

Climate forcing and malaria dynamics

Mercedes Pascual

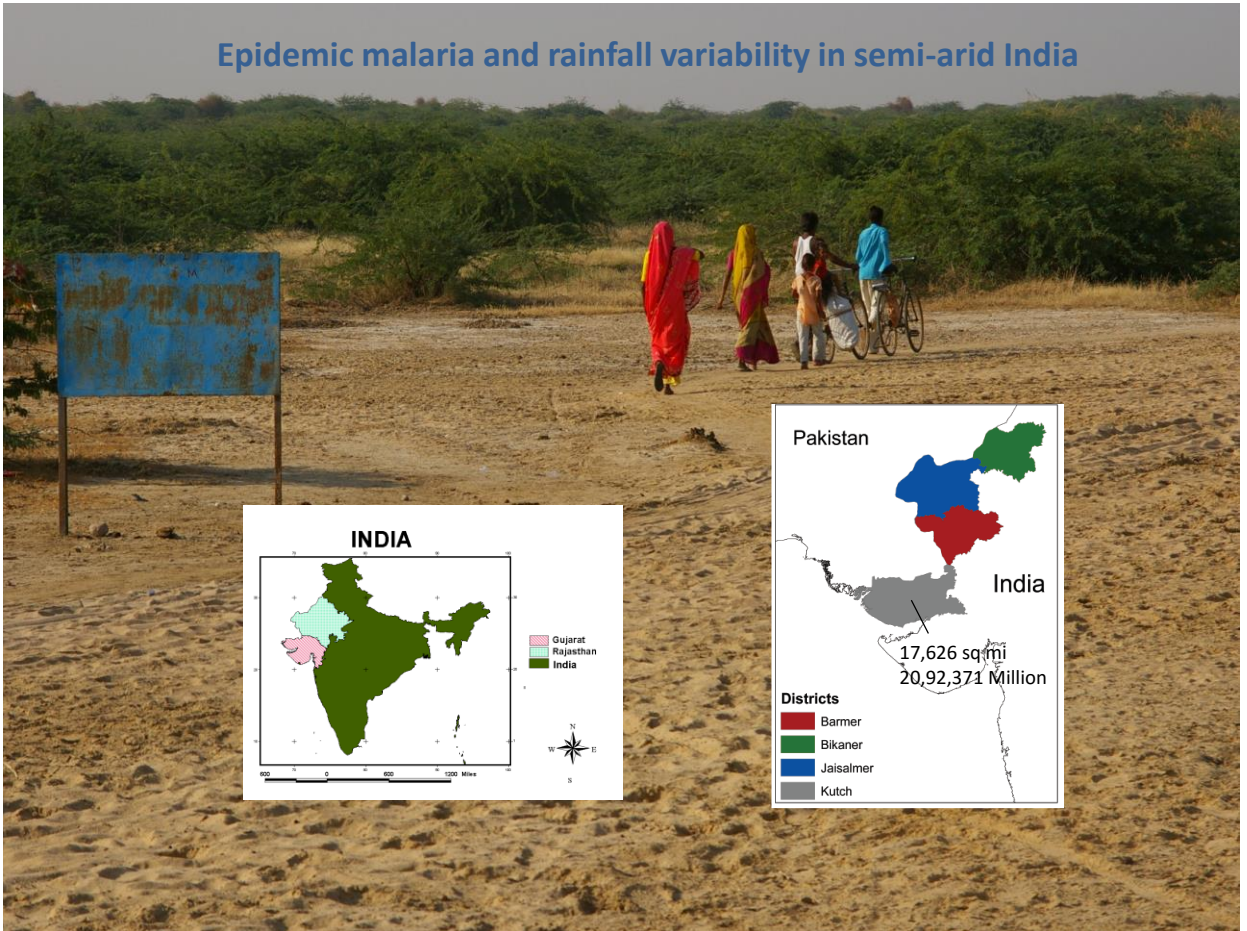
University of Chicago

and

The Santa Fe Institute

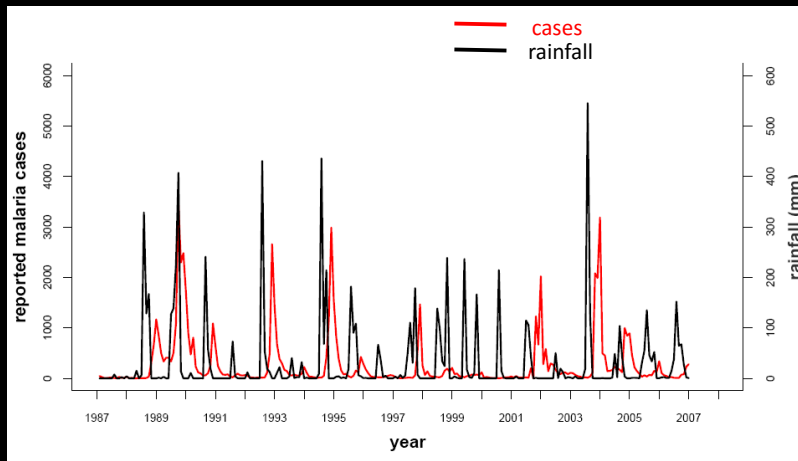


Epidemic malaria and rainfall variability in semi-arid India



Typical epidemic behavior of *P. falciparum* cases

District of Kutch:
30 years
monthly cases

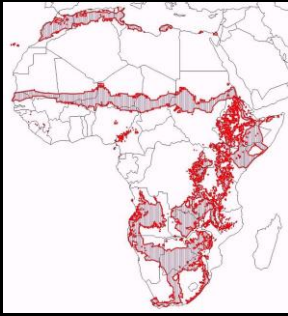


Laneri *et al.* PLoS Computational Biology 2010

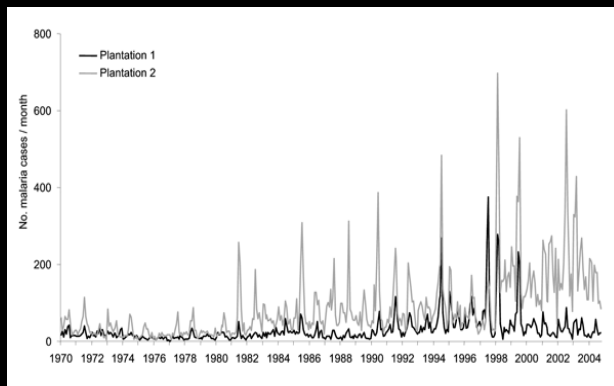
Highland malaria and climate change

~ 110 million Africans live in areas at risk of epidemic malaria

Estimated 110 000 deaths each year (Africa Malaria Report)

