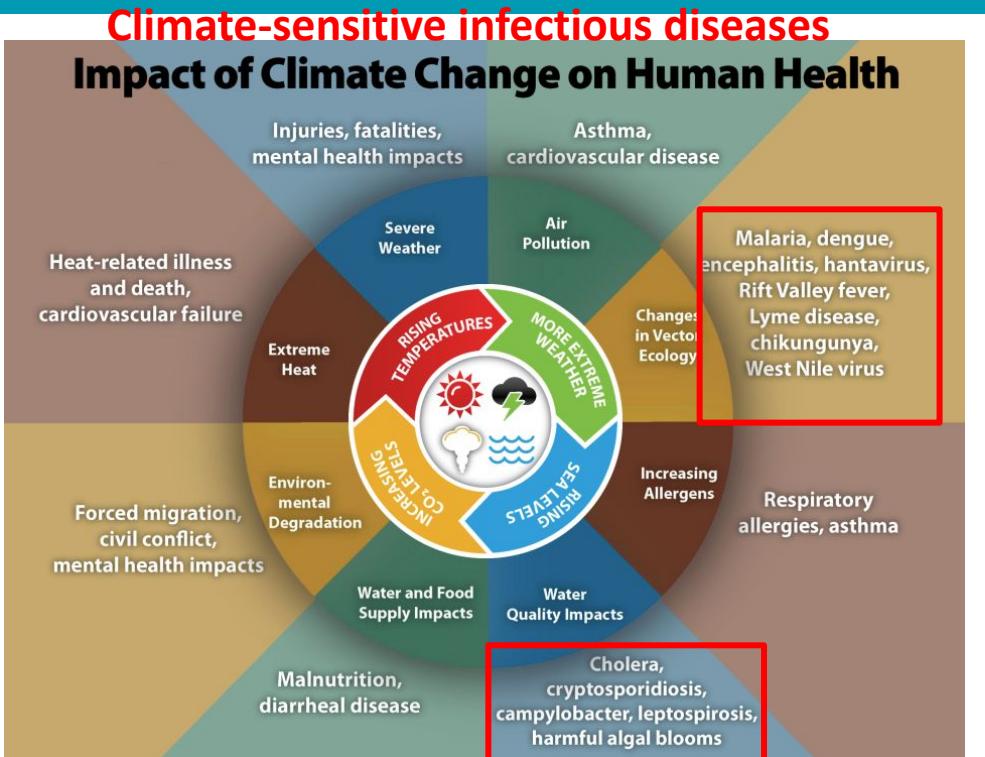
I am a data scientist at the Global Health Resilience group at the BSC. My background is in statistics and geoinformatics, and I have a PhD in spatial modelling for exposure assessment.

I am currently working on developing R packages for climate-sensitive data processing and modelling.

1. Climate sensitive-diseases.
2. Health impact of climatic variables and climate change.
3. Climate-informed early warning systems.
4. Data structure, pre-processing and exploration.
5. Questions at the end!

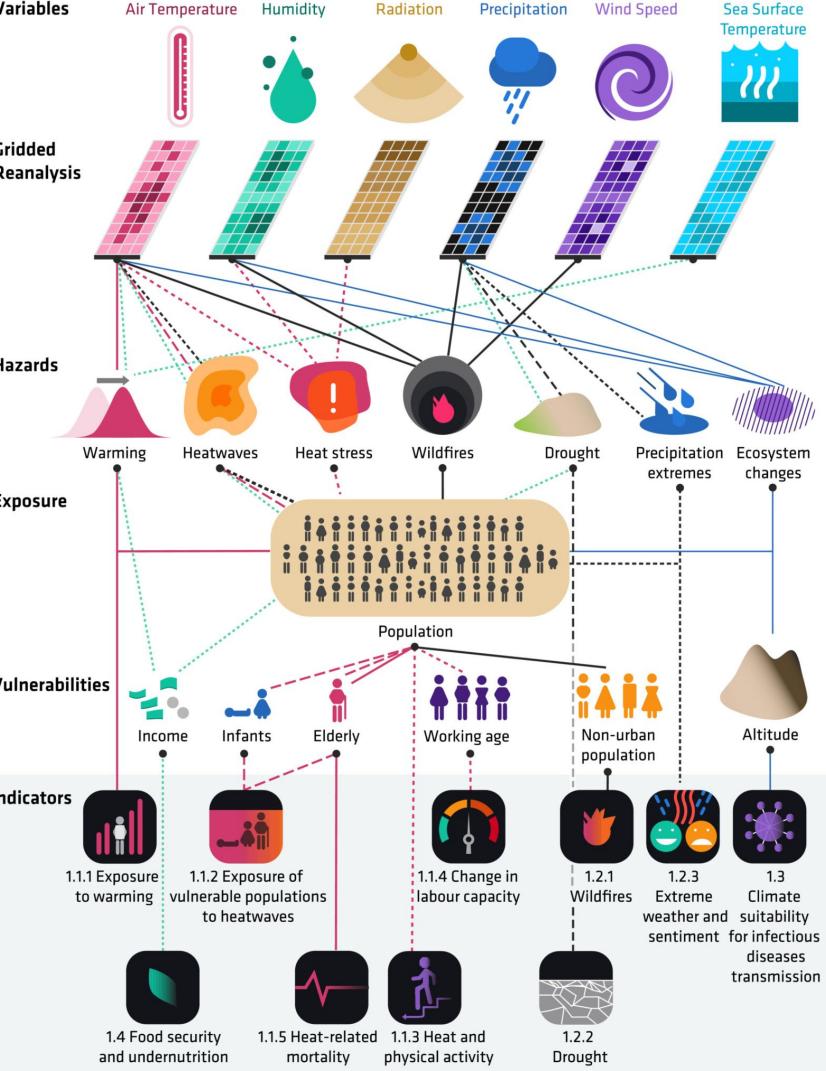
Climate-sensitive diseases

Health impacts of climate change

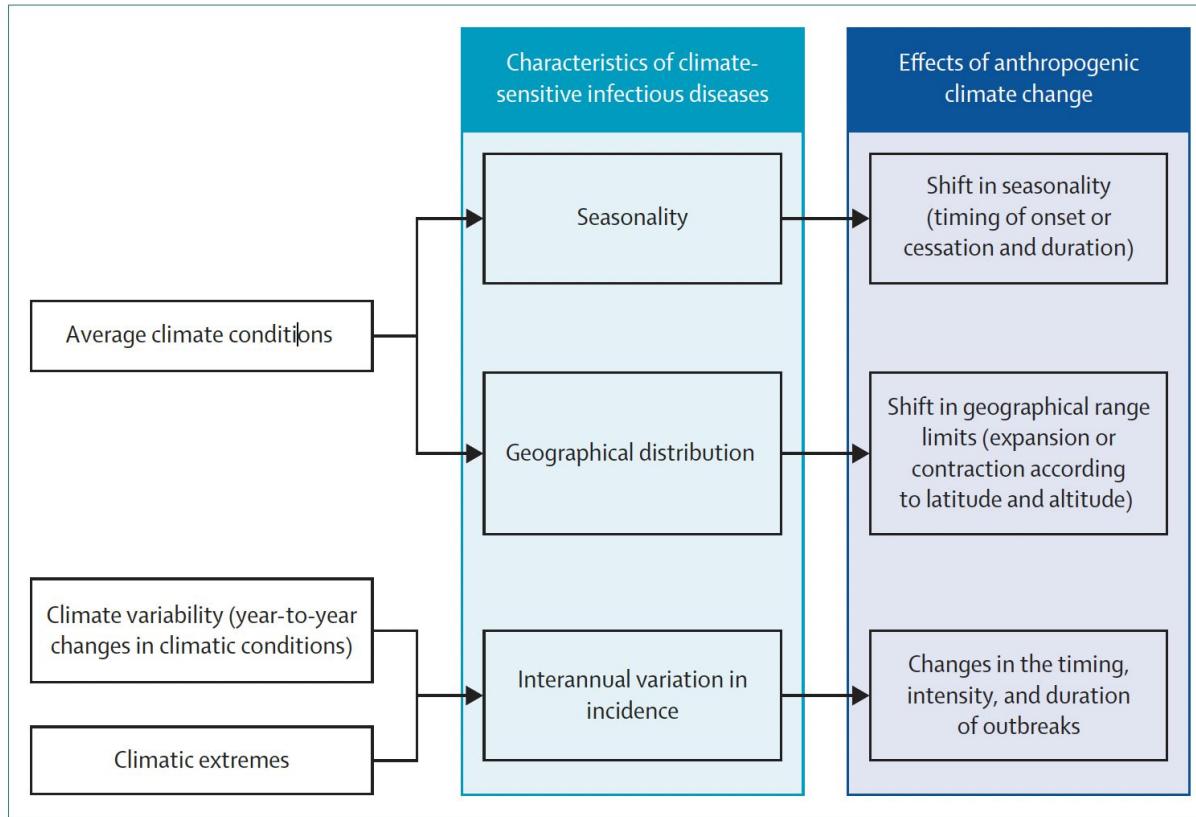


George Luber adapted from Patz *et al.* *et Nature* 2005

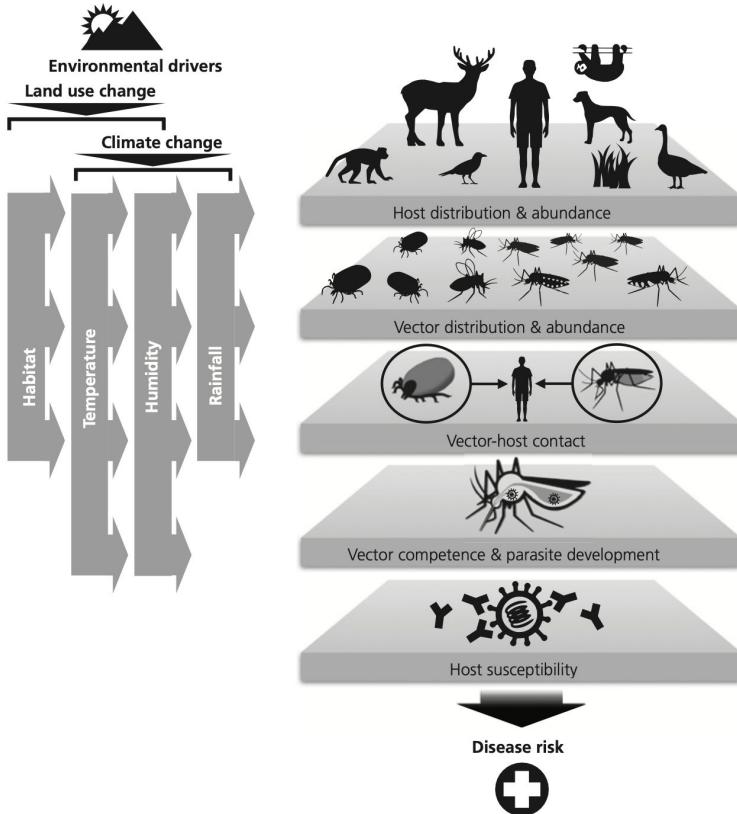
Di Napoli *et al.* *Meteorological Applications* 2022



