



Prendre en compte notre environnement
fluctuant pour la modélisation des maladies
vectorielles.

Bernard Cazelles

MAMBA, INRIA, Paris

IBENS, CNRS-ENS, Paris

UMMISCO, IRD-Sorbonne Université, Paris



Cette intervention est faite en toute indépendance vis-à-vis de l'organisateur de la manifestation.

Je n'ai pas de conflit d'intérêts en lien avec le sujet traité.

Accounting for Non-Stationarity in Epidemiology

Overview

- **Non-stationarity and transients in Epidemiology**
- Accounting for Non-Stationarity in Statistical Analysis
- Accounting for Non-Stationarity in Modeling

Non-stationarity and transients in Epidemiology

- Modification of pathogens, their transmissibility, their virulence
- Characteristics of the epidemics can evolve due to vaccination or others public health interventions
- Climate can influence the propagation of a pathogen
- Societal responses and/or changing human behavior during the course of an epidemic
 - Changing their social network
 - Social distancing
 - Voluntary avoidance behavior

Non-stationarity and transients in Epidemiology

■ Example of measles and whooping cough in UK

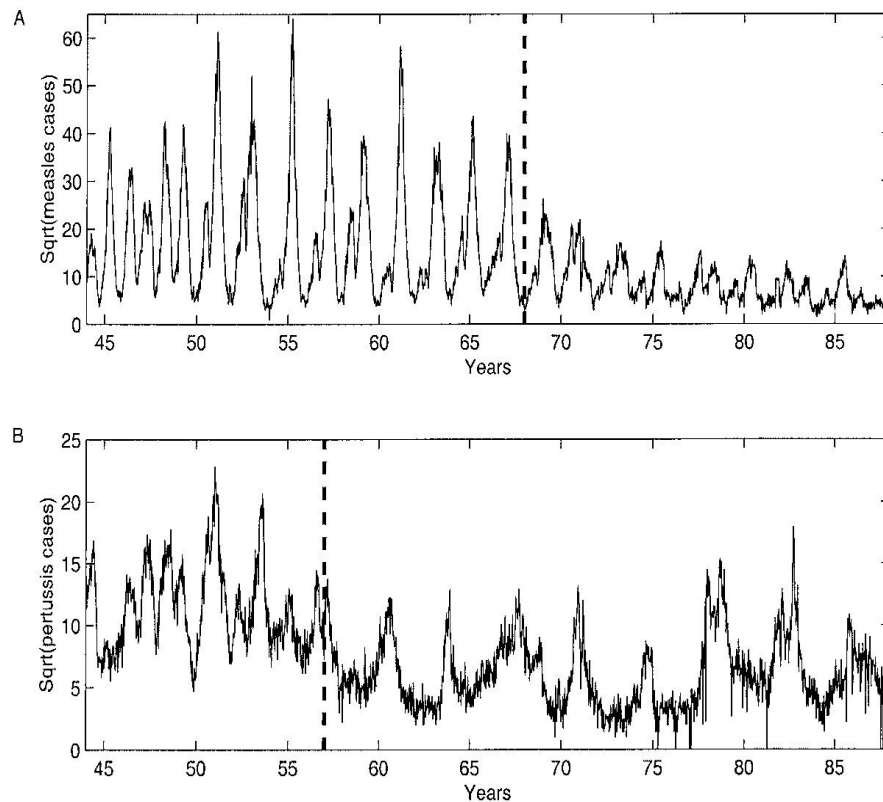
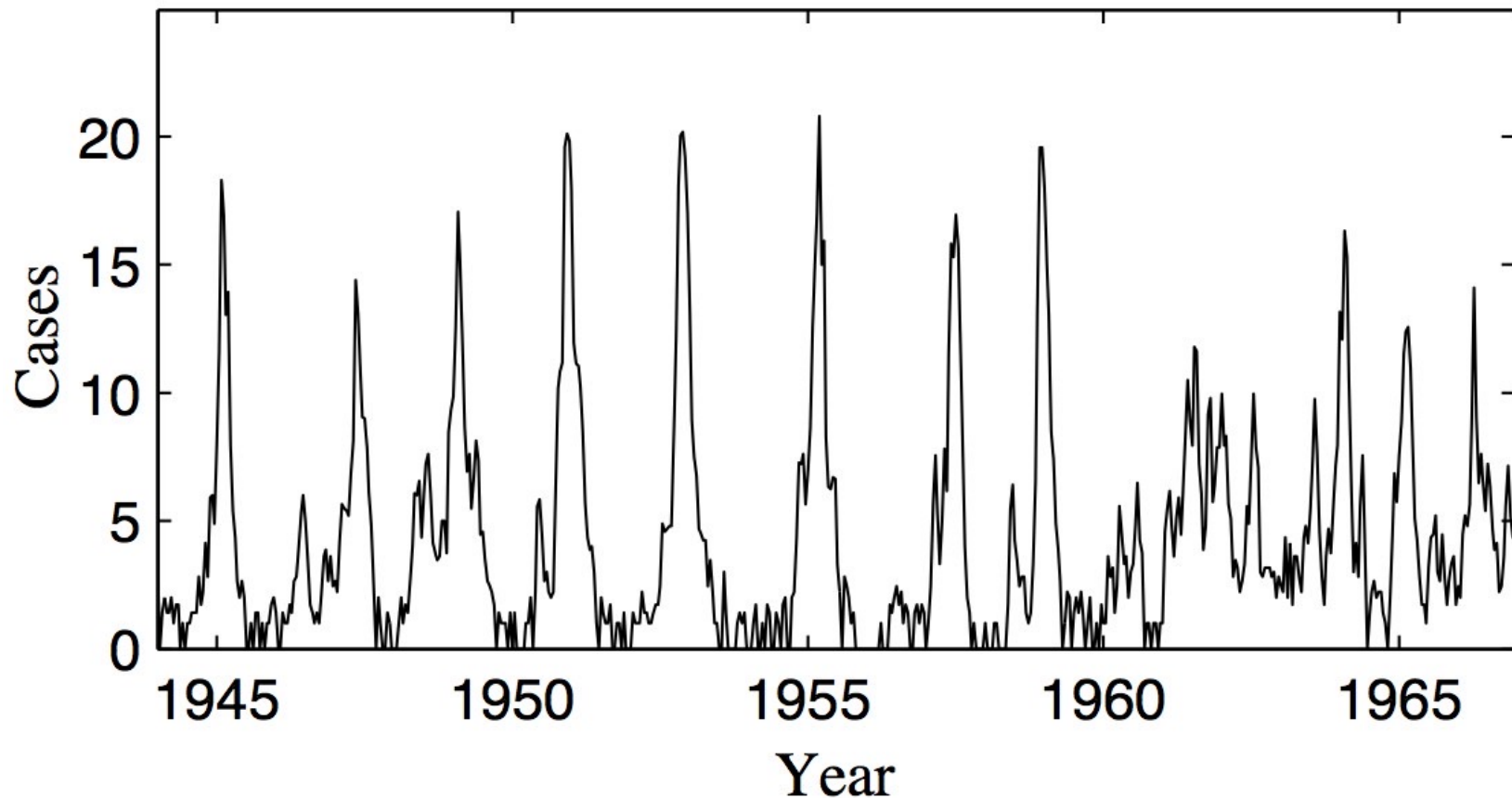


Figure 1: Aggregate measles and whooping cough notifications in England and Wales from 1944 to 1994; data obtained from the Registrar General's Weekly Returns. A, Time series for square root of measles cases in England and Wales, with vaccination starting in 1968 (*dotted line*). B, Square root of cases of whooping cough in England and Wales, with the onset of national vaccination indicated by the dotted line.

Non-stationarity and transients in Epidemiology

- An example of measles epidemics in York (UK)



Characteristics evolve with time => Non-Stationarity

Accounting for Non-Stationarity in Epidemiology

Overview

- Non-stationarity and transients in Epidemiology
- **Accounting for Non-Stationarity in Statistical Analysis**
- Accounting for Non-Stationarity in Modeling

Accounting for Non-Stationarity in Statistical Analysis

- For statistical approaches I have developed numerous tools using wavelet decomposition
- Wavelet analysis decomposes a signal into **time-space** and **frequency-space simultaneously**
- Wavelet analysis estimates the spectral characteristics of a time series as a function of time

Wavelet Analysis

Different wavelet functions

Mexican hat

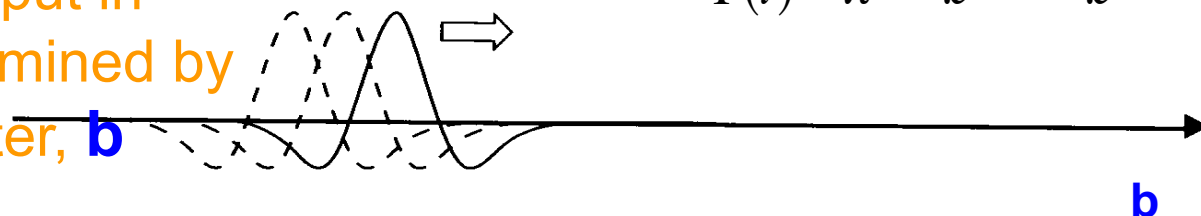
Morlet

$$\Psi(t) = (1 - t^2).e^{-t^2/2}$$

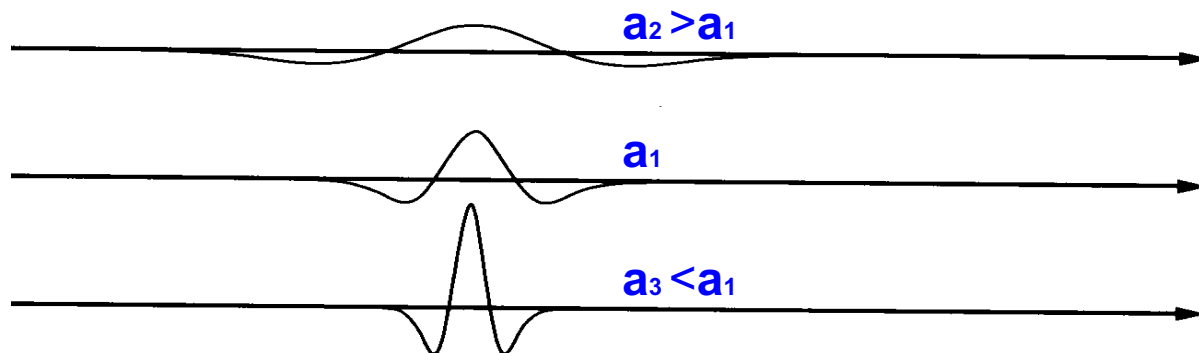


$$\Psi(t) = \pi^{-1/4} . e^{i.2\pi.f_0.t} . e^{-t^2/2}$$

The wavelet function is put in different positions determined by the translation parameter, **b**