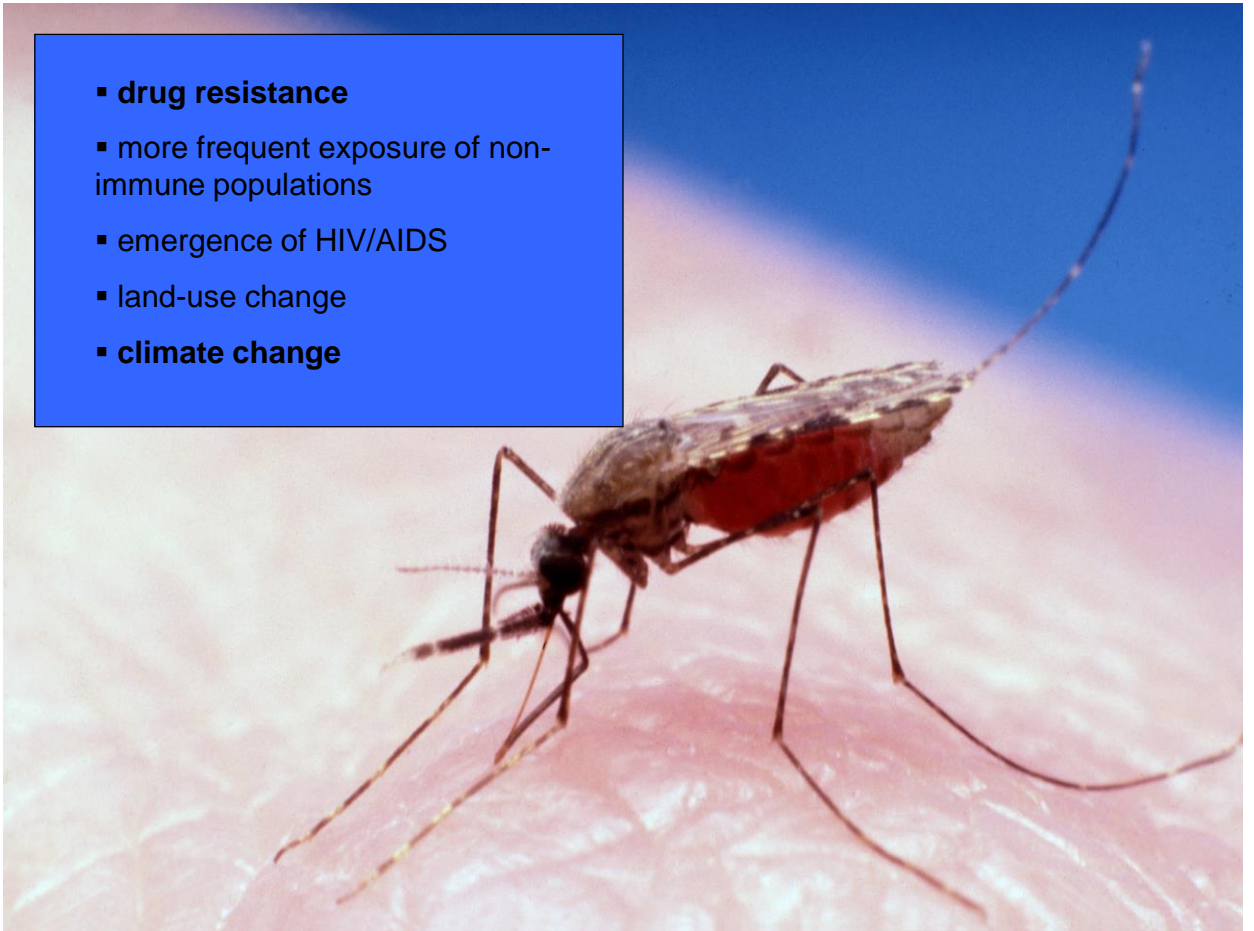


Areas at risk of epidemic malaria
From Grover-Kopec et al, Mal. J. 2005



From Shanks et al. EID 2005

- **drug resistance**
- more frequent exposure of non-immune populations
- emergence of HIV/AIDS
- land-use change
- **climate change**

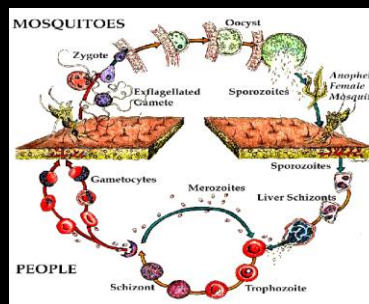


Testing hypotheses on disease dynamics and climate forcing by comparing mechanistic models

Best disease models
with no climate



Best disease models
with climate variability

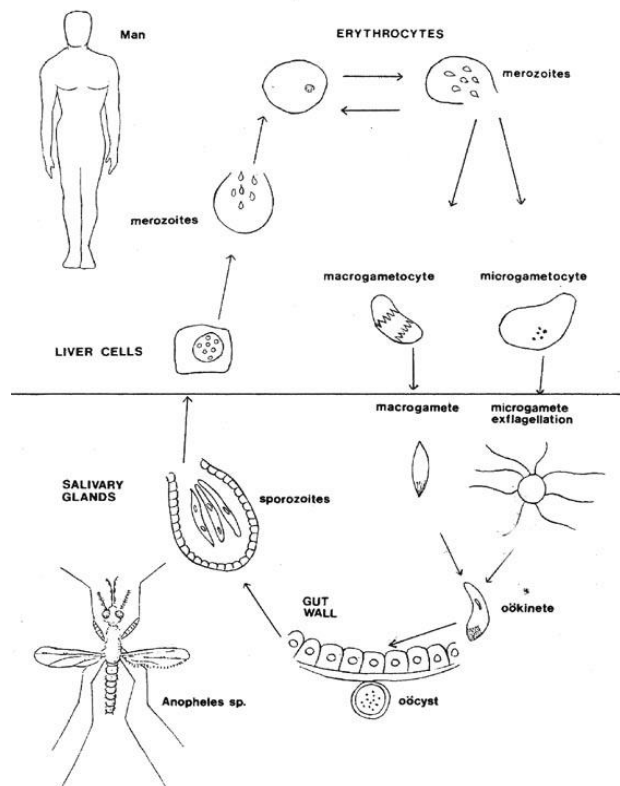


Conceptual outline

- The effect of climate forcing will be most apparent where climate factors act as strong limiting factors (at the edge of the spatial distribution of the disease, in highland and semi-arid regions). But here, by definition, transmission is low, and therefore, population immunity, is most unlikely to play a strong dynamical role.
- We will see that epidemiological processes matter primarily at seasonal and not interannual scales, and that 'reactive control' can act as a nonlinear feedback and generate multiannual cycles.
- Prediction needs to take into account non-stationary conditions.

Model by Ross and McDonald (1916-1957)

- proportion of the human population infected
- proportion of the female mosquito population infected



Ross-McDonald model:

Proportion mosquitoes infected, y

$$\frac{dx}{dt} = \left(\underset{\substack{\text{Success of bites} \\ \text{Biting rate}}}{abM} / \underset{\substack{\text{Number of mosquitoes} \\ \text{Number of hosts}}}{N} \right) y(1-x) - \underset{\text{Recovery rate}}{r}x$$