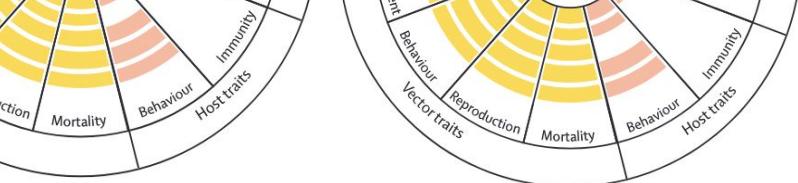
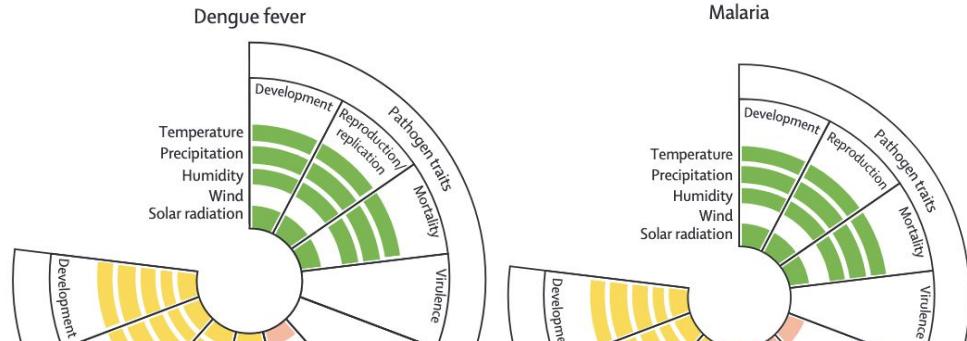
Climate-sensitive infectious diseases



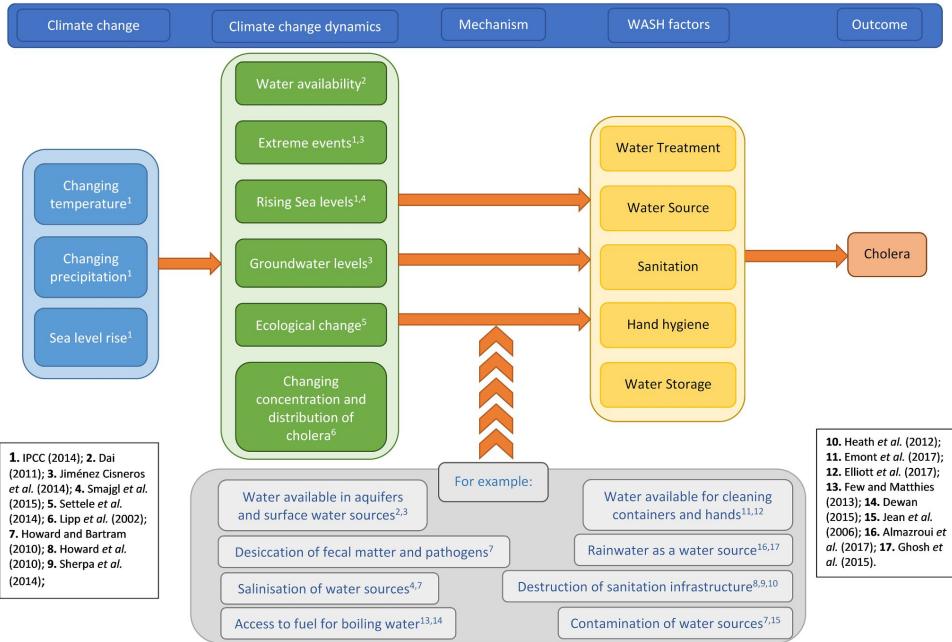
Climate-sensitive infectious diseases - vector-borne diseases



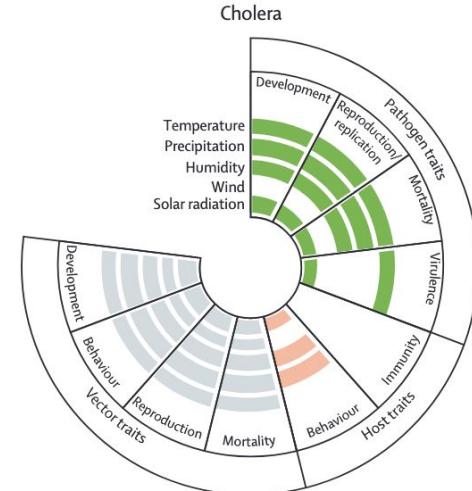
- Evidence exists for the climate-pathogen trait association
- Evidence exists for the climate-host trait association
- Evidence exists for the climate-vector trait association
- The trait is not relevant for the disease
- Limited, insufficient, or unavailable evidence



Climate-sensitive infectious diseases - water-borne diseases



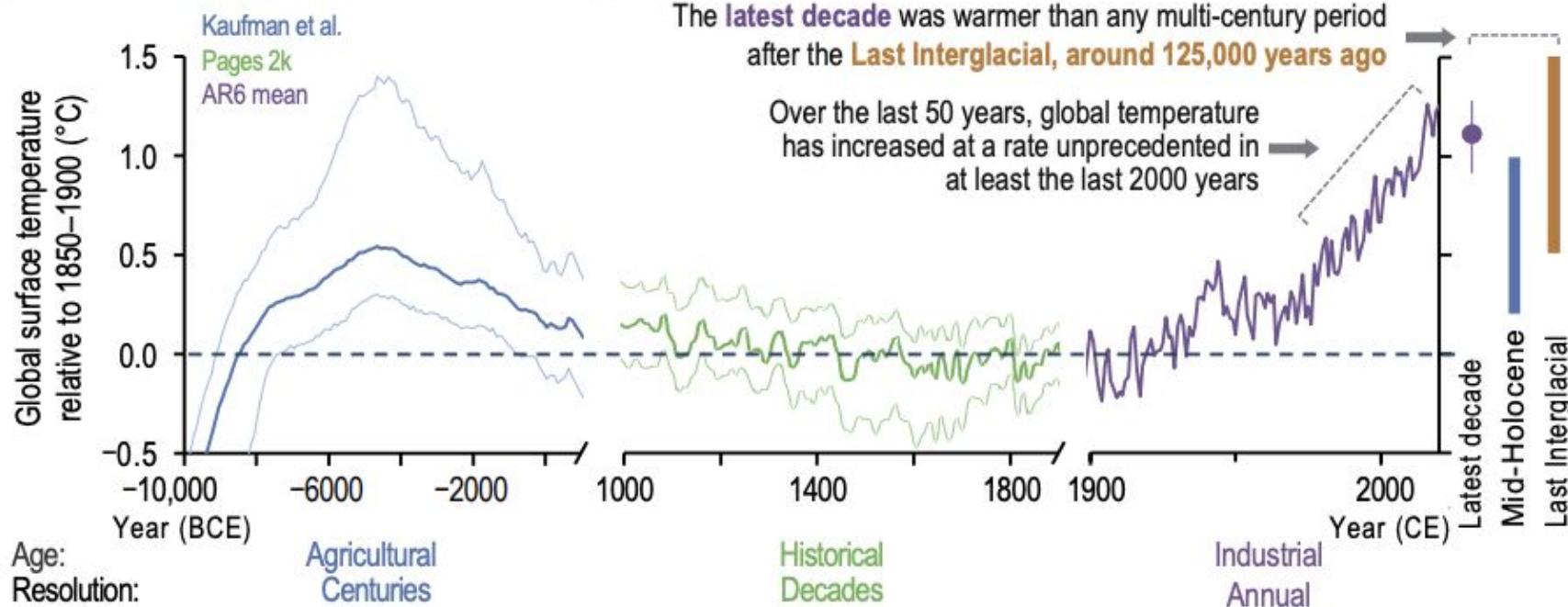
- Evidence exists for the climate-pathogen trait association (Green)
- Evidence exists for the climate-host trait association (Orange)
- Evidence exists for the climate-vector trait association (Yellow)
- The trait is not relevant for the disease (Grey)
- Limited, insufficient, or unavailable evidence (White)



Health impact of climatic variables and climate change

Changes in surface temperature

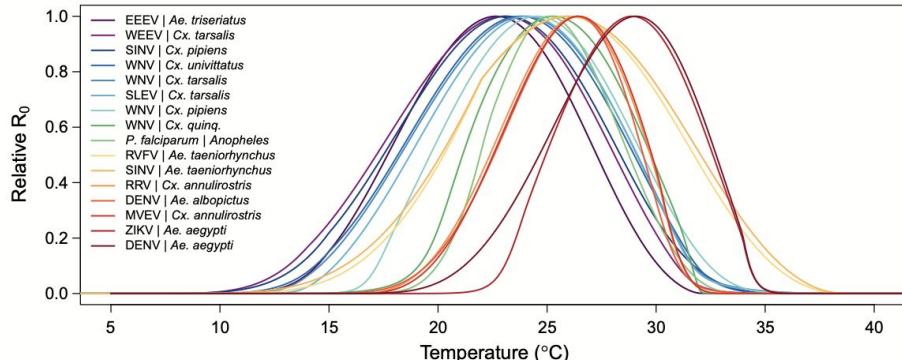
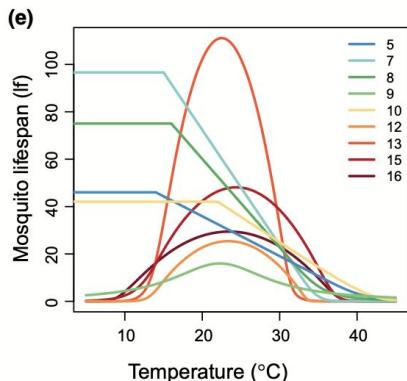
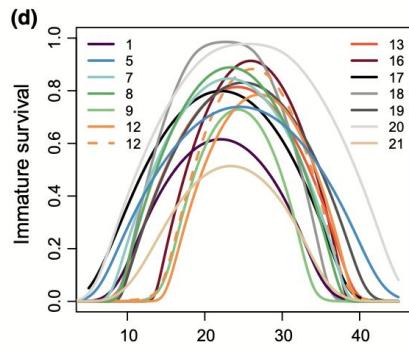
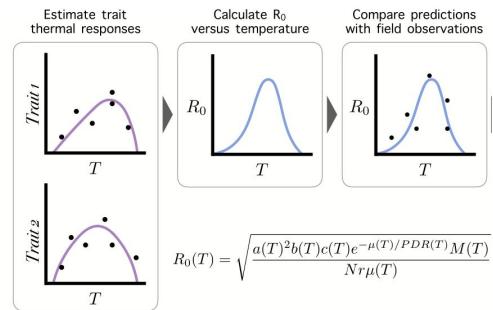
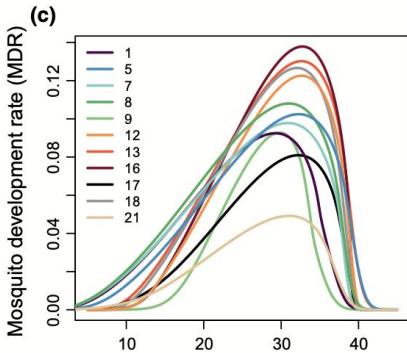
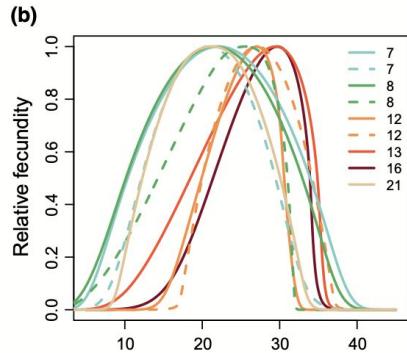
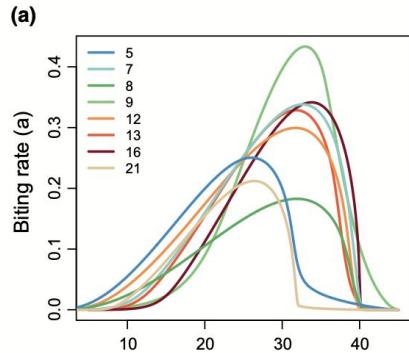
(a) Global surface temperatures are more likely than not unprecedented in the past 125,000 years