and with different dilatation parameter, **a**

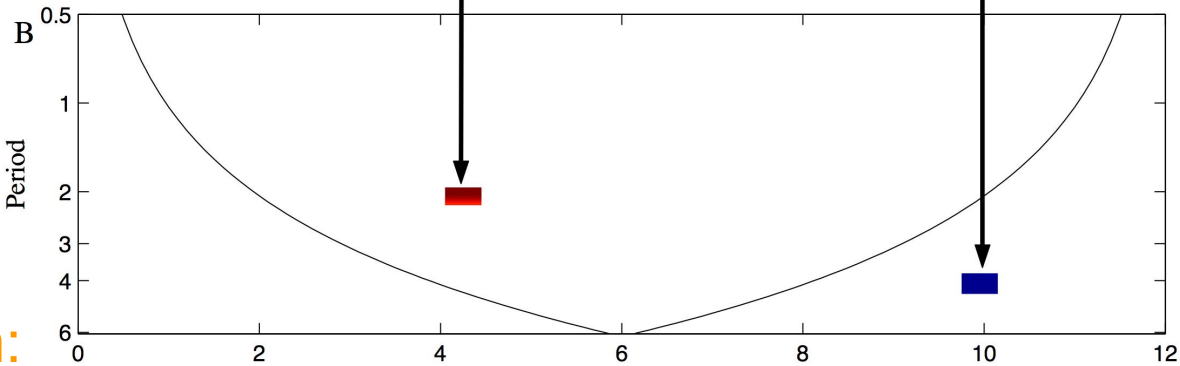
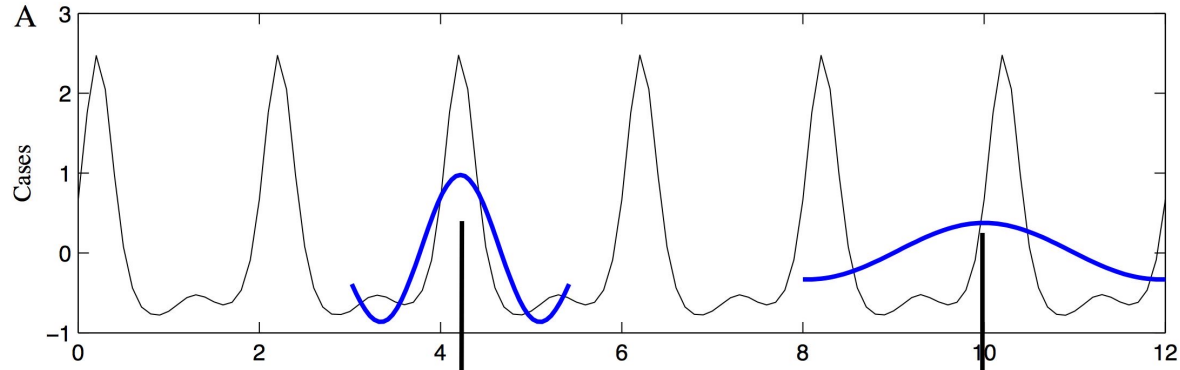


$$\Psi(a,b) = \Psi\left(\frac{t-b}{a}\right)$$

Wavelet Analysis

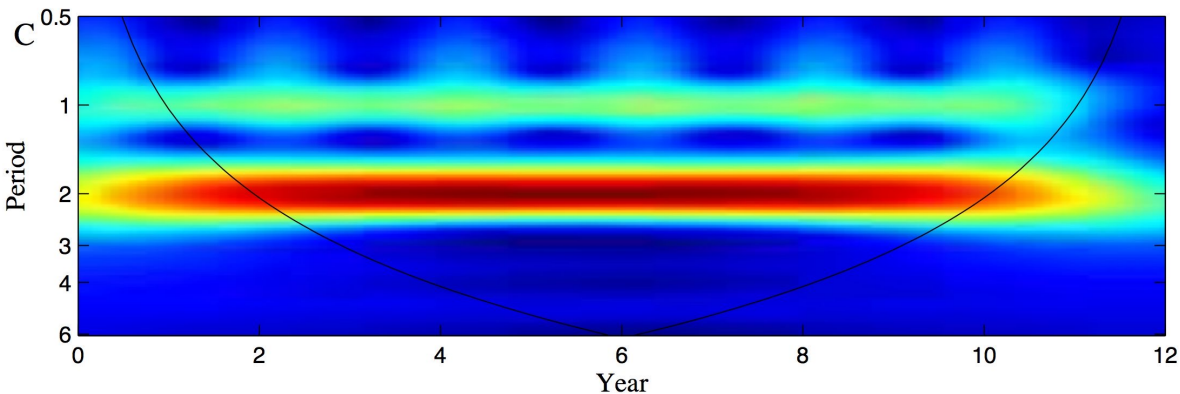
Wavelet Transform:

$$T(a,b) = \frac{1}{\sqrt{a}} \int_{-\infty}^{\infty} x(t) \cdot \Psi^* \left(\frac{t-b}{a} \right) dt$$



Wavelet Power Spectrum:

$$W_x(a,b) = |T(a,b)|^2$$



Wavelet Analysis

Wavelet Transform:
(and Inverse Wavelet Transform)

$$T(a,b) = \frac{1}{\sqrt{a}} \int_{-\infty}^{\infty} x(t) \cdot \Psi^* \left(\frac{t-b}{a} \right) dt$$

Wavelet Amplitude, Imaginary Part and
Wavelet Modulus:

$$\frac{\operatorname{Re}(T(a,b)) \quad \operatorname{Im}(T(a,b))}{|T(a,b)|}$$

Wavelet Power:

$$W_x(a,b) = |T(a,b)|^2$$

Averaged Wavelet Power:
(by analogy with the Fourier Power)

$$P_W(f) = \frac{1}{\tau f_c C_g} \int_0^{\tau} |T(a,b)|^2 db$$

CrossWavelet Power:

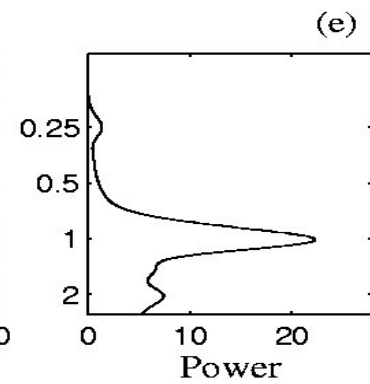
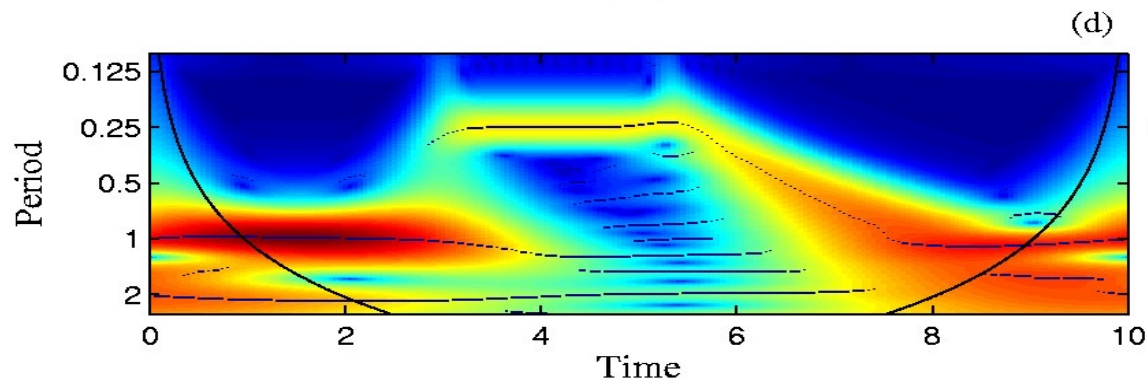
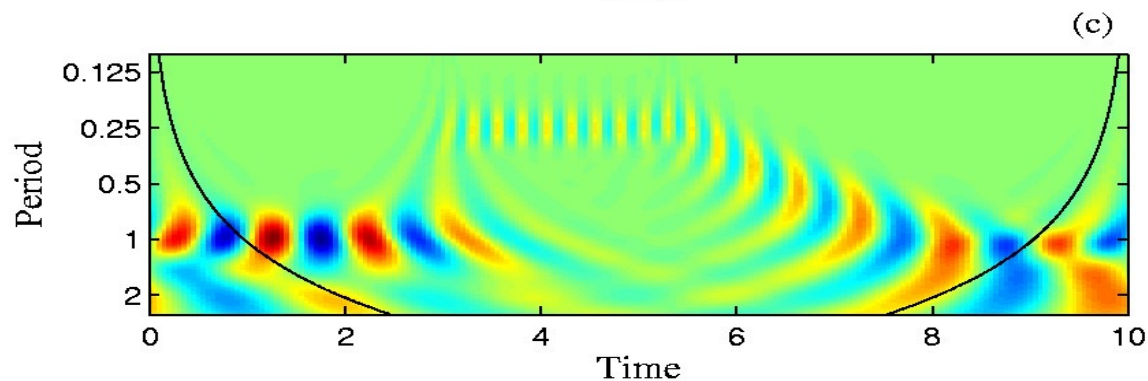
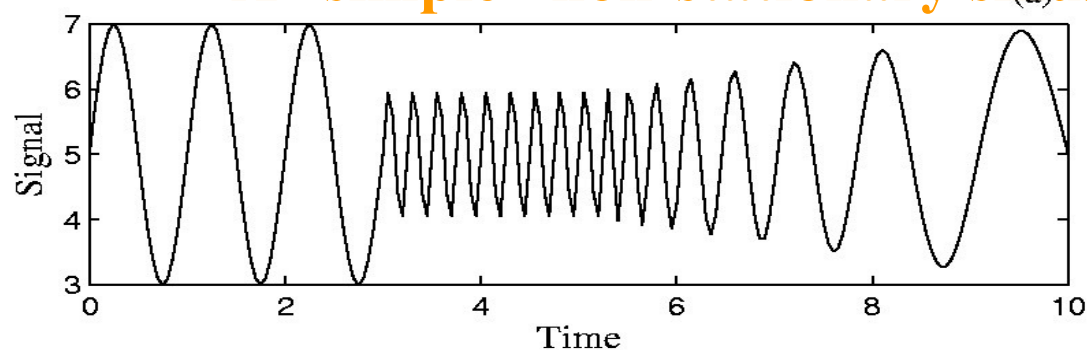
$$W_{xy}(a,b) = \frac{W_x(a,b) \cdot W_y(a,b)^*}{\sigma_x \cdot \sigma_y}$$