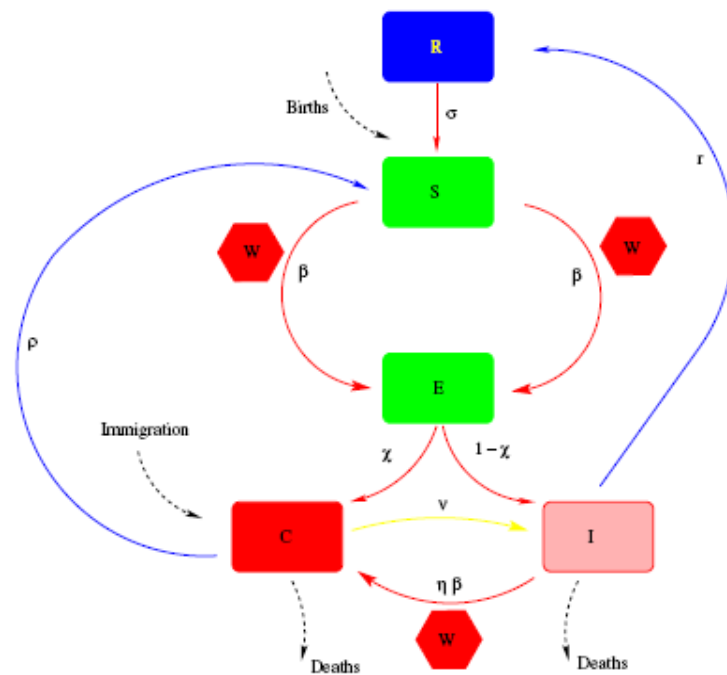
$$\frac{dy}{dt} = \underset{\text{Biting rate}}{ax}(1-y) - \underset{\text{Mosquito death rate}}{\mu}y$$

Proportion humans infected, x

Coupled mosquito-human transmission model

- Larvae
- Adults in three classes:

uninfected
exposed
infectious



Alonso, Bouma and Pascual, *Proc. R. Soc. London B* 2011

Human model: β , the force of infection

$$\frac{dS}{dt} = B - \beta S + \sigma R - \delta S + \rho C$$

$$\frac{dE}{dt} = \beta S - \delta E - \gamma E$$

$$\frac{dI}{dt} = (1 - \xi)\gamma E - \eta\beta I + \nu C - r I - \delta I$$

$$\frac{dR}{dt} = -\sigma R + r I - \delta R$$

$$\frac{dC}{dt} = \xi\gamma E + \eta\beta I - \nu C - \rho C - \delta C$$

$$\beta = b a \frac{W}{H} + \beta_0$$

Mosquito model: Population dynamics

L , larval stage and M adult stage

$$\begin{aligned}\frac{dL}{dt} &= f M \left(\frac{K-L}{K} \right) - \delta_L L - d_L L \\ \frac{dM}{dt} &= d_L L - \delta M\end{aligned}$$

Ahumada and Dobson 2009

Mosquito carrying capacity is controlled by water availability...

$$\frac{dK}{dt} = K_A P - K_E K$$

Mosquito sub-model:

$$\begin{aligned}\frac{dL}{dt} &= f M \left(\frac{K - L}{K} \right) - \delta_L L - d_L L \\ \frac{dX}{dt} &= -c a y X - \delta_M X + d_L L \\ \frac{dV}{dt} &= +c a y X - \gamma_P V - \delta_M V \\ \frac{dW}{dt} &= \gamma_P V - \delta_M W\end{aligned}$$

where y is the fraction of infectious humans:

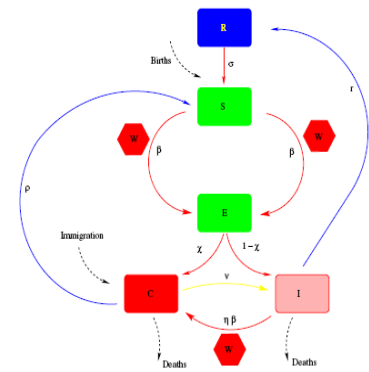
$$y = \frac{C + I}{H}$$

Temperature

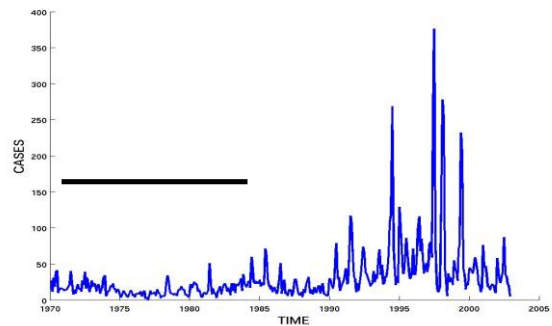


Rainfall

- larva development (T)
- Plasmodium development (T)
- Adult and larval survival (T, R)
- Gonotrophic Cycle (biting rate, T)
- Carrying capacity (R)



See E. Mordecai, *Ecology Letters* 2013:
Optimal temperature for malaria
transmission is dramatically lower than
previously predicted