Temperature



Effect on disease transmission-relevant vector traits



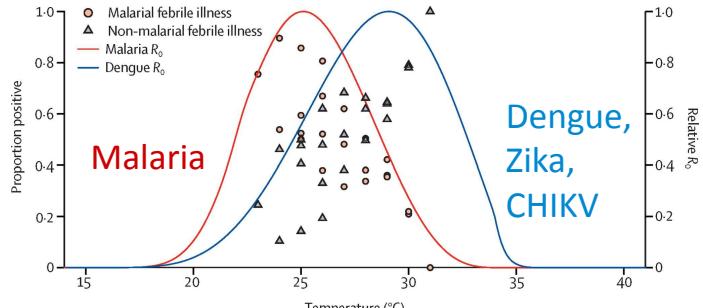
Temperature

Shifts in geographical distribution of disease

Observed

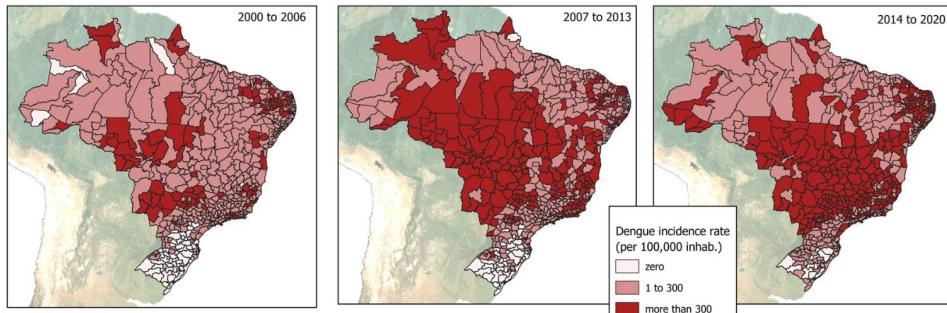
Projected

Kenyan children from 2014 to 2018



Mordecai *et al.* *Lancet Planetary Health* 2020

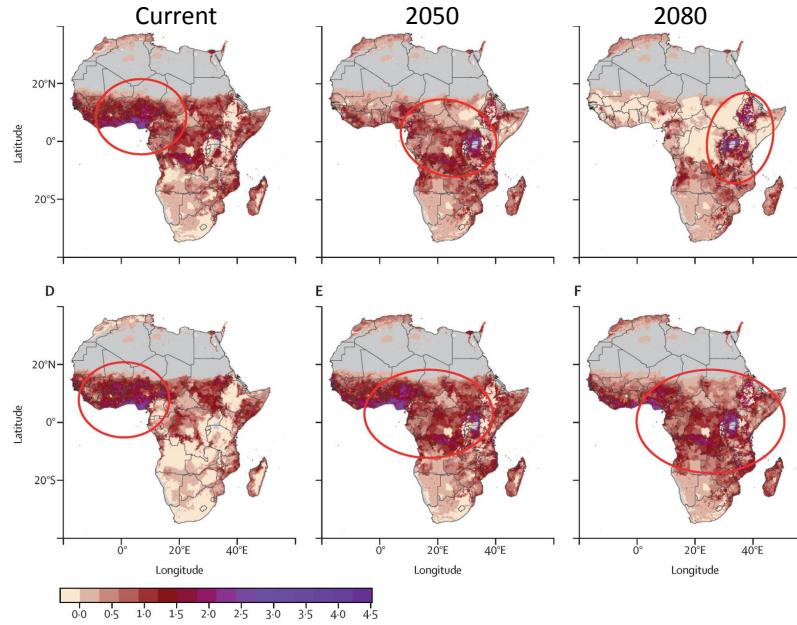
Brasil - dengue incidence



Barcellos *et al.* *Nature Scientific Reports* 2024

Malaria

Dengue, Zika, CHIKV

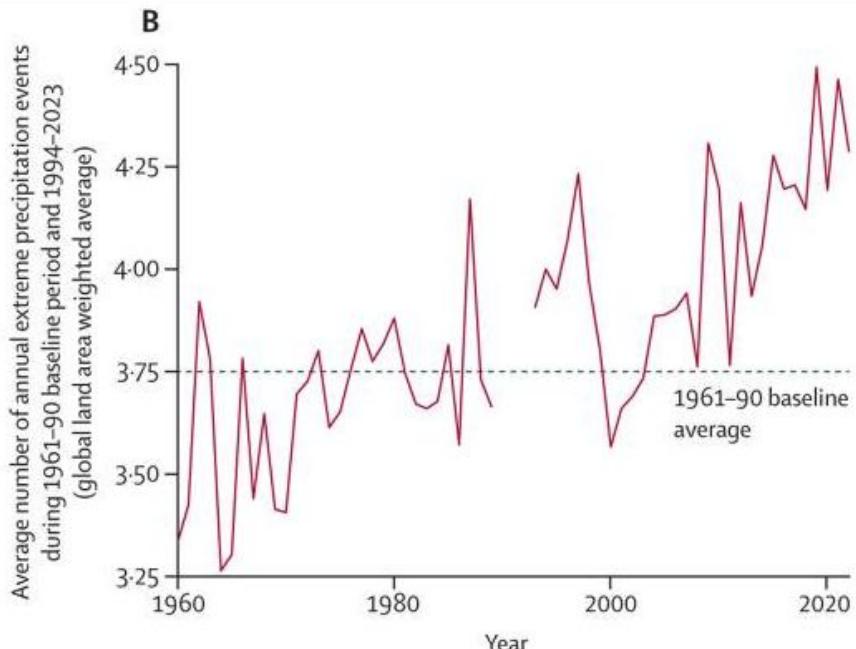


Mordecai *et al.* *Lancet Planetary Health* 2020

Extreme precipitation and drought

Extreme precipitation events over time

Average number of annual extreme precipitation events per 79 km²
average land area in baseline years (1961–90) and during the most recent
30-year period (1994–2023).

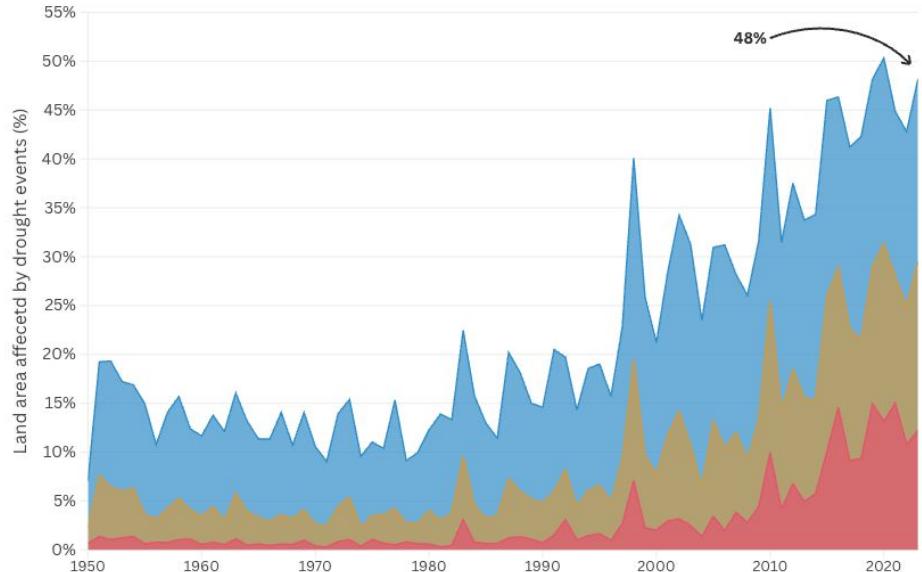


Romanello *et al.* *Lancet* 2024

Land Affected by Extreme Droughts

Percentage of global land area affected by one, three, six or twelve months of extreme drought per year, from 1950–2023

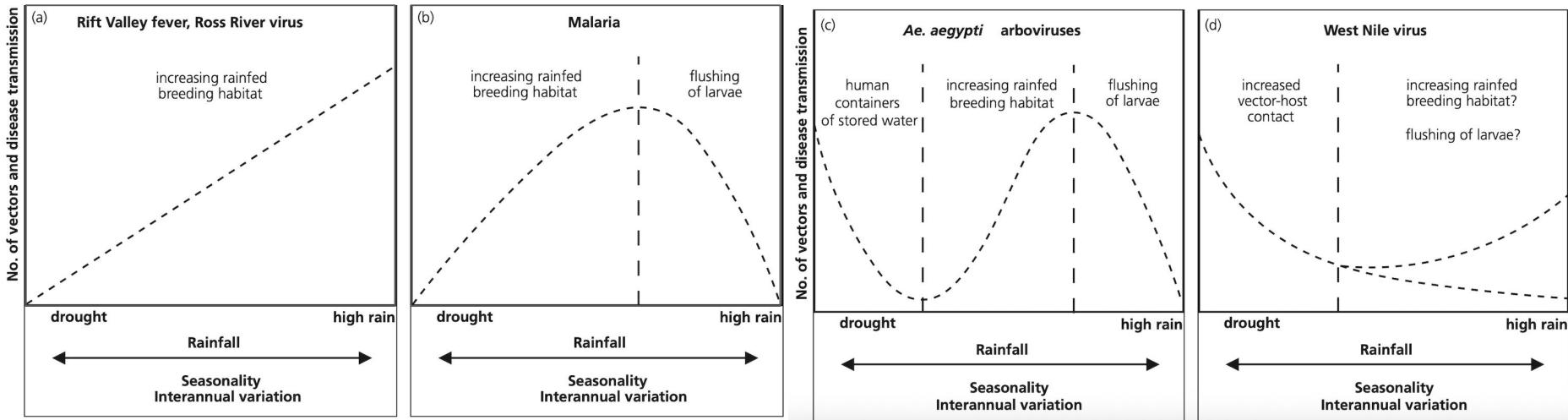
Months of Drought:  1 month  3 months  6 months



Please reference the 2024 Report of the Lancet Countdown if using this data.
For a full description of the indicator, see the 2024 report of the Lancet Countdown at lancetcountdown.org