Square Wavelet Coherency:

$$CW_{xy} = \frac{|\langle W_{xy}(a,b) \rangle|^2}{|\langle W_x(a,b) \rangle| \cdot |\langle W_y(a,b) \rangle|}$$

Wavelet Analysis

A “simple” non-stationary sinusoidal signal



Dengue in Thailand

Travelling waves in the occurrence of dengue haemorrhagic fever in Thailand

Derek A.T. Cummings^{1,2}, Rafael A. Irizarry³, Norden E. Huang⁴, Timothy P. Endy⁵, Ananda Nisalak⁶, Kumnuan Ungchusak⁷ & Donald S. Burke²

¹Department of Geography and Environmental Engineering, Johns Hopkins University, Baltimore, Maryland 21218, USA

²Department of International Health, and ³Department of Biostatistics, Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland 21205, USA

⁴Laboratory for Hydrospheric Processes/Oceans and Ice Branch, NASA Goddard Space Flight Center, Greenbelt, Maryland 20771, USA

⁵Virology Division, United States Army Medical Research Institute in Infectious Disease, Fort Detrick, Maryland 21702, USA

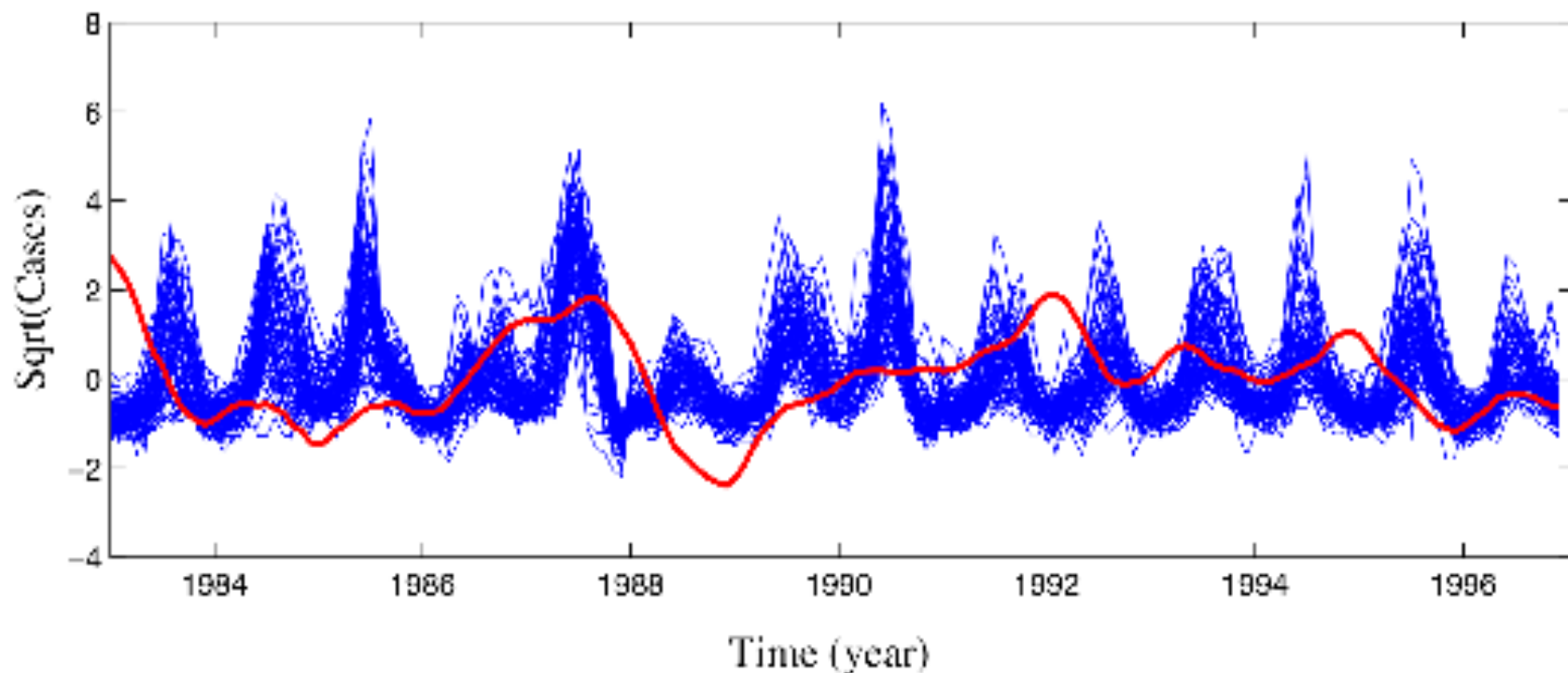
⁶Department of Virology, Armed Forces Research Institute of Medical Sciences, Bangkok 10400, Thailand

⁷Bureau of Epidemiology, Ministry of Public Health, Nonthaburi 11000, Thailand

Dengue fever is a mosquito-borne virus that infects 50–100 million people each year¹. Of these infections, 200,000–500,000 occur as the severe, life-threatening form of the disease, dengue haemorrhagic fever (DHF)². Large, unanticipated epidemics of DHF often overwhelm health systems³. An understanding of the spatial–temporal pattern of DHF incidence would aid the

But any links with climate!

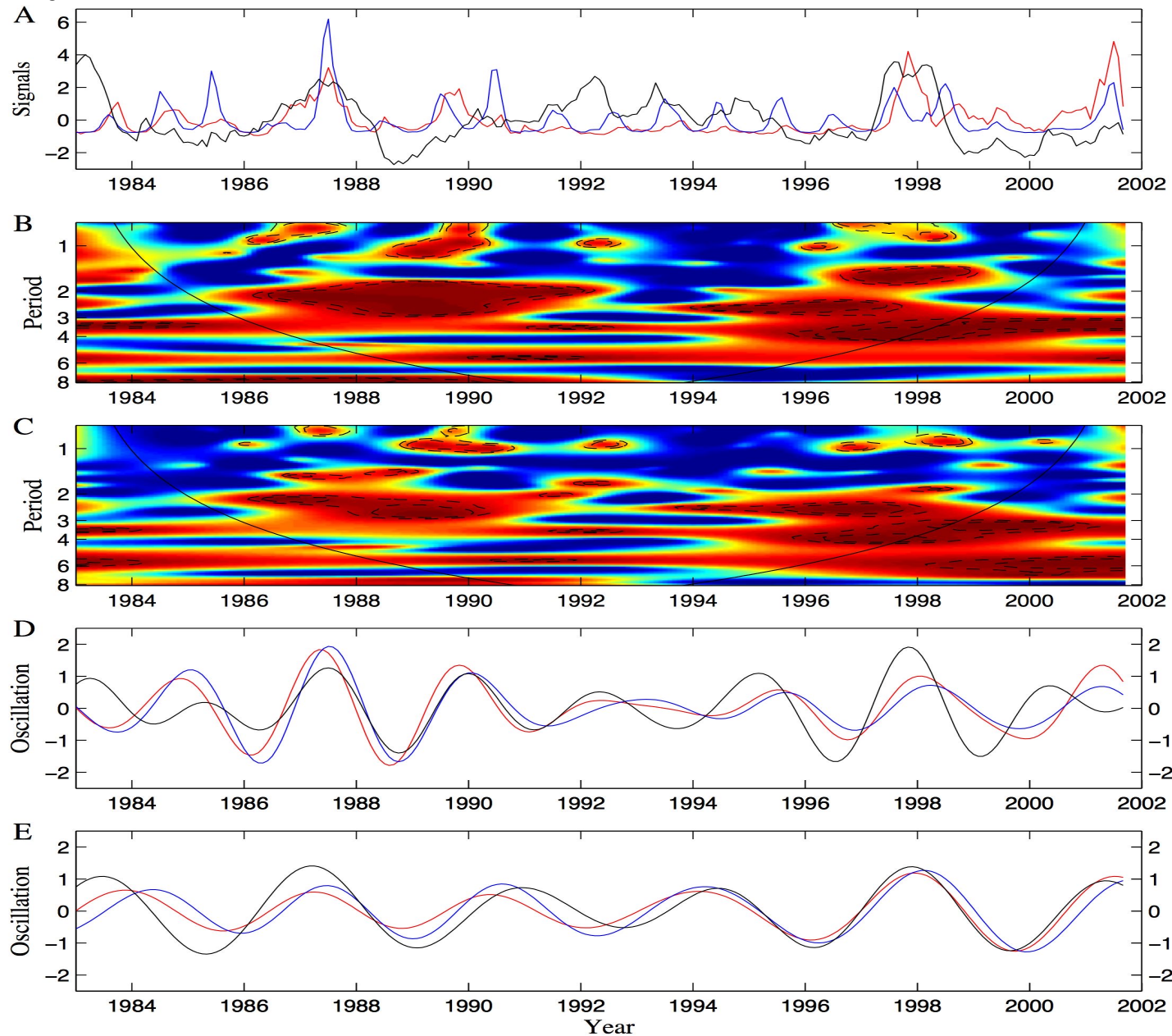
Dengue in Thailand



- Monthly DHF cases reported in the 76 provinces of Thailand
- Focus on the incidence in Bangkok and the averaged incidence for the rest of Thailand