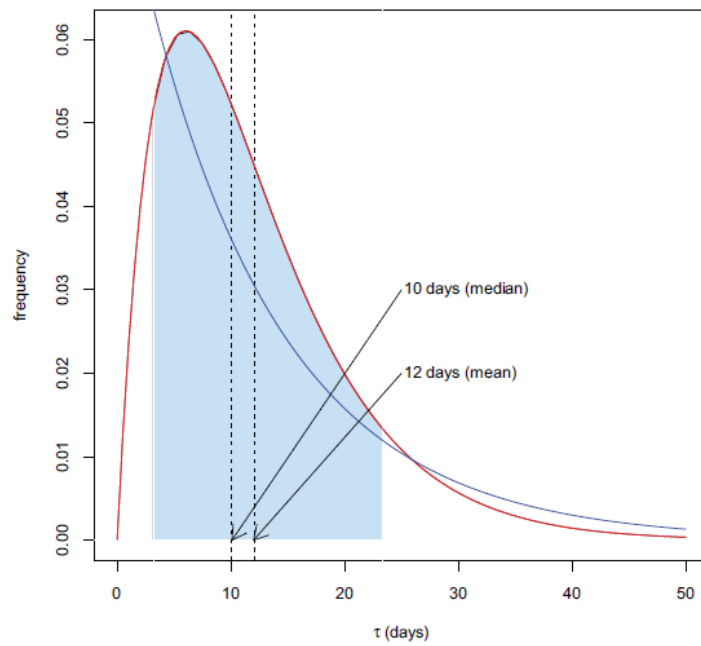
Alonso, Bouma and Pascual, *Proc. R. Soc. London B* 2011

$$\begin{aligned}
\frac{dS}{dt} &= B - \beta S + \sigma R - \delta S + \rho C \\
\frac{dE_1}{dt} &= \beta S - \delta E_1 - n_H \gamma_H E_1 \\
\frac{dE_2}{dt} &= n_H \gamma_H E_1 - n_H \gamma_H E_2 - \delta_H E_2 \\
&\dots = \dots \\
\frac{dE_n}{dt} &= n_H \gamma_H E_{n-1} - n_H \gamma_H E_n - \delta_H E_n \\
\frac{dI}{dt} &= (1 - \xi) n_H \gamma_H E_n - \eta \beta I + \nu C - r I - \Psi I - \delta I \\
\frac{dR}{dt} &= -\sigma R + r I - \delta R \\
\frac{dC}{dt} &= \xi n_H \gamma_H E_n + \eta \beta I - \nu C - \rho C - \alpha C - \delta C
\end{aligned}
\tag{\delta = \delta_H}$$

$$f_H(\tau) = \frac{n_H \gamma_H}{\Gamma(n_H)} \exp(-n_H \gamma_H \tau) \tau^{n_H-1}$$

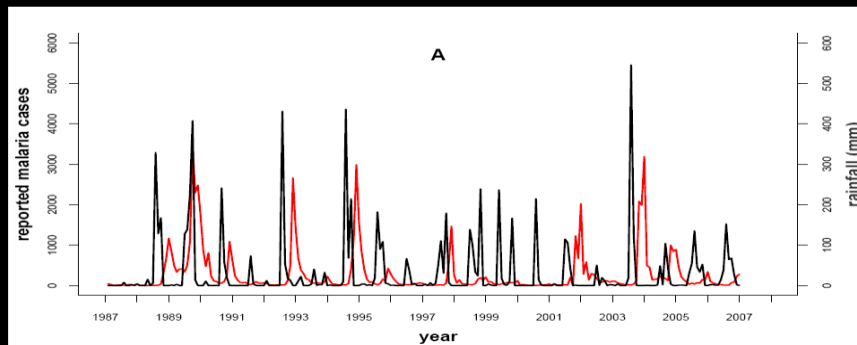
of average $1/\gamma_H$ and variance $1/(n_H \gamma_H^2)$.

Gamma distributed 'incubation' time

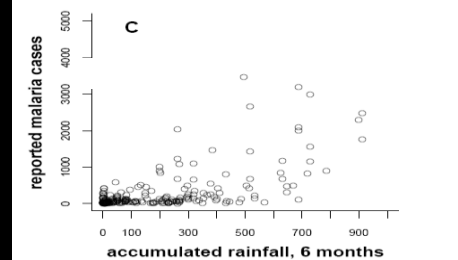


$$\begin{aligned}
\frac{dL}{dt} &= f M \left(\frac{K - L}{K} \right) - \delta_L L - d_L L \\
\frac{dX}{dt} &= -c a y X - \delta_M X + d_L L \\
\frac{dV_1}{dt} &= +c a y X - n_P \gamma_P V_1 - \delta_M V_1 \\
\frac{dV_2}{dt} &= n_P \gamma_P V_1 - n_P \gamma_P V_2 - \delta_M V_2 \\
&\dots \dots \dots \\
\frac{dV_n}{dt} &= n_P \gamma_P V_{n-1} - n_P \gamma_P V_n - \delta_M V_n \\
\frac{dW}{dt} &= n_P \gamma_P V_n - \delta_M W
\end{aligned}$$