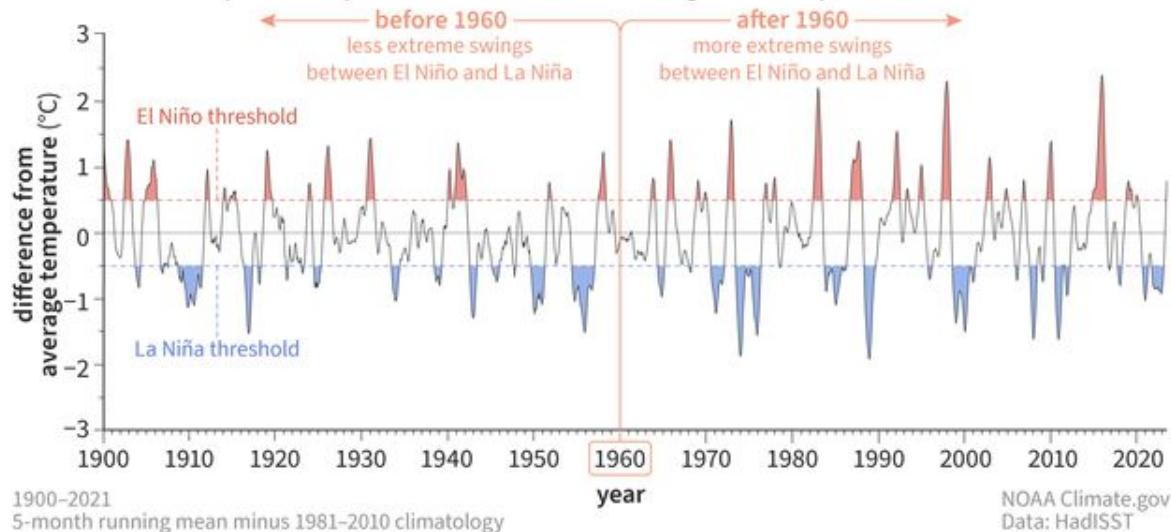
Extreme precipitation and drought

Effect on vector-borne diseases



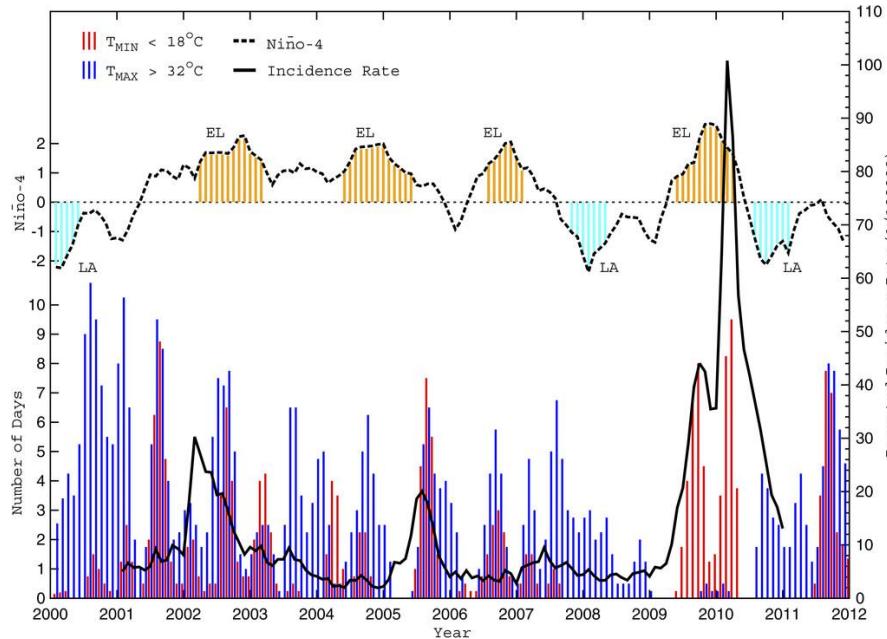
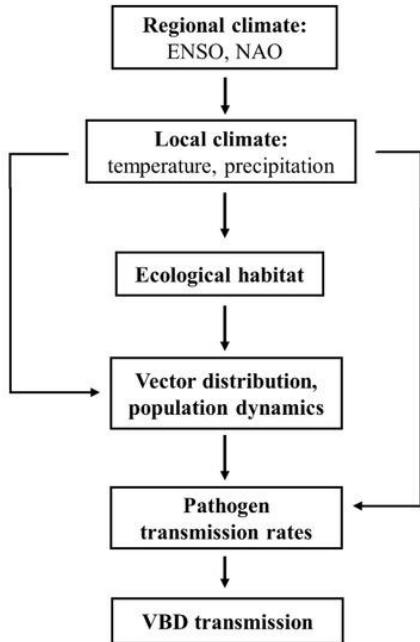
Climate change is expected to influence ENSO by changing ocean–atmosphere interactions in the tropical Pacific, potentially affecting its intensity, rainfall patterns, and global impacts, but there is still considerable uncertainty about how ENSO frequency and strength will change in the future (IPCC AR6 WGI, 2021)

Sea surface temperature patterns in the Niño-3.4 region of tropical Pacific



Oceanic indices

ENSO (El Niño Southern Oscillation) and **IOD** (Indian Ocean Dipole) are potential drivers of interannual variation in climate-sensitive infectious disease patterns.



In this study in Cali, Colombia, dengue incidence often peaked 4–6 months after an El Niño event associated with above-average local temperatures and below-average local rainfall and humidity.

Non-linear, lagged and interacting effects

Air Temperature



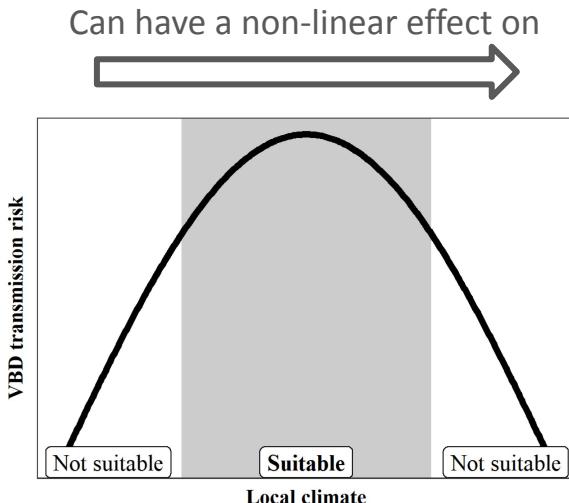
Precipitation



Humidity



Non-linear effects



Ma *et al.*, *Biology* 2022

Pathogen traits (e.g. viruses, bacteria and parasites):

- Replication and development rates
- Survival and mortality

Ectothermic vector traits (e.g. mosquitoes, ticks, sandflies, midges, blackflies, and snails):

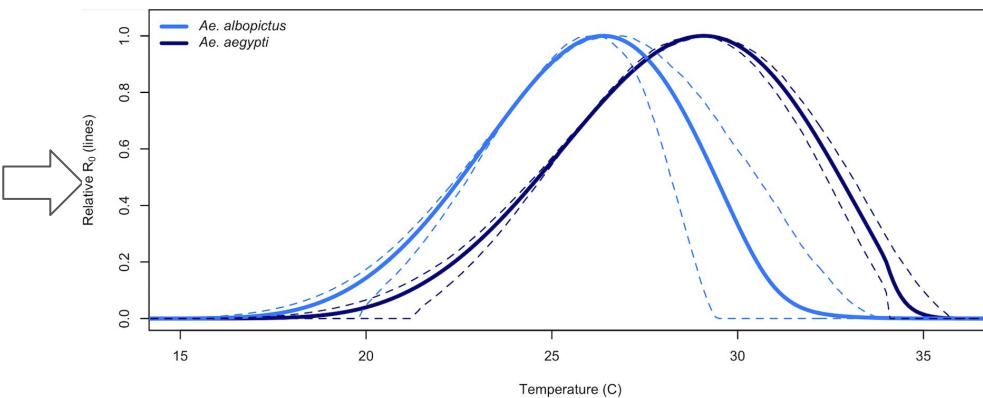
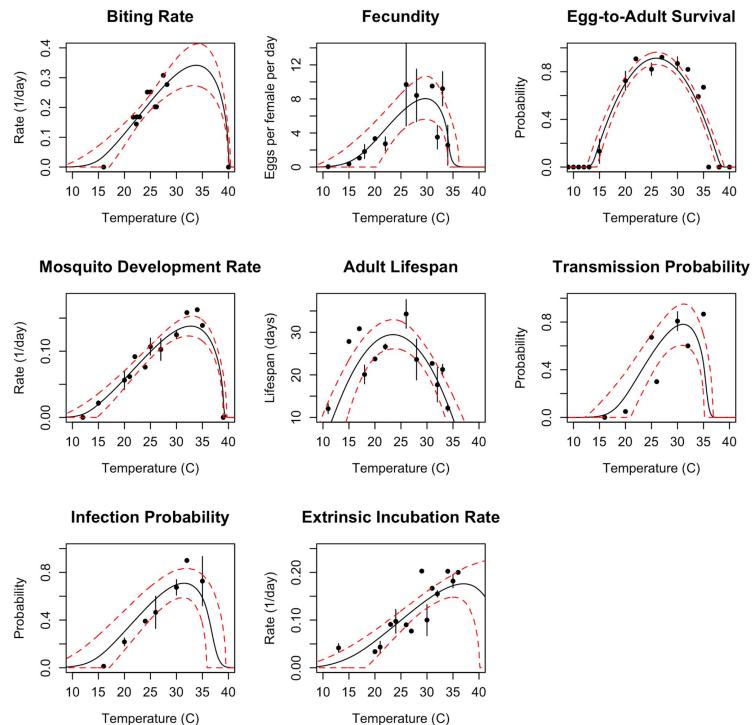
- Reproduction and development rates
- Survival and mortality
- Biting rate
- Vector competence (the probability of becoming infectious after biting an infectious host)

Non-linear, lagged and interacting effects

Non-linear effects

The effect of temperature on dengue virus transmission is non-linear.

Thermal responses of *Ae. aegypti* and DENV traits that drive transmission



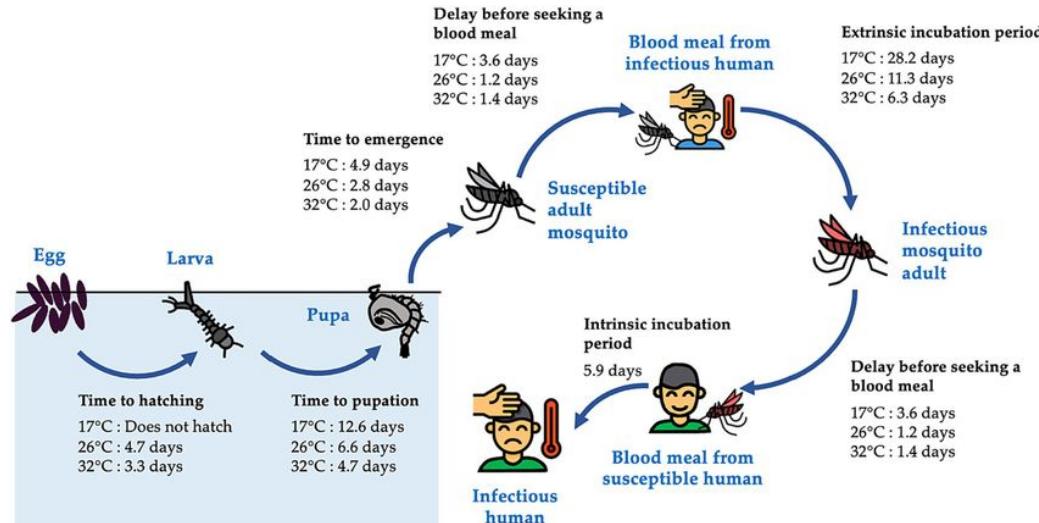
Ae. aegypti transmission peaks at 29.1°C.

Lagged effects

Climate-sensitive infectious diseases often have a delayed response to climate drivers. Delays might be due to:

- vector or non-human host lifecycles
- the extrinsic and intrinsic incubation period of the pathogen
- the time between symptom onset and health-seeking behaviour.

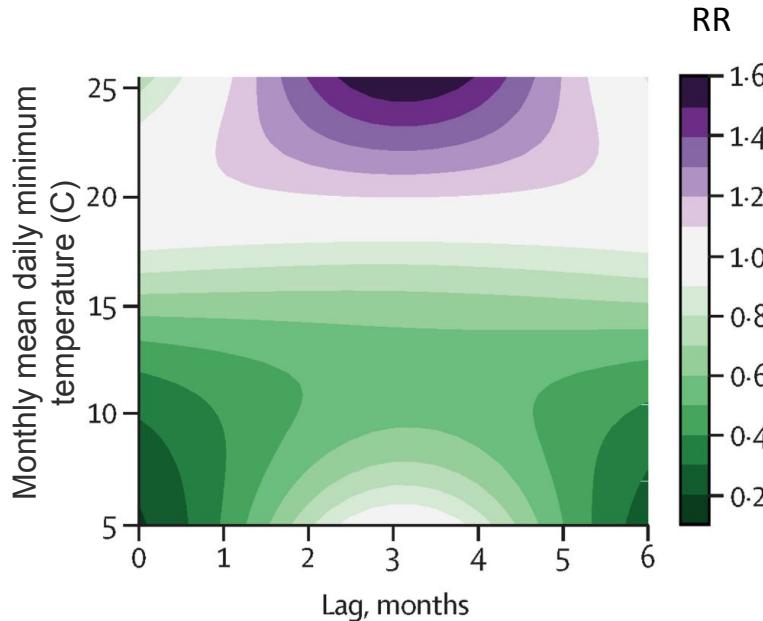
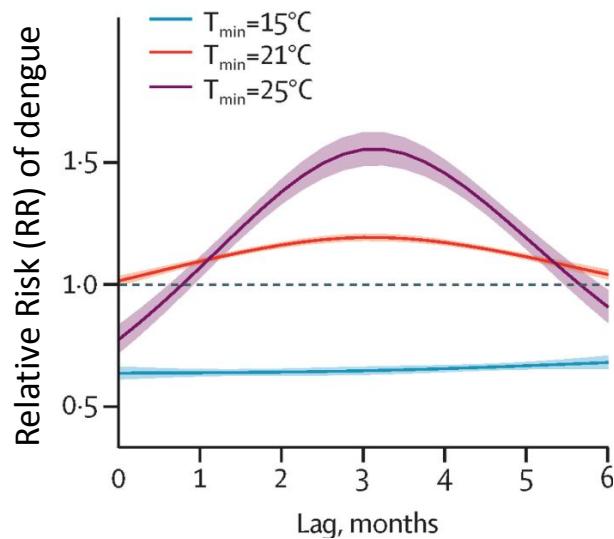
Life stages of *Aedes* mosquitoes and transmission of dengue virus



Lagged effects

The effect of temperature on dengue virus transmission is non-linear and lagged.