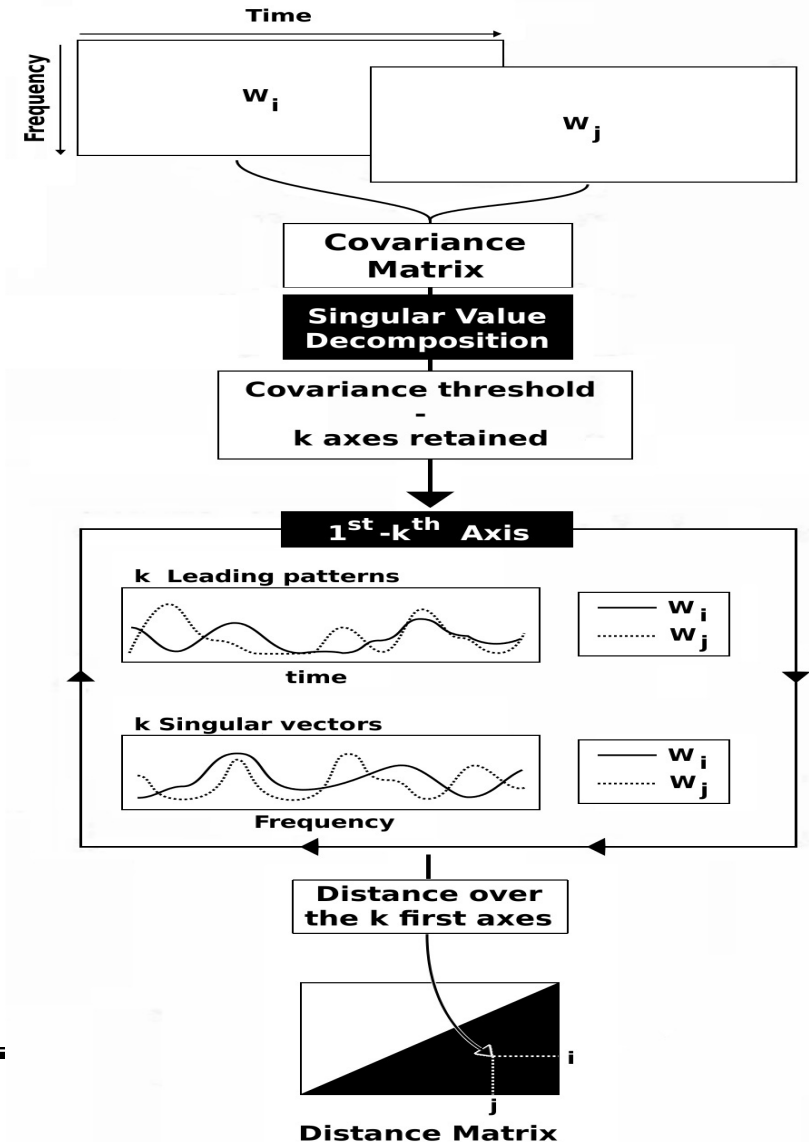
Wavelet Coherency

Dengue in Thailand and ENSO



The Rouyer's Method

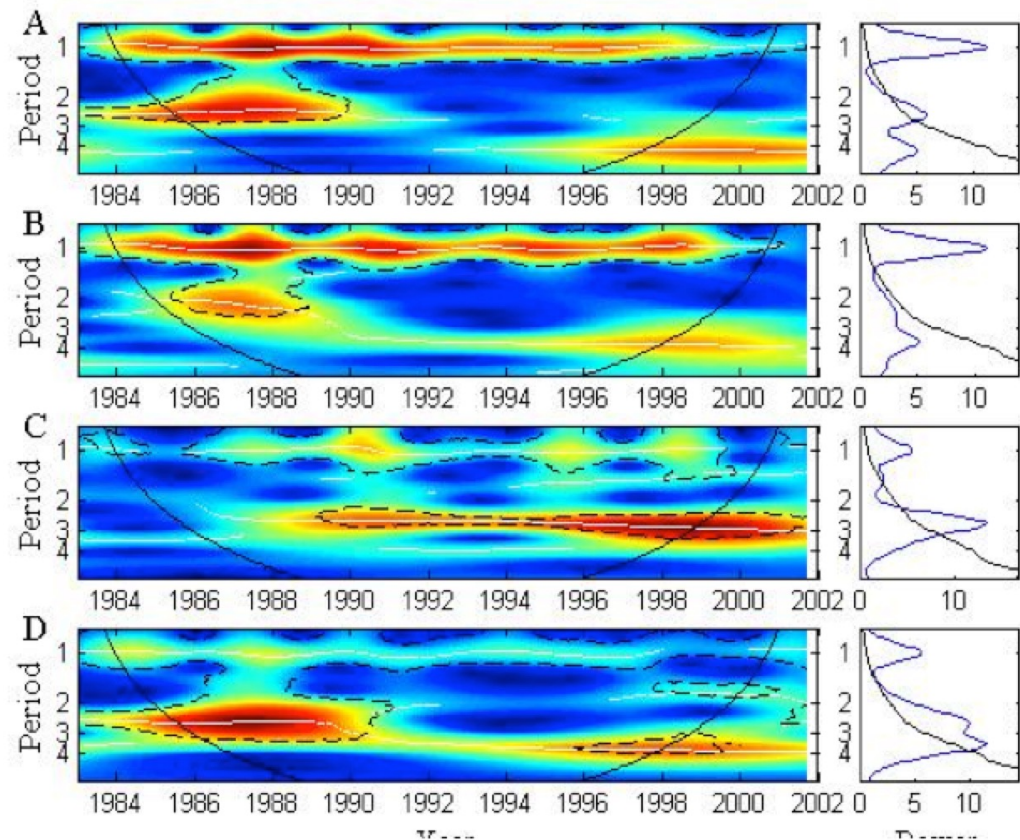
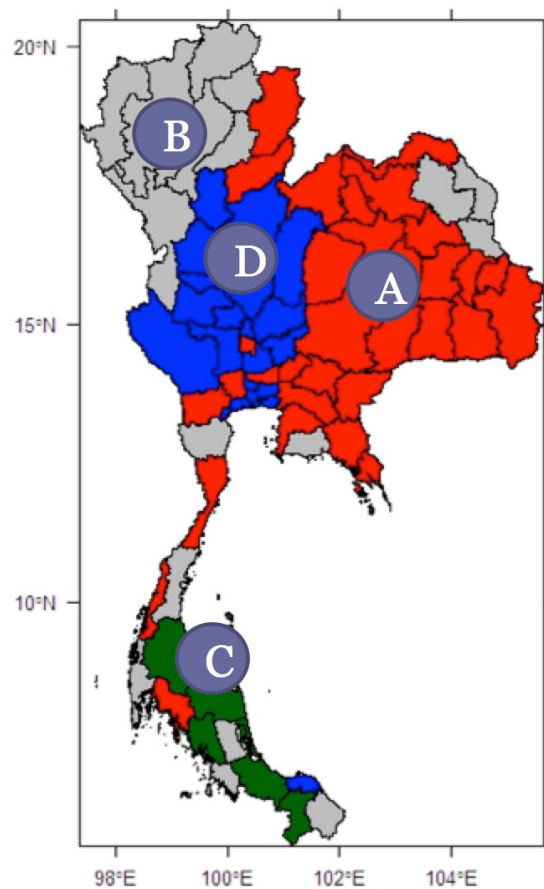
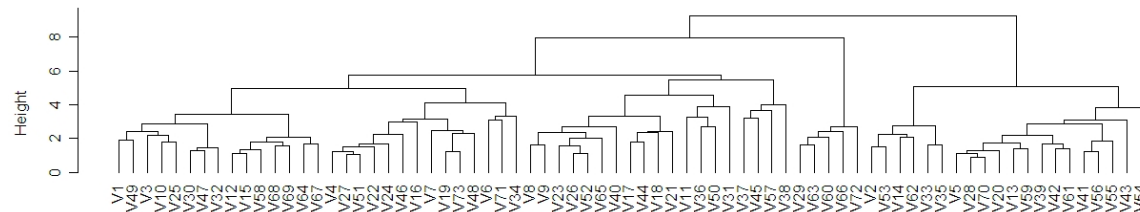
Rouyer, T., Fromentin, J.M., Stenseth, N.C. & Cazelles, B., 2008.
Analysing multiple time series and extending significance testing
in wavelet analysis. *Marine Ecology Progress Series*, 359, 11-23.