A simple 'coupled' model: malaria in Kutch, India

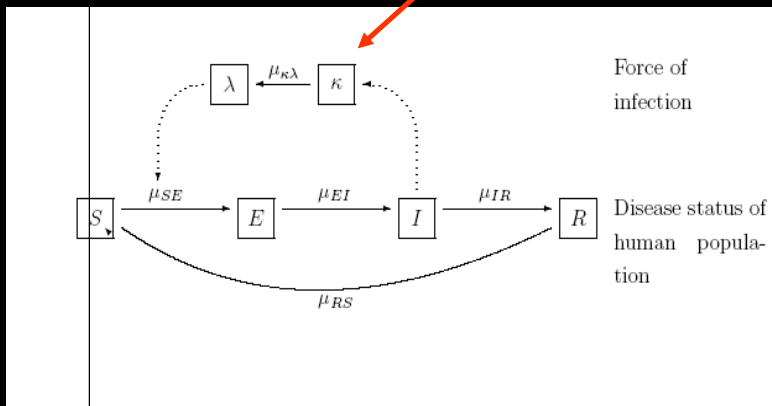


- does population immunity play a role in the response to climate variability?
- how predictable is the size of outbreaks based on transmission models driven by climate?



Malaria model

$$f(t) = \frac{I(t)}{N(t)} \exp\{\beta_{seas} + \beta \cdot Rain(t)\} Noise$$



Latent force of infection

$$\kappa = \lambda_0$$



$$\lambda_1$$



$$\lambda_2$$



$$\lambda$$

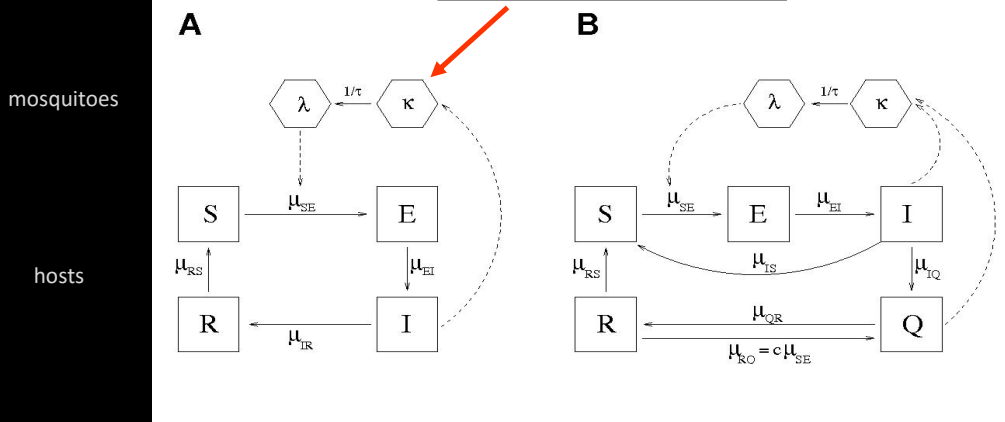
Force of infection

Parasite's development in surviving mosquitoes

Two possible structures for human component

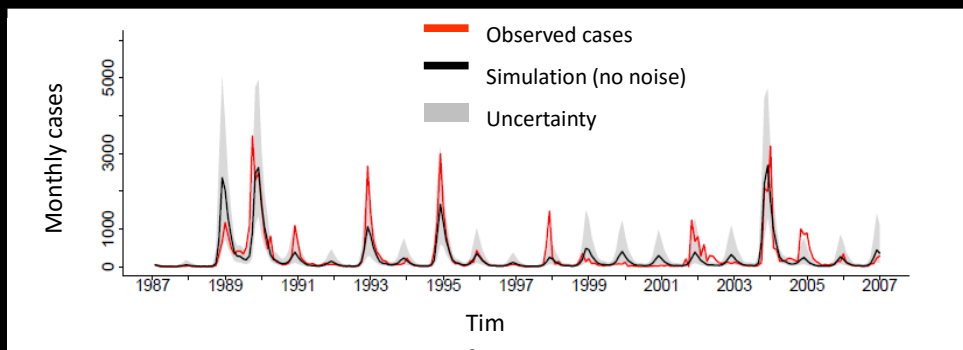
Force of infection:
a function of rainfall

$$f(t) = \frac{I(t)}{N(t)} \exp\{\beta_{seas} + \beta \cdot \text{Rain}(t)\} \text{Noise}$$