In this study in Brazil the greatest relative risk (RR) of dengue was found at T_{\min} of 25.5°C at a lag of 2–4 months.



Interacting effects

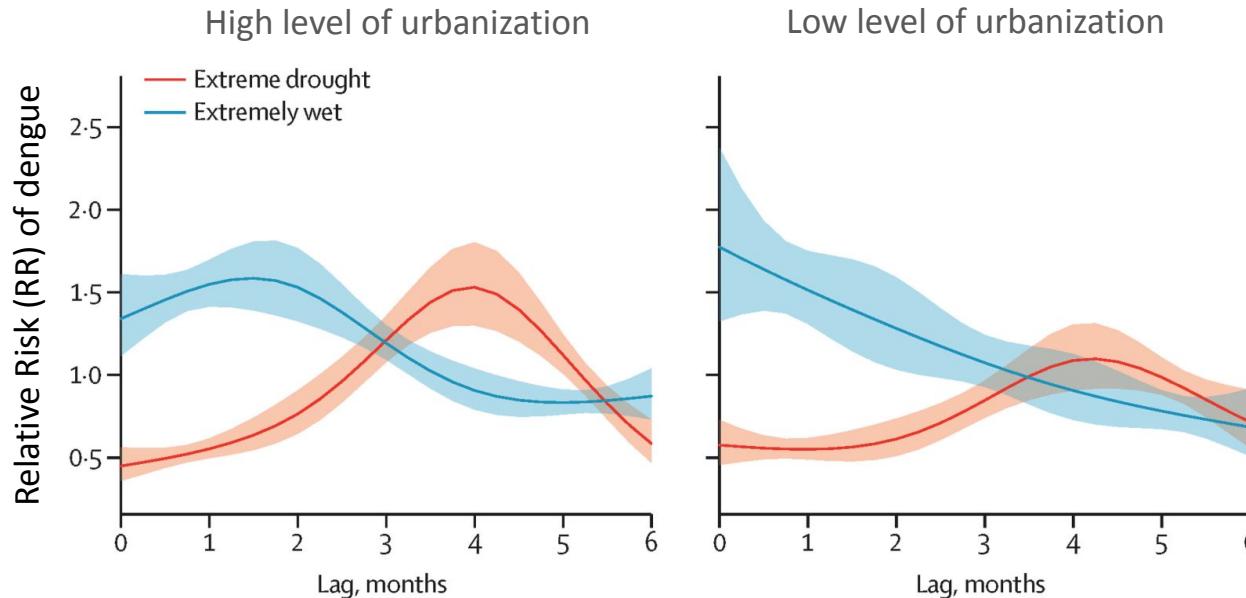
Non-climatic factors can modulate disease transmission:

- Habitat degradation and land-use change:
 - influence the density and contact among reservoirs, vectors, and humans
 - modify local climatic and environmental conditions.
- Urbanisation and human infrastructure (e.g. sanitation and water systems)
- Population demographics and socio-economic factors: can influence susceptibility and vulnerability to disease.
- Behavioural factors (such as clothing, use of insect repellent, time spent outdoors)



Interacting effects

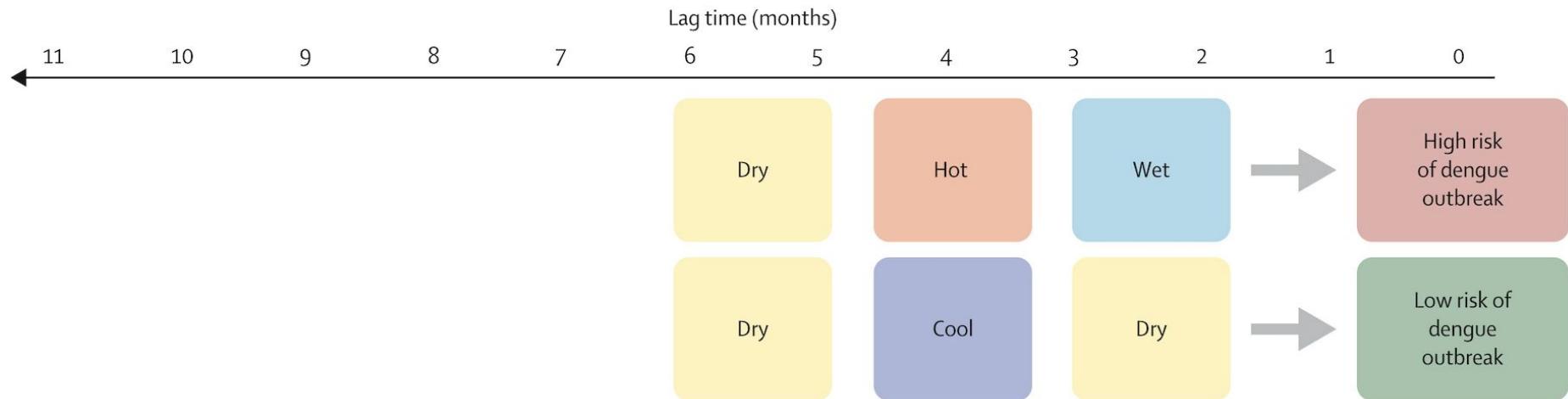
In this study in Brazil there was an interacting effect between urbanization level and extreme values of PDSI: the greater the level of urbanisation, the higher the risk of dengue after extreme drought, while at lower levels of urbanisation the relative risk of dengue after extremely wet conditions was greater and more immediate.



Non-linear, lagged and interacting effects

Climatic factors can interact with each other at short and long lags leading to a compound effect on disease risk. Excessive rainfall following a period of abnormally low rainfall or drought is important for outbreaks of cholera, diarrheal diseases, and dengue.

In this study in Barbados, long-lag dry (lagged by 5 months), mid-lag hot (lagged by 3 months), and short-lag wet (lagged by 1 month) conditions led to the greatest dengue risk



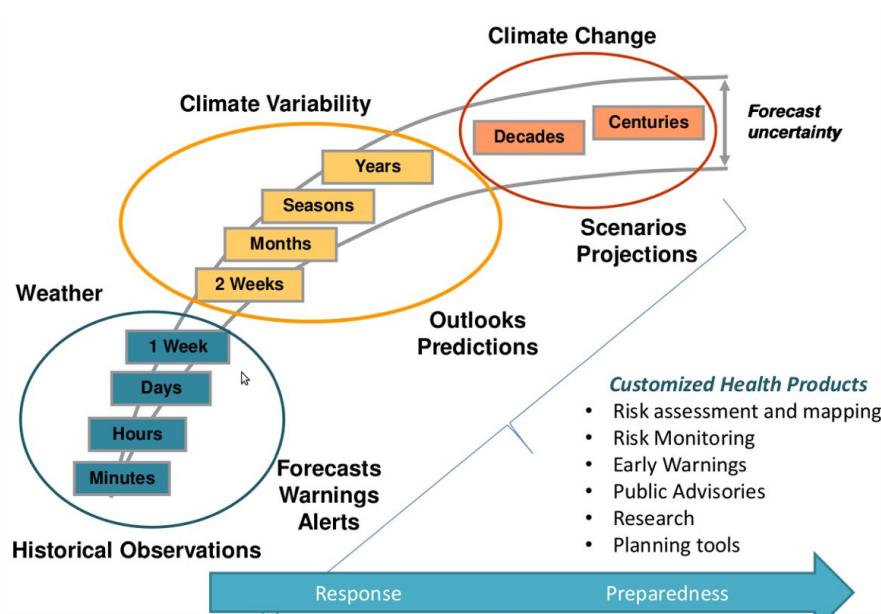
Climate variables associated with infectious diseases

CSID	Climate/weather trigger	Confidence	References
Cholera	Abrupt and heavy/extreme rainfall, extensive flooding	High agreement, medium evidence	De Magny et al. 2012; Passeto et al. 2018; Sinyange et al 2018; Cann et al 2013
	Higher temperatures/heatwave (one study reported drier conditions being a risk modifier, whilst the other reported rainfall was a significant risk modifier)	Medium agreement, low evidence	Charnley et al. 2021; Wu et al. 2018
Diarrheal diseases	Heavy and extended rainfall (torrential) and flooding	High agreement, high evidence	Ding et al 2013; Chen et al 2012; Thompson et al. 2014; Kim et al. 2013; Deng et al 2015; Jones et al 2016; CDC 1983; Wade et al. 2003; Saulnier et al. 2017
	High temperatures	High agreement, Medium evidence	Kraay et al. 2020; Levy et al. 2016; Mertens et al. 2019; Thompson et al. 2014; Xu et al. 2014
	Heavier and longer duration of flood, than a flash flood	Low evidence	Ding et al 2013
	Excessive rainfall especially when following a dry period or period of low rainfall	High agreement, Medium evidence	Despande et al 2020; Levy et al 2016; Carlton et al. 2014; Mertens et al 2019
Malaria	Amount and duration of heavy rainfall, leading to extensive flooding	High agreement, high evidence	Boyce et al. 2018; Maes et al. 2014; Ding et al. 2014; Okaka et al. 2018; Himeidan et al. 2007
Dengue	Extreme heavy rain	High agreement, medium evidence	Chen et al. 2012; Lowe et al. 2018; Lowe et al. 2021; Cheng et al. 2021
	Above normal temperatures or drought followed by extremely high rainfall, 1-3 months lag time	High agreement, medium evidence	Lowe et al. 2018; Lowe et al. 2021; Cheng et al. 2021
	Drought conditions, especially with increased temperatures	High agreement, medium evidence	Stanke et al 2013; Ayambe et al. 2014; Lowe et al. 2018; Lowe et al. 2021
	Heatwaves delay timing and increase magnitude of dengue outbreaks, up to a point as longer duration of heatwaves (number of heatwave days) may reduce dengue outbreak risk	Medium agreement, medium evidence	Cheng et al 2020; Cheng et al. 2021; Seah et al. 2021

Climate-informed early warning systems

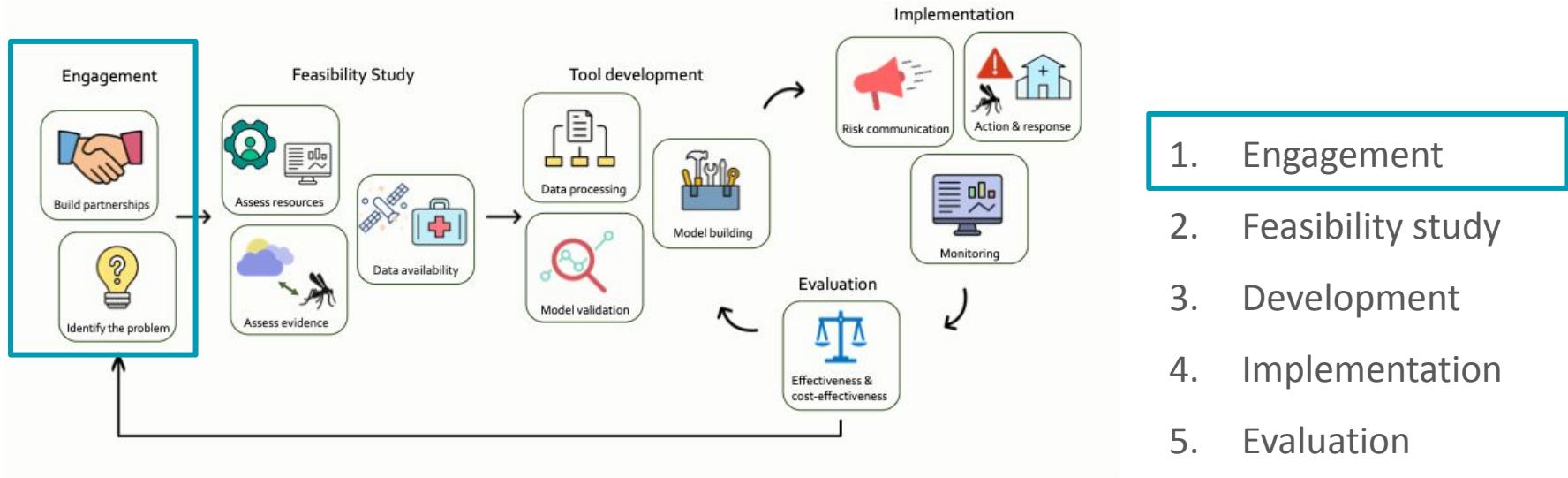
Climate-informed early warning systems: lead times

Climate-informed early warning systems integrate climate data to predict the risk of an infectious disease outbreak.