Classical clustering based
on this Distance Matrix

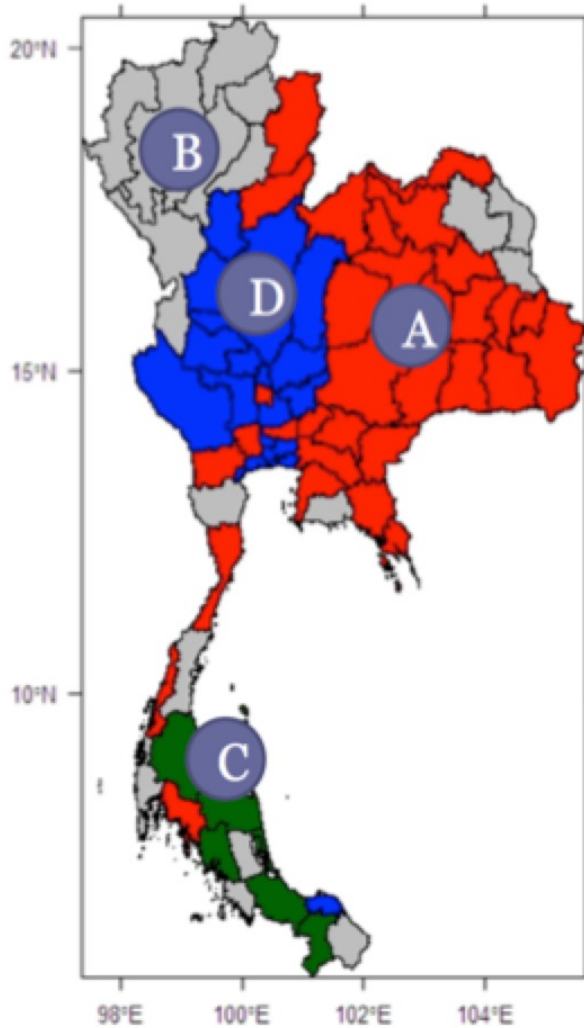
Wavelet Clustering

Cluster Dendrogram

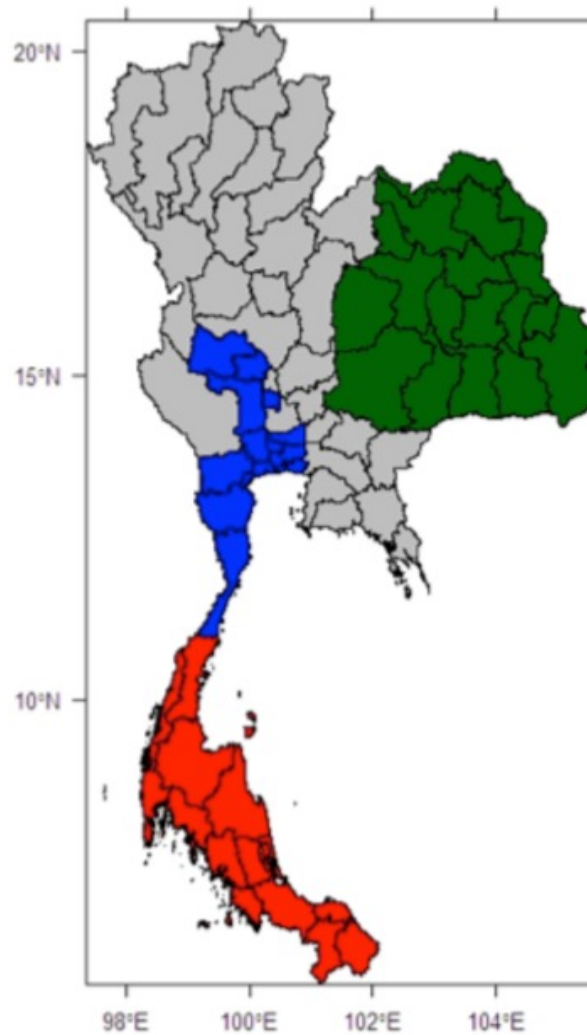


Wavelet Clustering

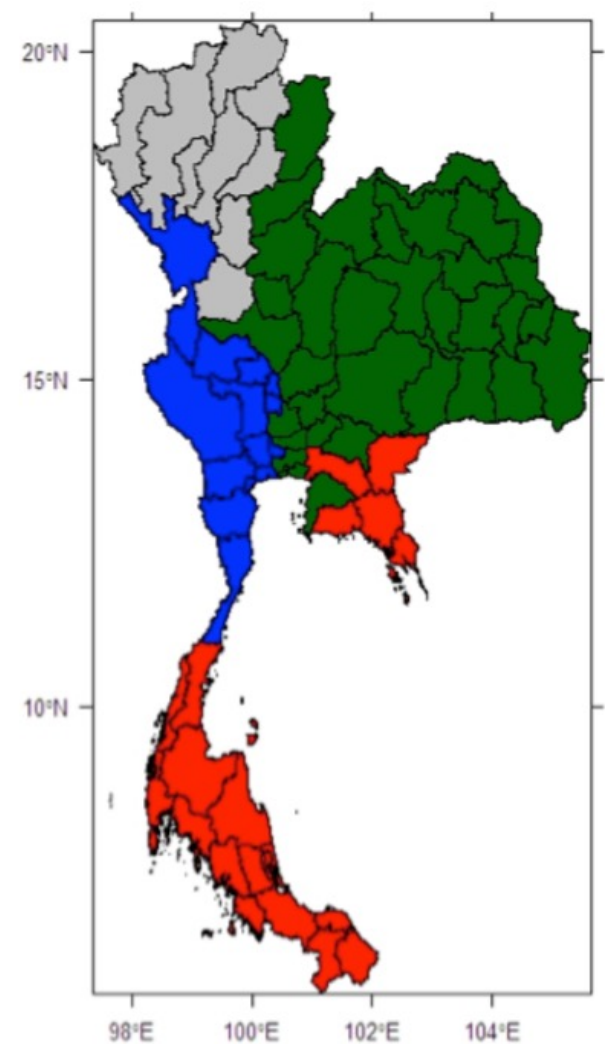
Dengue



Rainfall



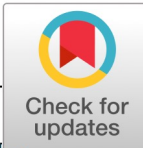
Average temperature



Disentangling local and global climate drivers in the population dynamics of mosquito-borne infections

Bernard Cazelles^{1,2}, Kévin Cazelles^{3,4}, Huaiyu Tian⁵, Mario Chavez^{6*}, Mercedes Pascual^{7,8*}

Identifying climate drivers is essential to understand and predict epidemics of mosquito-borne infections whose population dynamics typically exhibit seasonality and multiannual cycles. Which climate covariates to consider varies across studies, from local factors such as temperature to remote drivers such as the El Niño–Southern Oscillation. With partial wavelet coherence, we present a systematic investigation of nonstationary associations between mosquito-borne disease incidence and a given climate factor while controlling for another. Analysis of almost 200 time series of dengue and malaria around the globe at different geographical scales shows a systematic effect of global climate drivers on interannual variability and of local ones on seasonality. This clear separation of time scales of action enhances detection of climate drivers and indicates those best suited for building early-warning systems.



Copyright © 2023 The Authors, some rights reserved; exclusive licensee American Association for the Advancement of Science. No claim to original U.S. Government Works. Distributed under a Creative Commons Attribution NonCommercial License 4.0 (CC BY-NC).

Disentangling the local and global climatic influences on dengue

Application to Dengue in Thailand

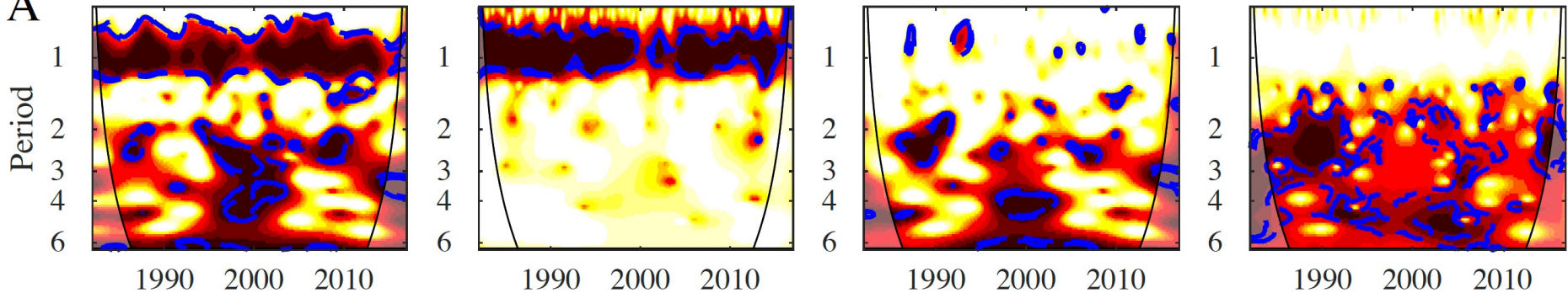
WC D-LC

PWC D-LC|GC

WC D-GC

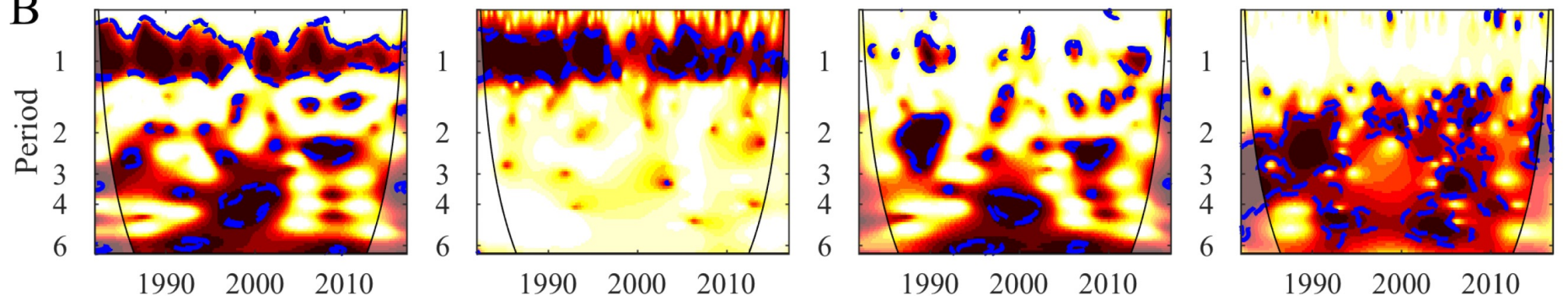
PWC D-GC|LC

A



Whole Thailand: dengue, mean temperature and ONI

B



Zone 4: dengue, rainfall and MEI

ionarity in Statistical Analysis

Kutch district (India): Malaria, Rainfall, DMI.

Application to Malaria

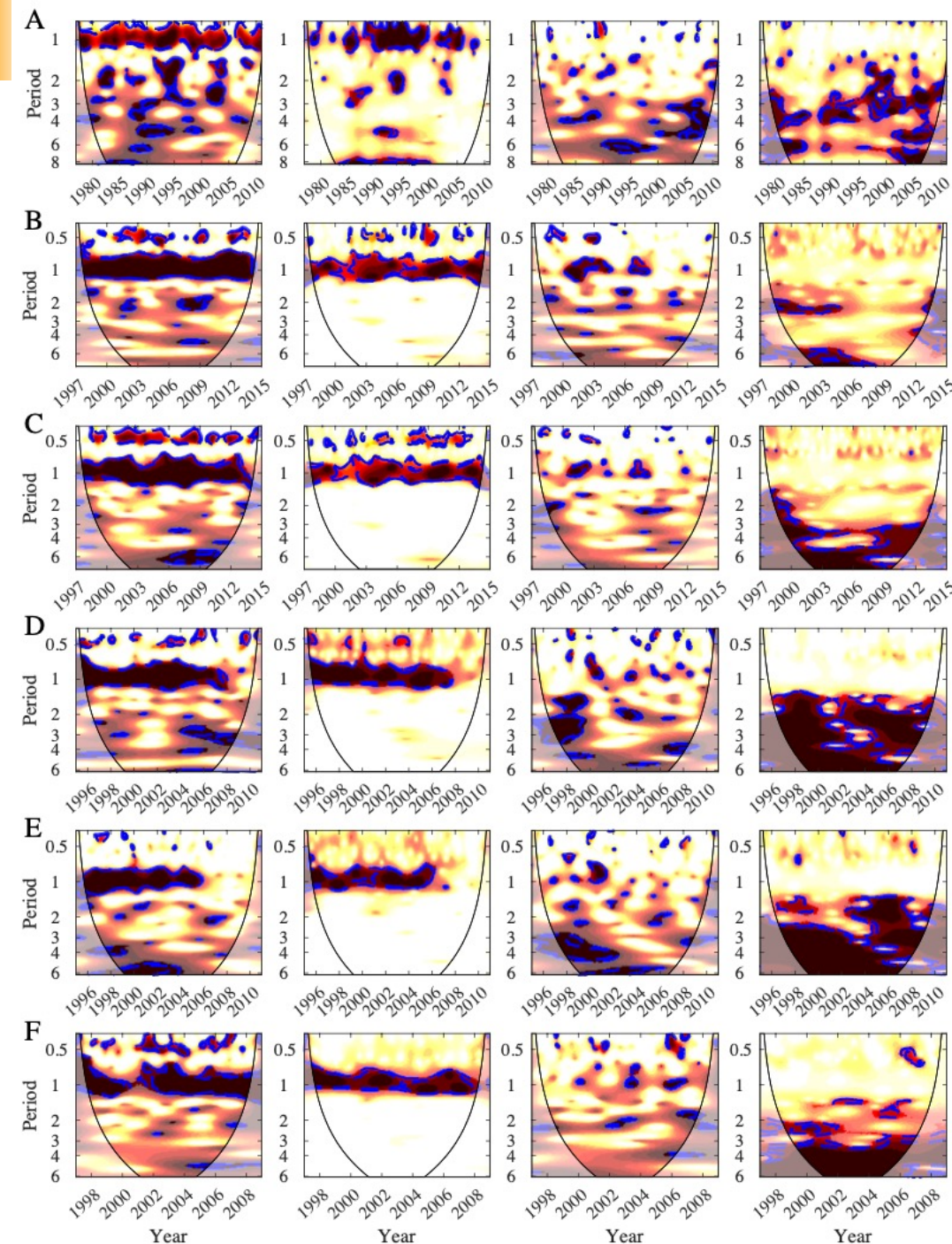
Surat (India): Malaria, Relative Humidity, SOI

Ahmedabad (India): Malaria, Temperature, MEI.