Both rainfall and clinical immunity are included in the 'best' model

- Clinical immunity is important at seasonal scales
- This model outperforms a 'standard' non-mechanistic, linear autoregressive, model that includes rainfall



Laneri *et al.* PloS Computational Biology 2010
Bhadra *et al.* J. American Statistical Association 2011

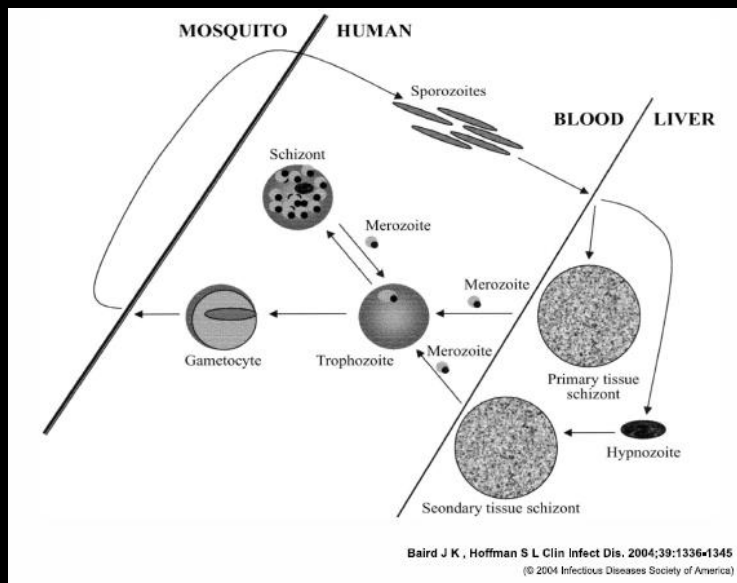
Model comparison

Table S1. Table of log-likelihood (ℓ) and AIC of the fitted models for Kutch and Barmer.

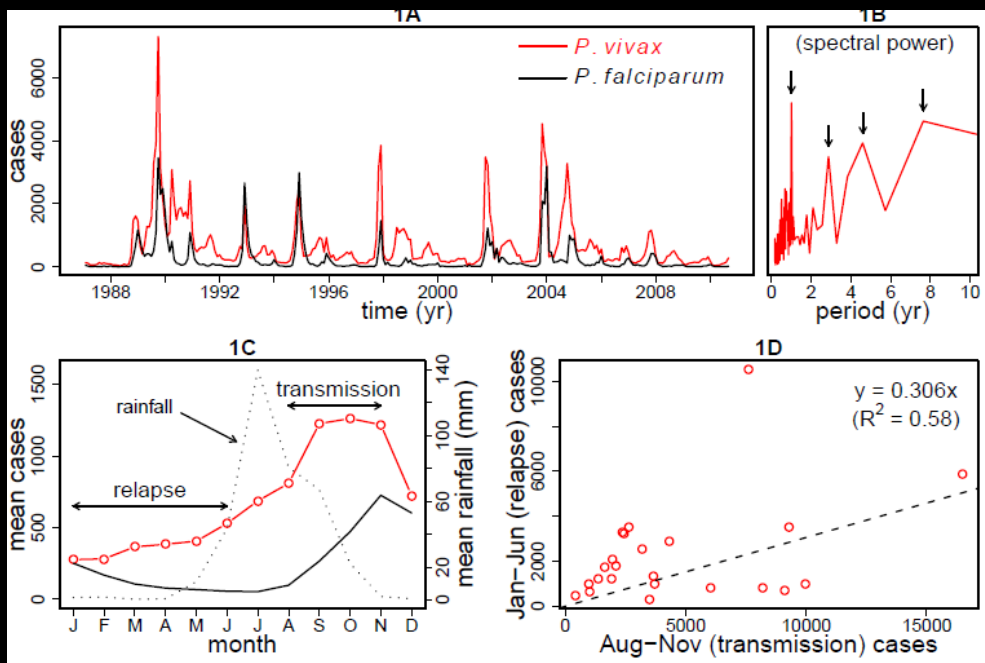
model	p	log-likelihood (ℓ)		AIC	
		Kutch	Barmer	Kutch	Barmer
VSEIRS model without rainfall	19	-1275.0	-984.1	2588.0	2006.2
VSEIRS model with rainfall	20	-1265.0	-978.6	2570.0	1997.2
VS^2EI^2 model without rainfall	24	-1261.1	-975.3	2570.2	1998.6
VS^2EI^2 model with rainfall	25	-1251.0	-970.5	2552.0	1991.0
SARIMA $(1, 0, 1) \times (1, 0, 1)_{12}$ without rainfall	6	-1329.0	-983.7	2670.0	1979.4
SARIMA $(1, 0, 1) \times (1, 0, 1)_{12}$ with rainfall	7	-1322.6	-977.0	2659.2	1968.0

In the table “ p ” denotes the number of parameters for each model. AIC is computed by the formula $AIC = -2\ell + 2p$. The SARIMA model was fitted to the data on the log scale (see the supplement of [2] for a detailed description of this procedure).