For vector-borne diseases,
a typical lead time ranges
from 1 to 6 months

Stages involved in the co-creation of an early warning system for vector-borne disease

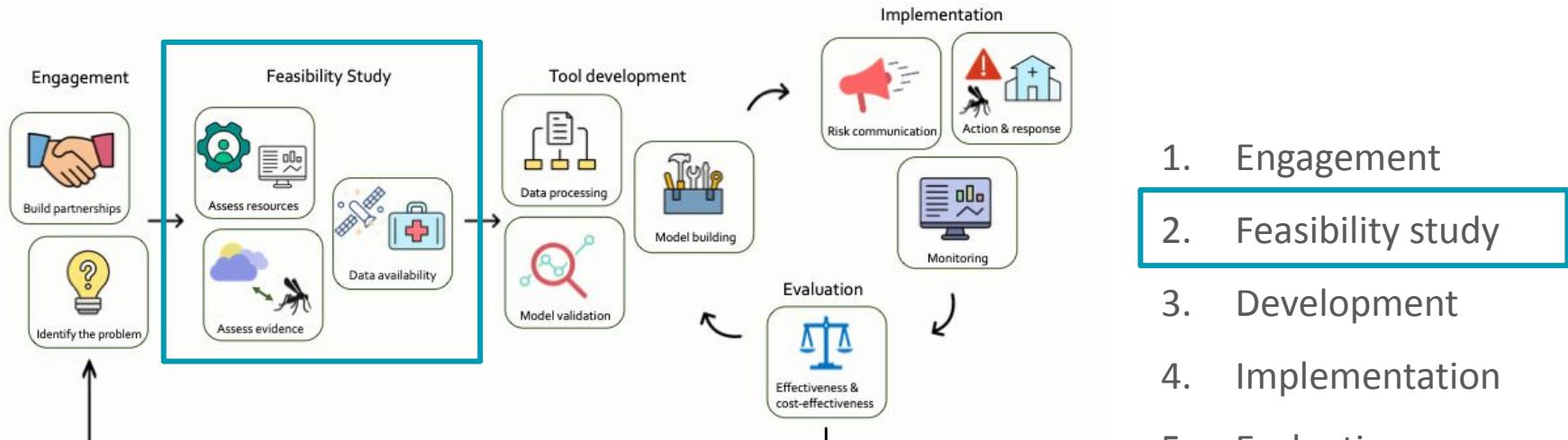


CHAPTER 12

Early warning systems for vector-borne diseases:
engagement, methods and implementation

Emilie Finch^{1,2*}, Martin Lotto Batista^{3,4}, Tilly Alcayna^{1,2,5}, Sophie A. Lee^{1,2},
Isabel K. Fletcher⁶ and Rachel Lowe^{1,2,4,7*}

Stages involved in the co-creation of an early warning system for vector-borne diseases

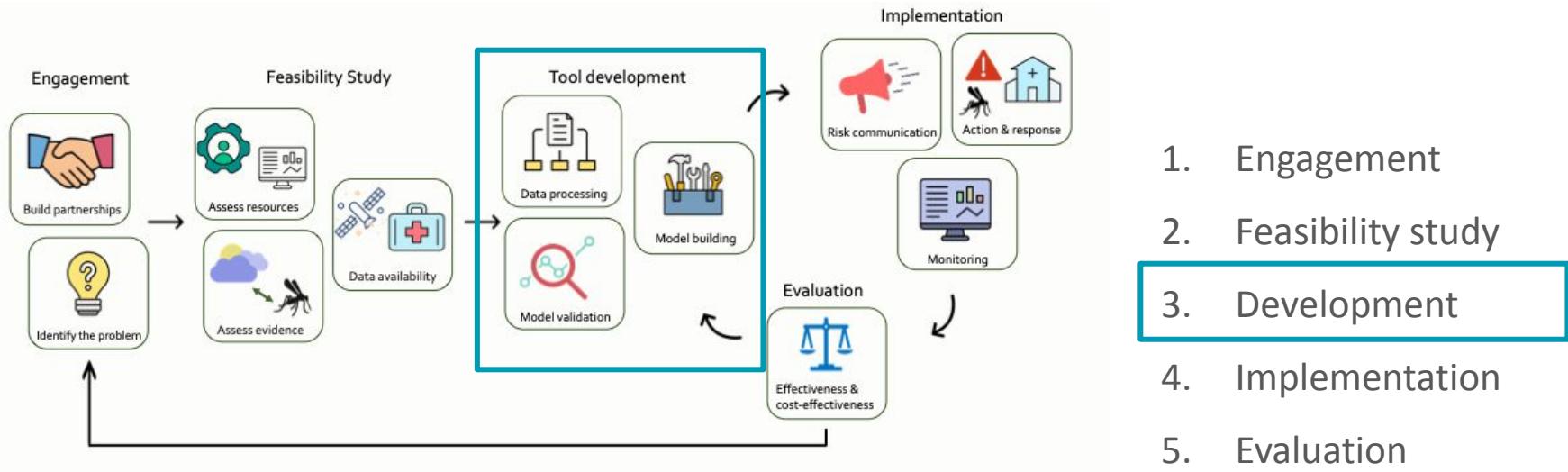


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Statistical

- Characterise the relationship between outcome and exposure under a set of assumptions.
- Require long time series.
- Can produce reliable uncertainty estimates.
- Examples: hierarchical models, ARIMA models.

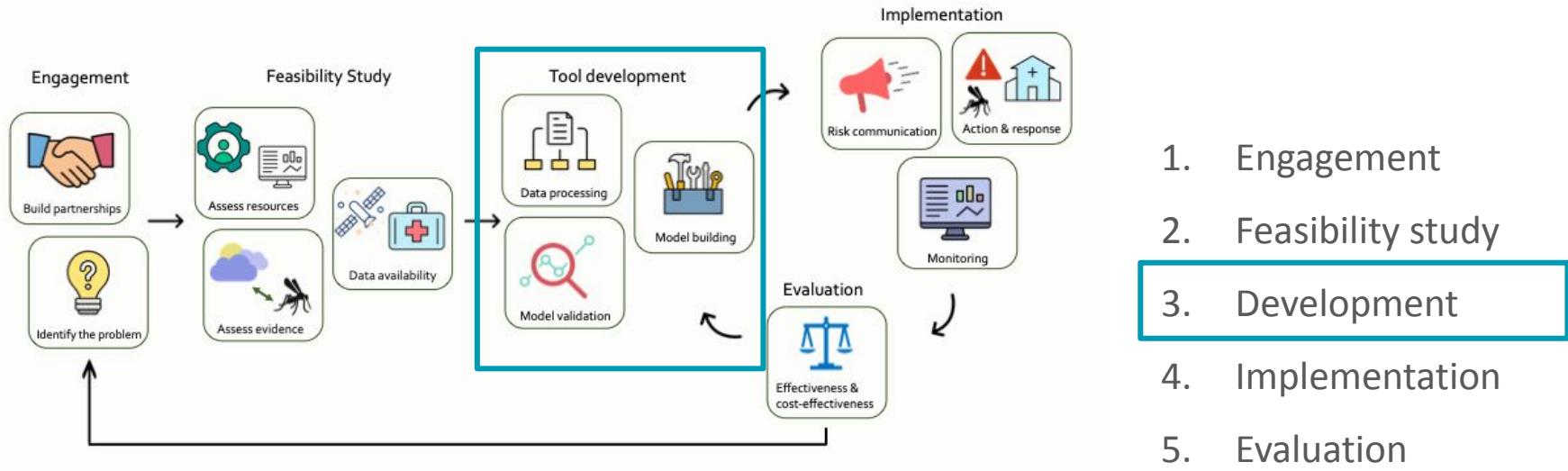
Machine learning

- More flexible than statistical models, can achieve better results but require more data.
- More difficult interpretation and uncertainty quantification.
- Examples: neural networks, random forest.

Mechanistic

- Capture the underlying biological processes, population immunity, contact patterns, population mobility and control measures.
- Require deep understanding and many assumptions, not always feasible/realistic.
- Examples: compartmental model, agent-based models

Stages involved in the co-creation of an early warning system for vector-borne disease

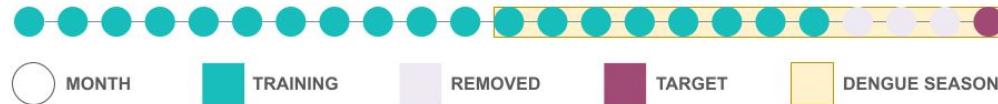


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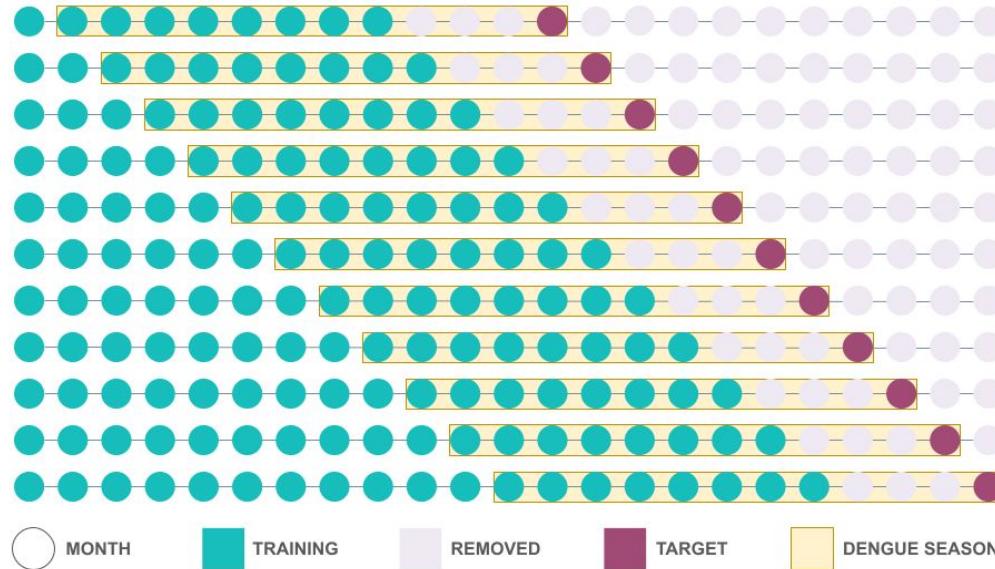
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Expanding window