Hainan province: Malaria, Rainfall, ONI.

Hainan province: Malaria, Temperature, MEI.

Anhui : Malaria, Temperature, Nino34.



Non-Stationnarité et Modélisation des Epidémies

Overview

- Non-stationarity and transients in Epidemiology
- Accounting for Non-Stationarity in Statistical Analysis
- **Accounting for Non-Stationarity in Modeling**

Accounting for Non-Stationarity in Modeling

- Reconstruction the time evolution of some key parameters without any specific hypothesis:
- We have used:
 - State space models

$$\begin{cases} \dot{x}_t = g(t, x(t), \theta) + u_t \\ y_t | x_t = f(h(x(t)), y_t, \theta) + v_t \end{cases}$$

- Parameters considered to be state variables that follow a diffusion process
- Inference tools as Kalman Filter or Bayesian approaches (MCMC, K-MCMC and P-MCMC)

Accounting for Non-Stationarity In Modeling

- State space models

$$\begin{cases} \dot{x}_t = g(t, x(t), \theta) + u_t \\ y_t | x_t = f(h(x(t)), y_t, \theta) + v_t \end{cases}$$

- System process: an epidemiological model
- Observational process: a probabilistic law with an observation rate, ρ
 - *Poisson*
 - *Negative Binomial*
 - *Normal*

Accounting for Non-Stationarity in Modeling

- State space models

$$\begin{cases} \dot{x}_t = g(t, x(t), \theta) + u_t \\ y_t | x_t = f(h(x(t)), y_t, \theta) + v_t \end{cases}$$

- Parameters considered to be state variables that follow a diffusion process

$$d\theta_t = \sigma dB_t$$

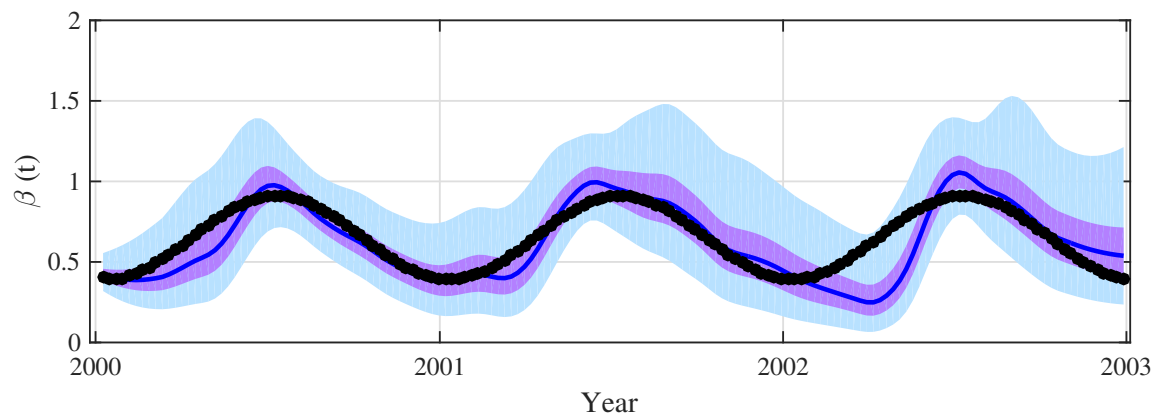
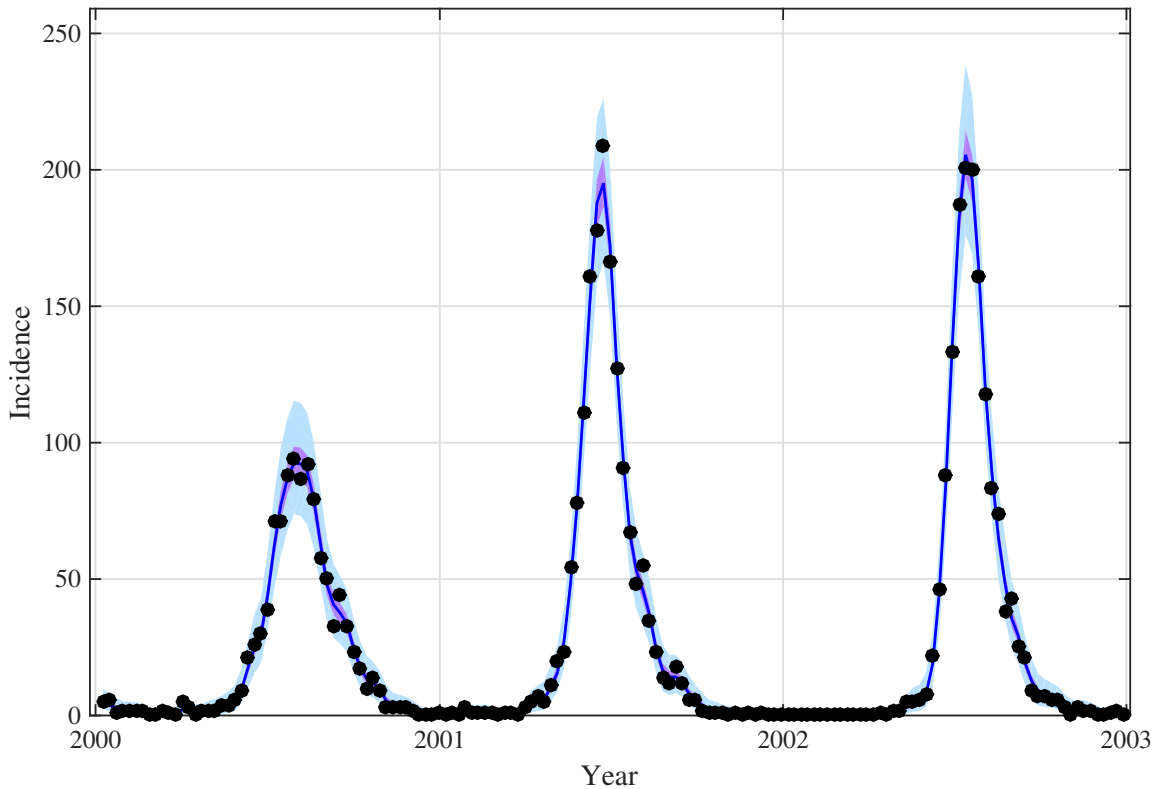
$$d \log(\theta_t) = \sigma dB_t$$

$$\theta_{t+1} = \theta_t + \sigma B_t$$

Accounting for Non-Stationarity in Modeling

- We used a stochastic framework with Markov jump process (or an approximation of it)
- In the stochastic framework, the likelihood is intractable thus EKF or SMC is used to compute it in the MCMC
- Thus we coupled time varying parameters approach with Bayesian methods coupling MCMC and EKF (K-MCMC) or SMC (P-MCMC) (Andrieu et al. 2010; Dureau et al. 2013)

A SIRS toy model



$$\frac{dS}{dt} = \mu \cdot (N - S) - \beta(t) \frac{S \cdot I}{N} + \alpha \cdot R$$

$$\frac{dI}{dt} = \beta(t) \frac{S \cdot I}{N} - (\gamma + \mu) \cdot I$$

$$\frac{dR}{dt} = \gamma \cdot I - (\alpha + \mu) \cdot R$$

$$\beta(t) = \beta_0 \cdot \left(1 + \beta_1 \cdot \sin\left(\frac{2\pi t}{365} + 2\pi\phi\right) \right)$$

Application to Dengue in Phnom Penh

SEIR model

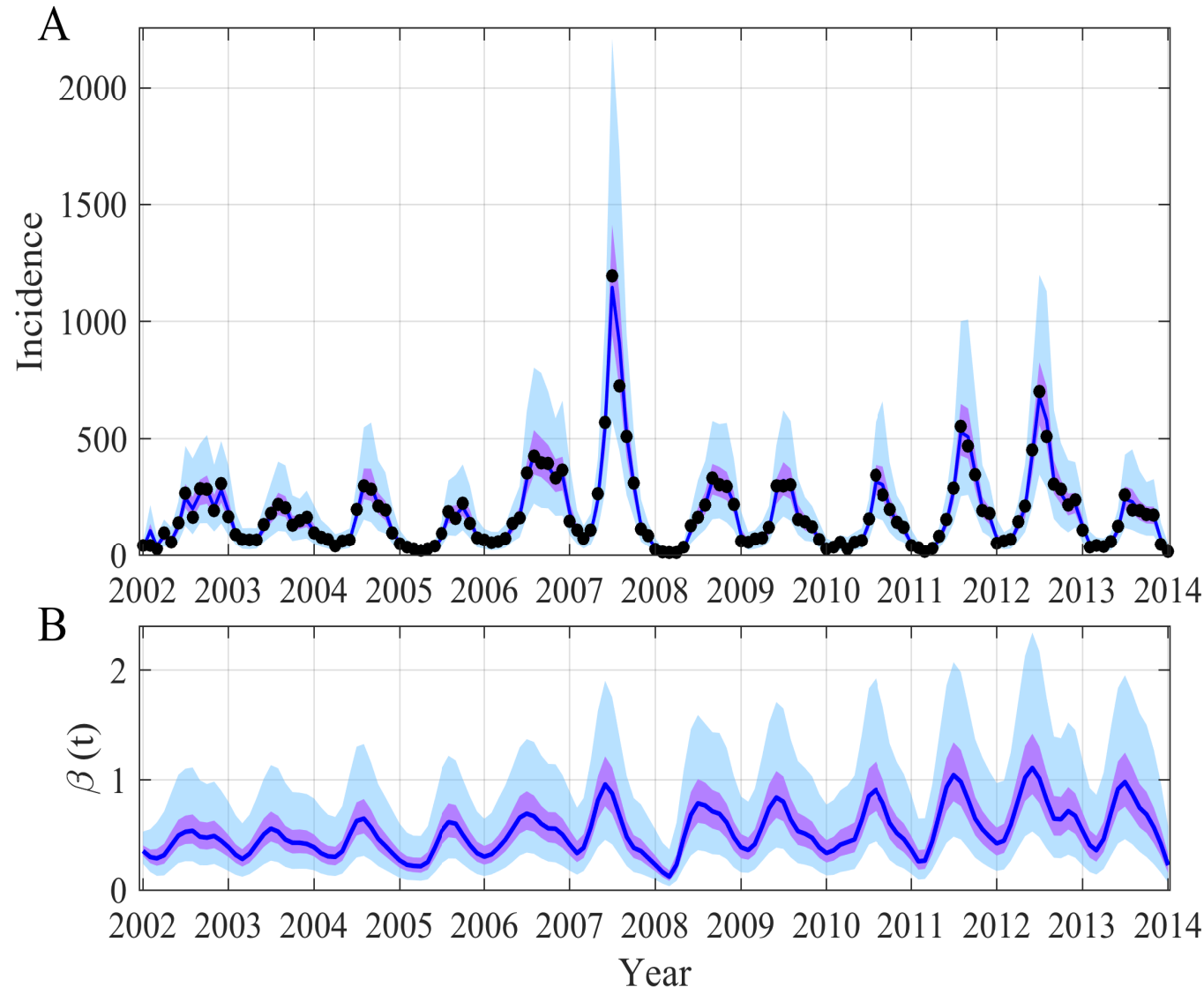
$$\frac{dS}{dt} = \mu \cdot (N - S) - \beta(t) \cdot \left(\frac{S \cdot I}{N} + i \right)$$

$$\frac{dE}{dt} = \beta(t) \cdot \left(\frac{S \cdot I}{N} + i \right) - (\delta + \mu) \cdot E$$

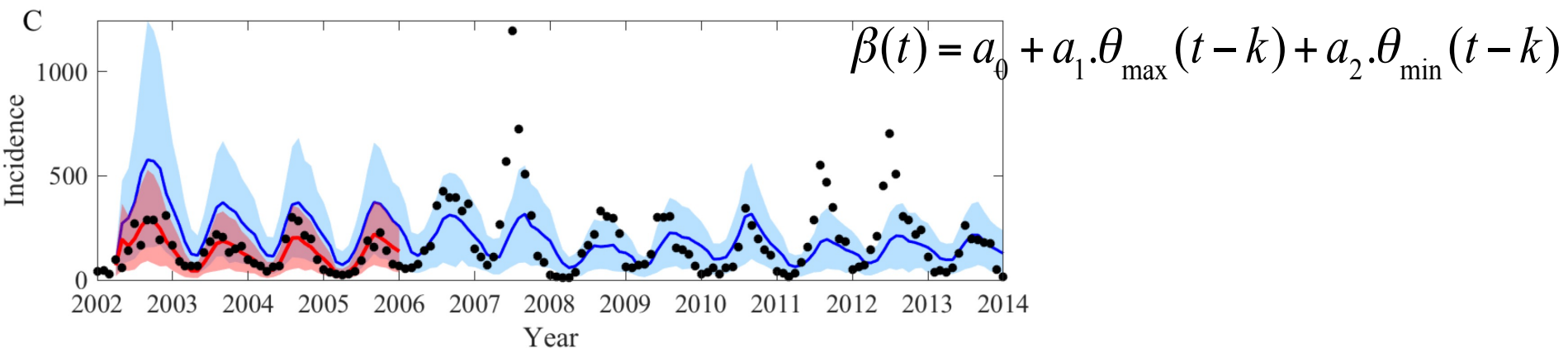
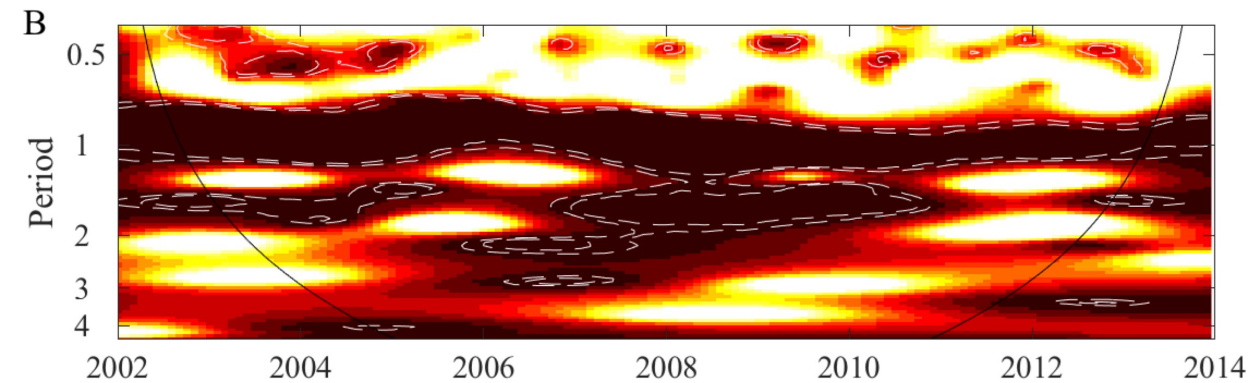
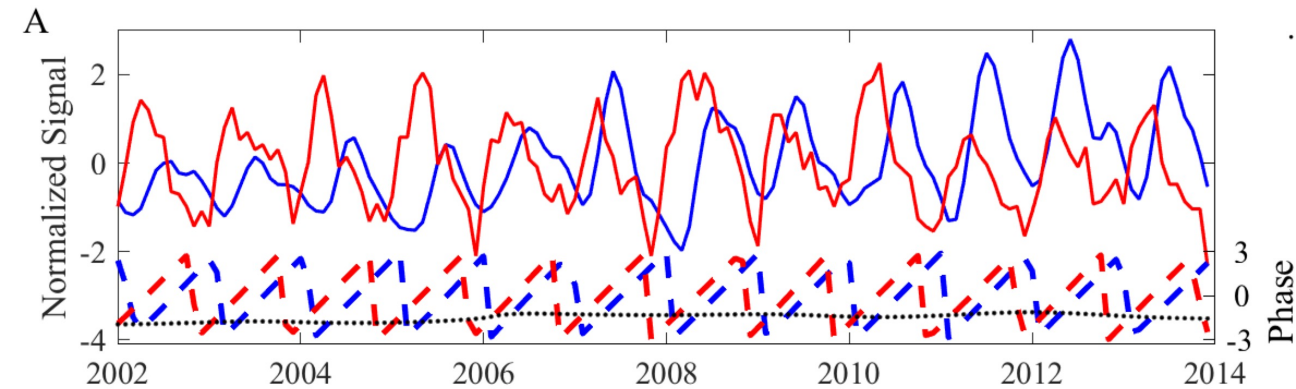
$$\frac{dI}{dt} = \delta \cdot E - (\gamma + \mu) \cdot I$$

$$\frac{dR}{dt} = \gamma \cdot I - \mu \cdot R$$

$$d \log(\beta(t)) = \sigma \cdot dB(t)$$



Application to Dengue in Phnom Penh



Concluding remarks

- **It is important to take into account non-stationarity when analyzing epidemiological datasets.**
- Wavelet analysis is one of the adapted statistical approaches considering these features of epidemiological time series
- Time-varying parameters modeled with a diffusion process is an alternative to the use of more complex model.
- Models with time-varying parameters can be easily used to predict an epidemic in real time.

Accounting for Non-Stationarity in Epidemiology

Thanks to all my collaborators and particularly to

- Mario Chavez from ICM, Paris and Mercedes Pascual from New-York University
- The members of the Eco-Evolutionary Mathematics team at IBENS
- Kévin Cazelles for all the R graphs and for the *future* Wavelet package in R





Maladies transmises par les moustiques, météo et climat : des liaisons dangereuses

Publié: 4 octobre 2023, 20:39 CEST

Aedes aegypti est une espèce d'insectes diptères, un moustique qui est le vecteur principal de la dengue, de l'infection à virus Zika, du chikungunya et de la fièvre jaune. U.S. NAVY

Partager par e-mail

X (anciennement Twitter)

Facebook 87

LinkedIn

Imprimer

Lors de la crise sanitaire liée au SARS-CoV-2, nos sociétés ont pris conscience de l'importance et de l'utilité des outils mathématiques et statistiques pour caractériser la propagation d'une maladie dans la population générale, prévoir ses conséquences en termes de santé publique et anticiper les répercussions économiques à court terme. Au-delà du Covid-19, les maladies propagées par les moustiques, dont l'aire de répartition s'élargit régulièrement sous l'influence du changement climatique, représentent aujourd'hui une menace émergente.