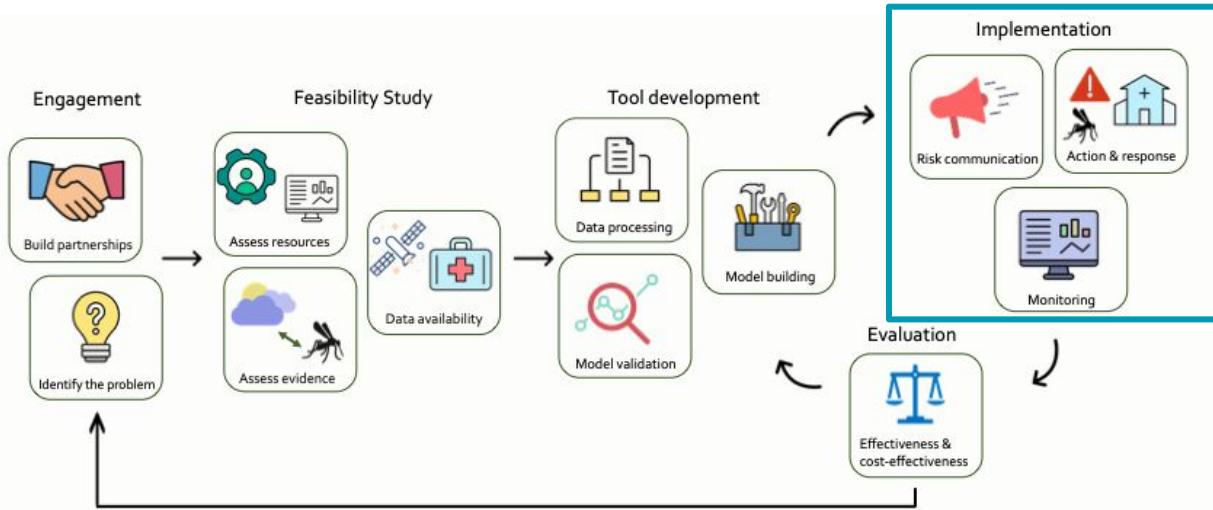


Expanding window cross-validation



Early warning systems: implementation

Stages involved in the co-creation of an early warning system for vector-borne disease



1. Engagement
2. Feasibility study
3. Development
- 4. Implementation**
5. Evaluation

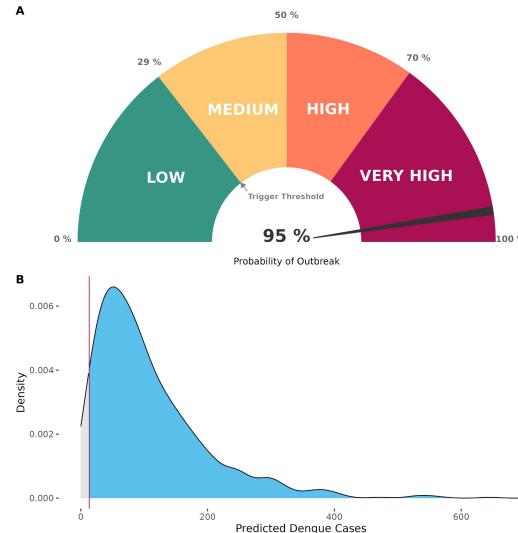
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Early warning systems: implementation

Barbados - Dengue forecast



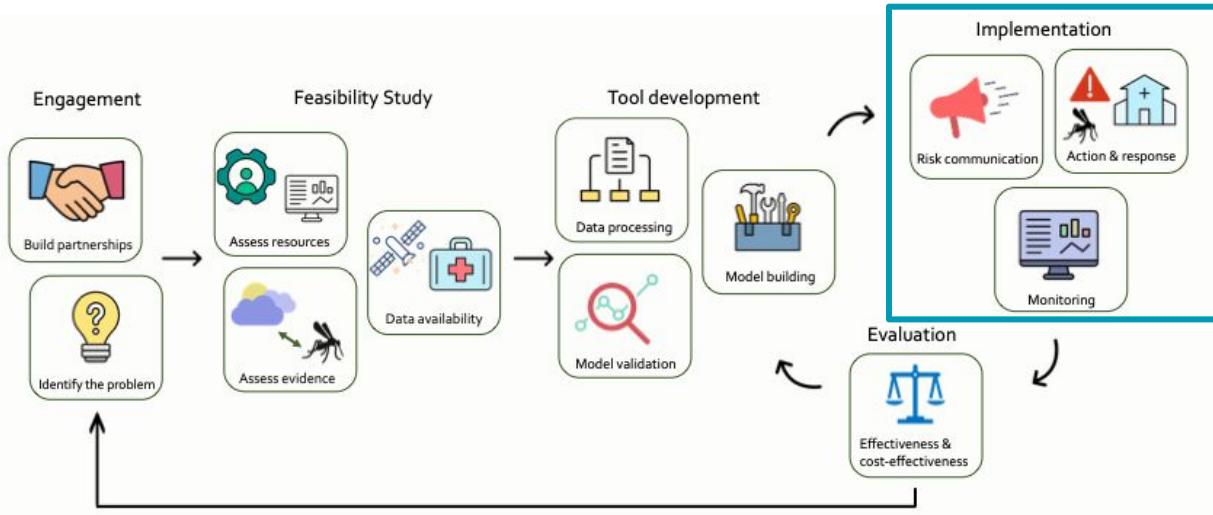
Associated
levels of
intervention

- Predicting cases vs. outbreaks.
- Translate complex results into clear insights.
- Conveying uncertainty.

Fletcher *et al.* *Lancet Planetary Health* 2025

Early warning systems: implementation

Stages involved in the co-creation of an early warning system for vector-borne disease



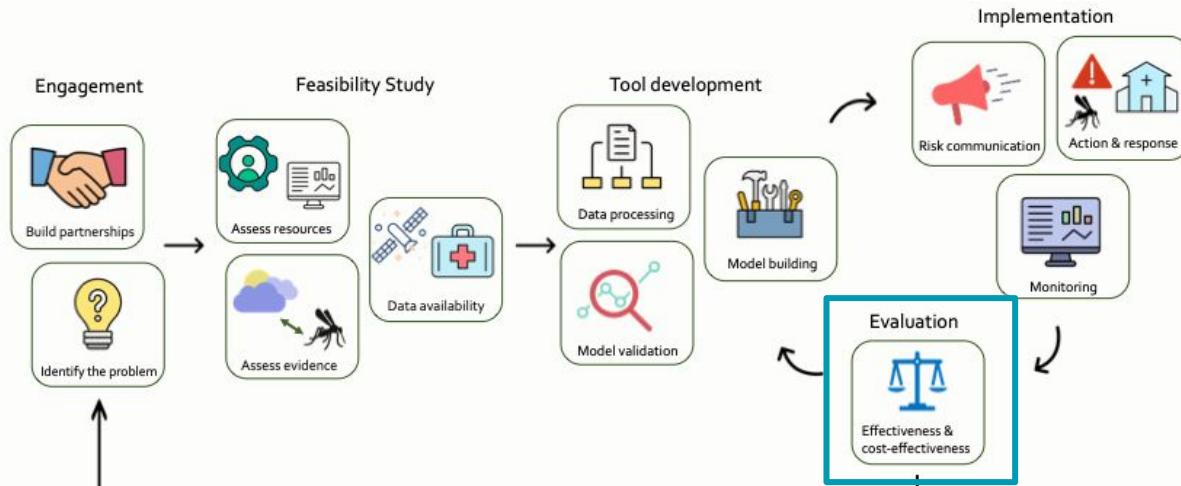
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Data structure, pre-processing and exploration

Types of data

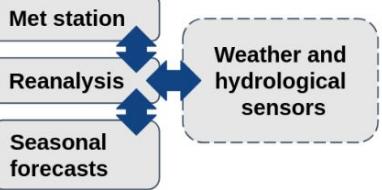


HARMONIZE



Barcelona
Supercomputing
Center
Centro Nacional de Supercomputación

Weather & Climate Data



Type depends on EWS

Temperature, precipitation, oceanic indices

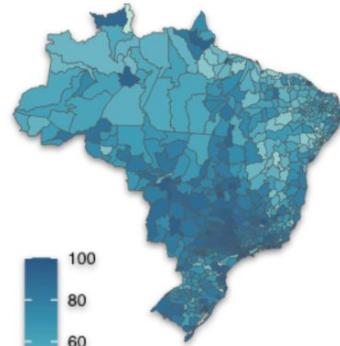
Land use and land cover data



Vegetation indices

Water indices

Demographic and socio-economic indicators

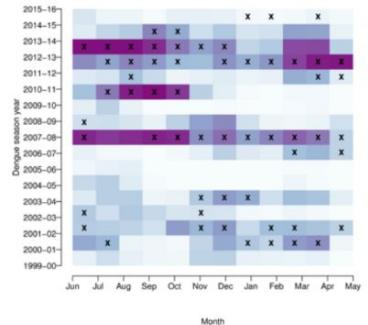


% residents living in urban areas

Vulnerability and inequality

Population at risk

Disease surveillance data



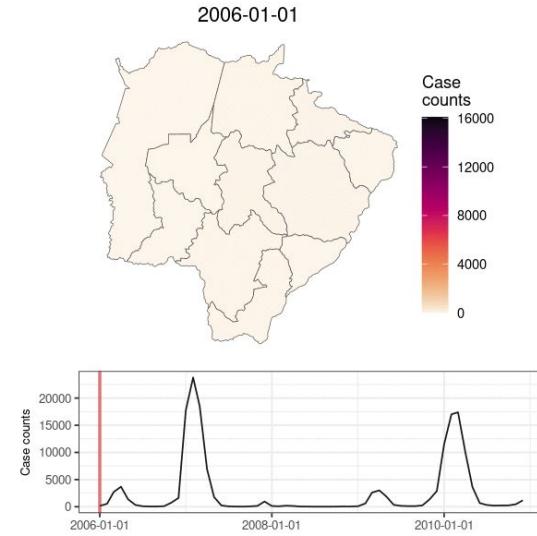
Variable of interest

Location-dependent

Data structure

Regular time series of one or several areas containing:

- Temporal ID (date)
- Spatial ID (area)
- Diseases counts for a specific time/area
- Population at risk for a specific time/area
- Predictors:
 - Spatio-temporal: climate
 - Temporal: ENSO indicator
 - Spatial: climate area (Köppen)



	date	geocode	cases	tas	prlr	nino
1	2010-01-01	24001	3	26.40411	66.1383667	1.52
2	2010-02-01	24001	2	27.24414	52.0459404	1.27
3	2010-03-01	24001	2	27.49093	49.4391060	0.95
4	2010-04-01	24001	5	26.88115	116.6783752	0.49
5	2010-05-01	24001	23	26.77081	44.9898071	-0.10
6	2010-06-01	24001	29	25.48106	70.8722610	-0.55
7	2010-07-01	24001	15	24.37067	91.5613556	-0.96
8	2010-08-01	24001	12	24.27328	28.0503407	-1.36

Cumulative climate data can capture factors that a single-point variable doesn't reflect:

- Temperature (average):
 - Duration of warm/cold conditions -> vector cycle
- Precipitation (sum):
 - Persistent low precipitation -> drought
 - Persistent high precipitation -> high soil moisture and humidity

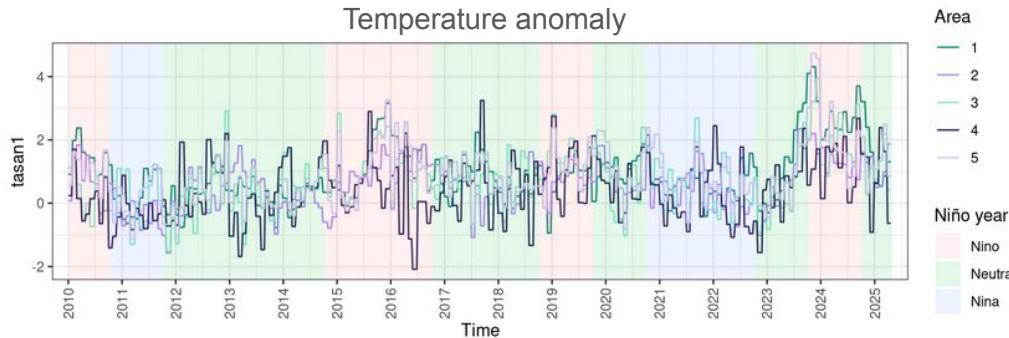
Date	Temperature	Temperature - 3 months
2015-01-01	12	
2015-02-01	13	
2015-03-01	15	13.3
2015-04-01	17	15
2015-05-01	20	17.3
2015-06-01	23	20

Averaging times depend on the health outcome.