auteur



Bernard Cazelles
Ecologie, Epidemiologie,
Sorbonne Université

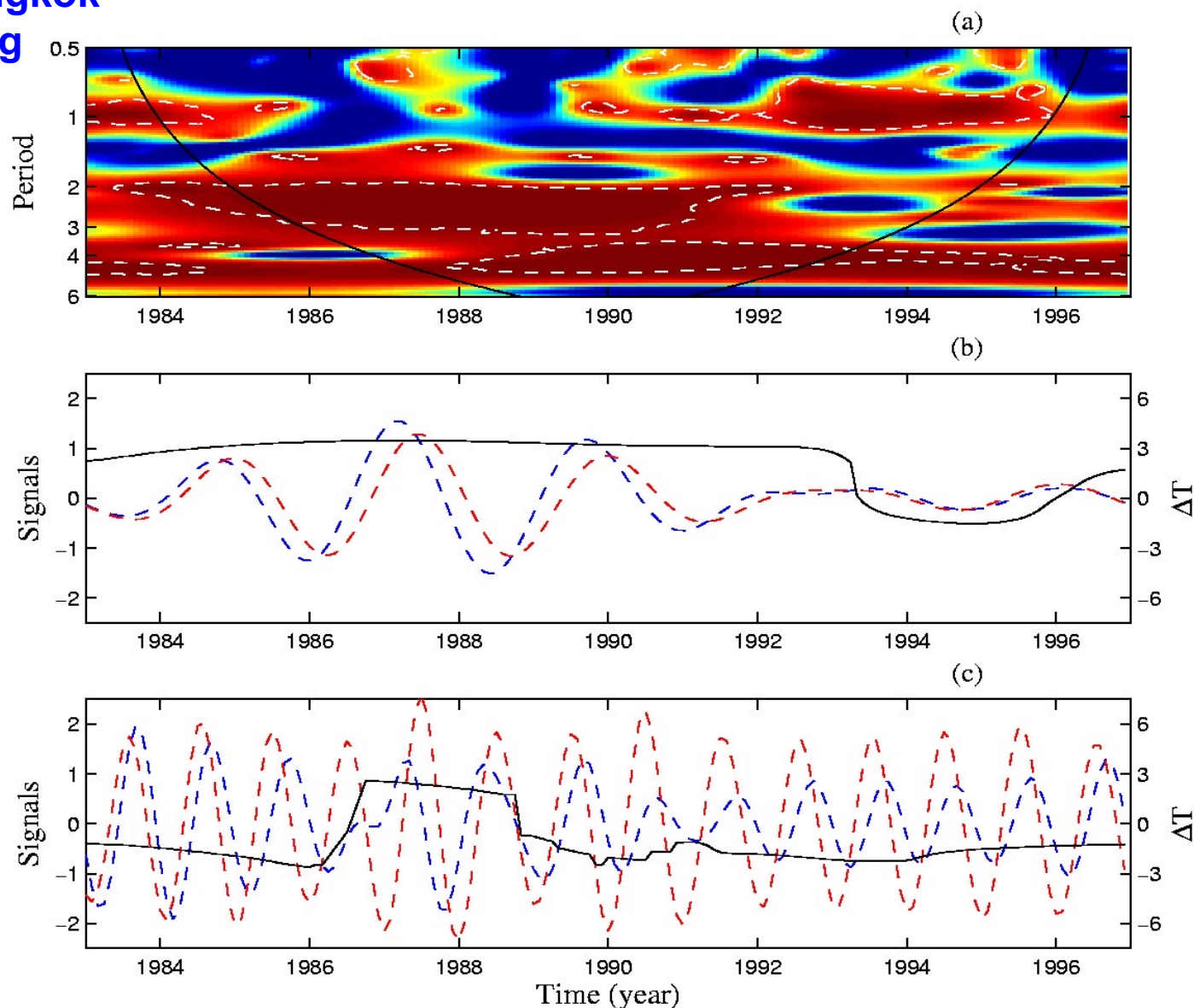
Déclaration d'intérêts

Bernard Cazelles a reçu des financements de l'Agence National de la Recherche (ANR).

Partenaires

Wavelet Analysis

Relation between Bangkok and the remaining Thailand



Importance
of non-
stationarity

Disentangling the

Whole Thailand: Dengue, Rainfall, SOI.

Application to Dengue In Thailand

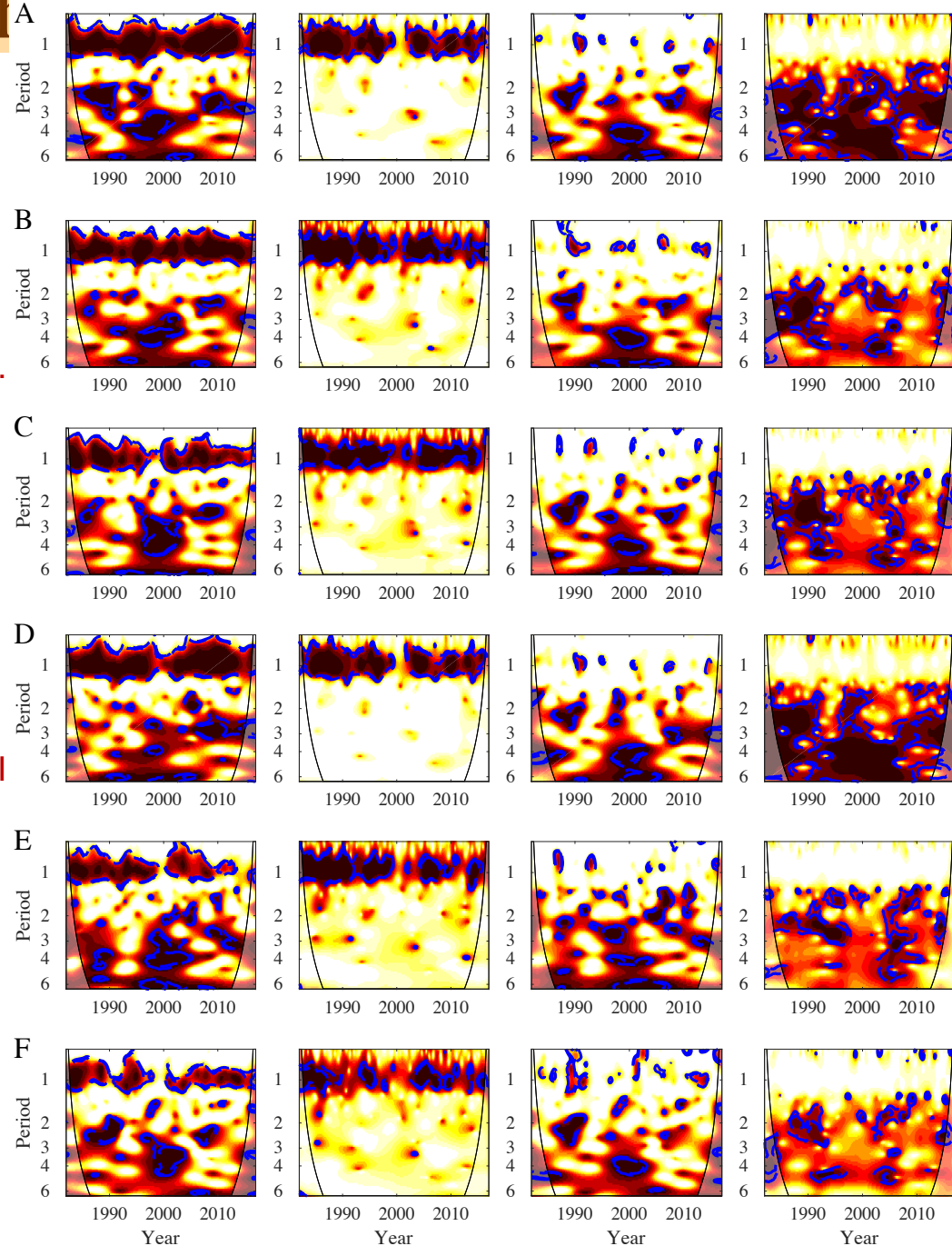
Northern Thailand: Dengue, Temperature MEI.

Central Thailand: Dengue, Temperature, ONI.

North-Eastern Thailand: Dengue, Temperature, ONI

Southern Thailand: Dengue, Rainfall, SOI.

Bangkok: Dengue, Temperature (min), MEI.



Monarity in Statistical Analysis

Tea Plantation (AHP): Rainfall (Kaisugu), ONI.

Application to Malaria In Kenya and Ethiopia

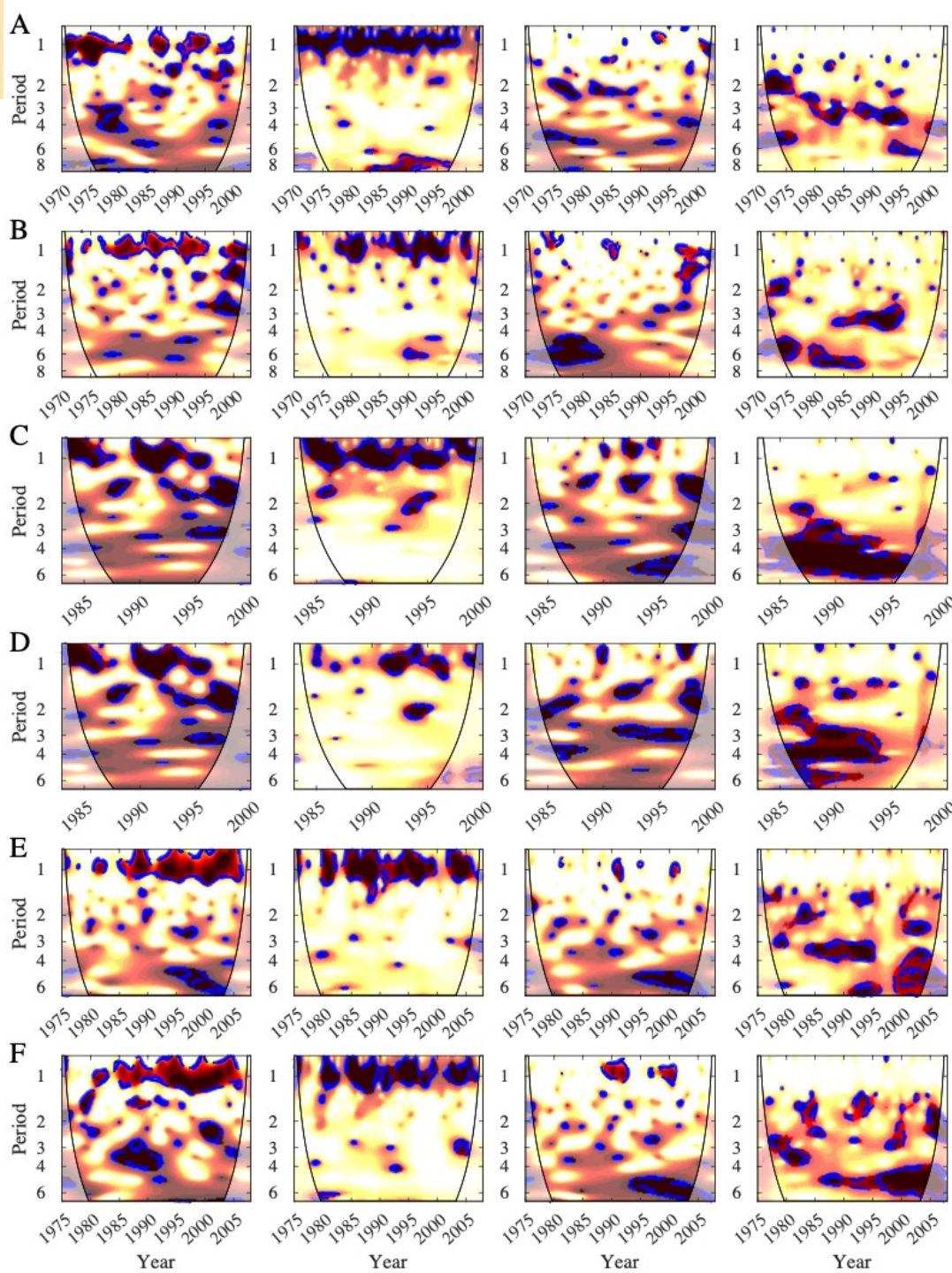
Tea Plantation (BBF): Rainfall (Kaisugu); DMI.

Kapsabet district: Rainfall, Nino3.

Kisii district: Rainfall, DMI.

Ethiopia: Temperature (min), SOI.

Ethiopia: Temperature (max), MEI.



Application to Dengue in Phnom Penh