Vector-borne: 3-12 months.

Pre-processing: temperature and precipitation anomalies

We can create **anomaly indices** to compare the precipitation for a given location and time to their respective climatologies (30+ years of historical data)



1. We compute historical monthly means and standard deviations for each location in our dataset.
2. We standardise the values (z-scores) in our dataset using these values.

Precipitation values require an extra standardization step due to their skewed distribution

<https://doi.org/10.1175/2009JCLI2909.1>

Pre-processing: lagging climate data

We want to include **lags of the exposures** because:

- Delayed effects.
- Be able to use observations in forecasts.
 - If we use a 3-month lag, we will be able to predict in 3 months using the observed weather today.

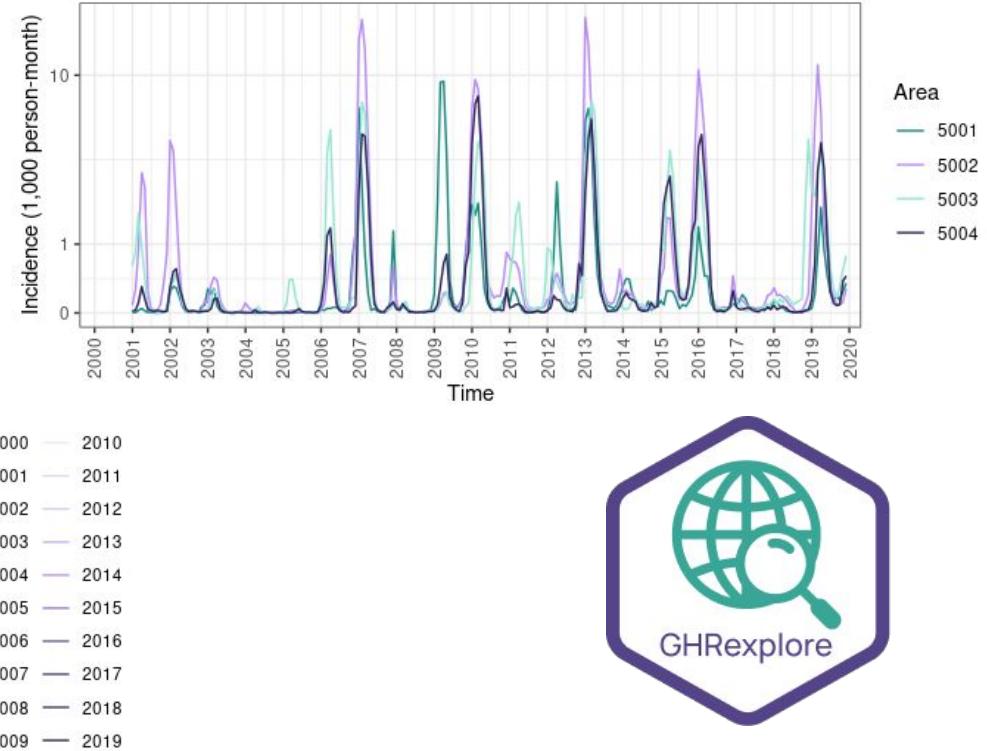
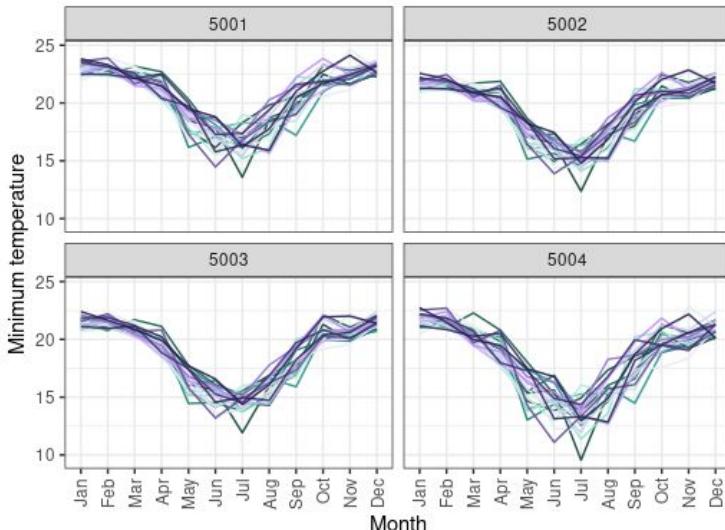
Date	Temperature	Temperature - lag 3
2015-01-01	12	
2015-02-01	13	
2015-03-01	15	
2015-04-01	17	12
2015-05-01	20	13
2015-06-01	23	15

The lag value will depend on the health outcome.

Vector-borne: 1-6 months.

Exploring: temporal component

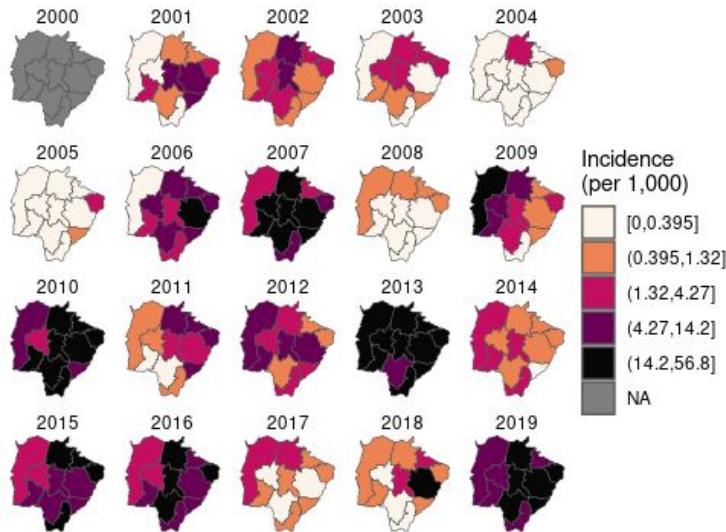
Time series



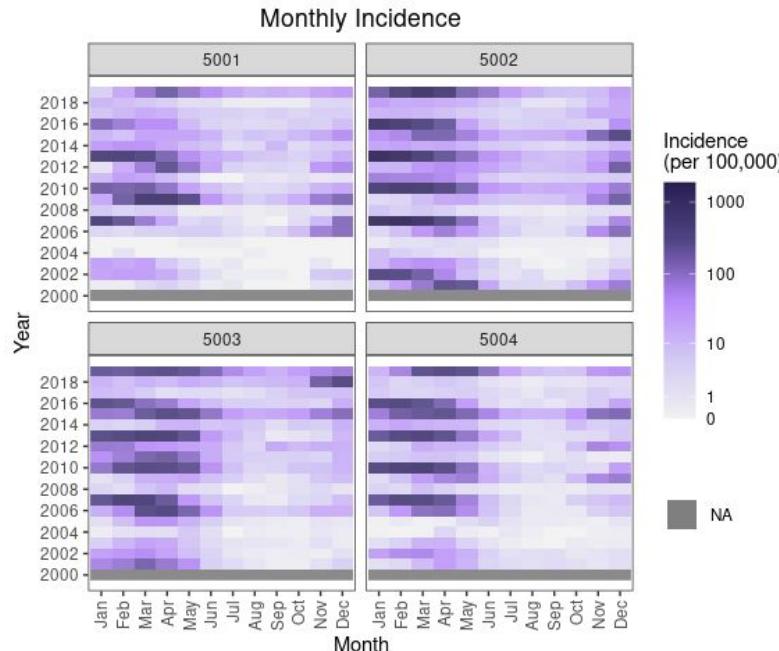
Seasonality plots

Exploring: spatial and spatio-temporal component

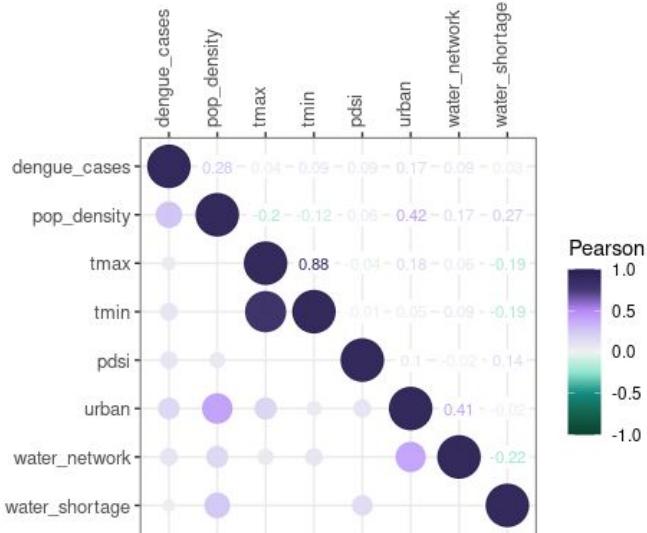
Choropleth maps



Heatmaps

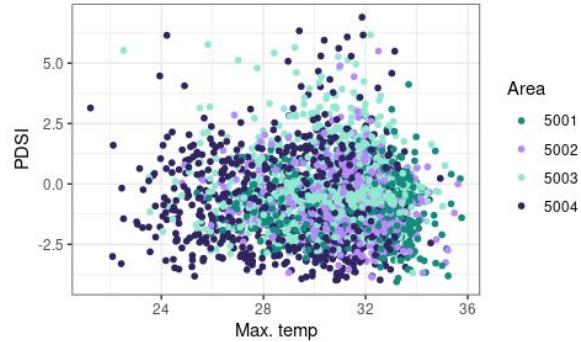


Exploring: correlation

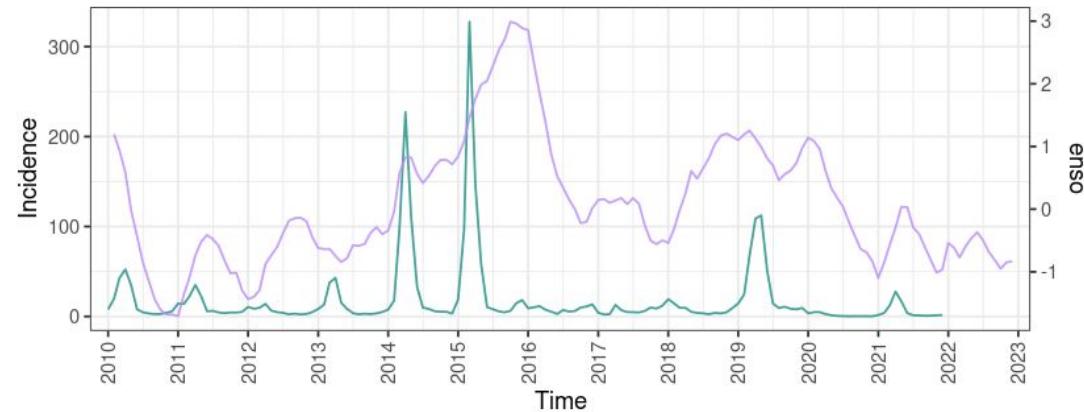


Correlation matrices

Scatterplots



Dual-axis time series



Variable Incidence enso



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The GHR team!!



PROF RACHEL LOWE
LEADING RESEARCHER



DR GIOVENALE MOIRANO
VISITING
RESEARCHER



TILLY ALCAYNA
VISITING
PHD CANDIDATE



DR MARTIN LOTTO
RESEARCHER



CHLOE FLETCHER
PHD CANDIDATE



DR BRUNO CARVALHO
VISITING
RESEARCHER

Time for questions

Ania Kawiecki Peralta (ania.kawiecki@bsc.es)

Carles Milà Garcia (carles.milagarcia@bsc.es)

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