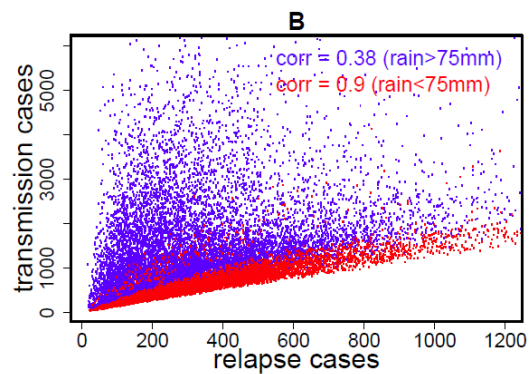
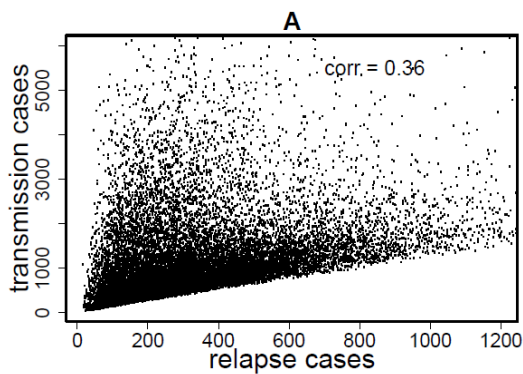
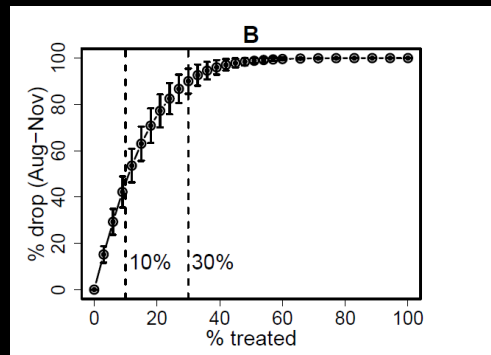
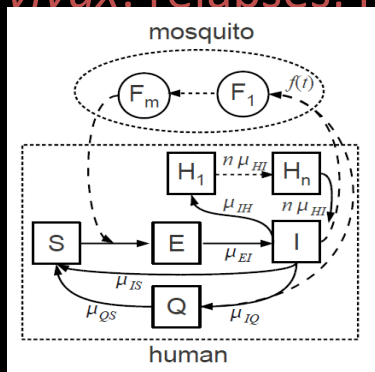
P. vivax malaria : relapses, rainfall and treatment



Inference on importance and duration of relapses for the population dynamics of the disease
Potential implications for treatment that focuses on this stage of the disease



P. vivax: relapses, rainfall, and treatment

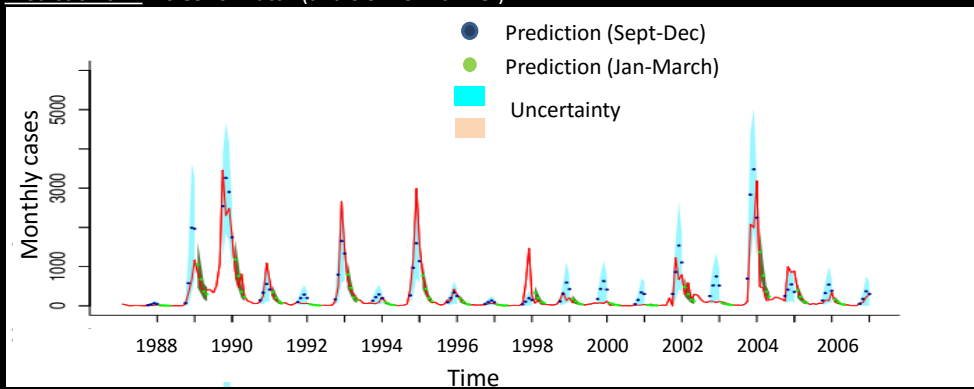


Roy *et al*, PLoS Neglected Tropical Diseases, *PLoS NTD*

Prediction

The rainfall-driven transmission model exhibits high prediction skill (retrospectively)

Prediction skill = 0.89 for Kutch (and 0.92 for Barmer)



Prediction performance: 4 months

$$\text{skill} = 1 - \frac{\sum_{i=1987}^{2006} (y_i - \hat{y}_i)^2 w_i}{\sum_{i=1987}^{2006} (y_i - \mu)^2 w_i}$$

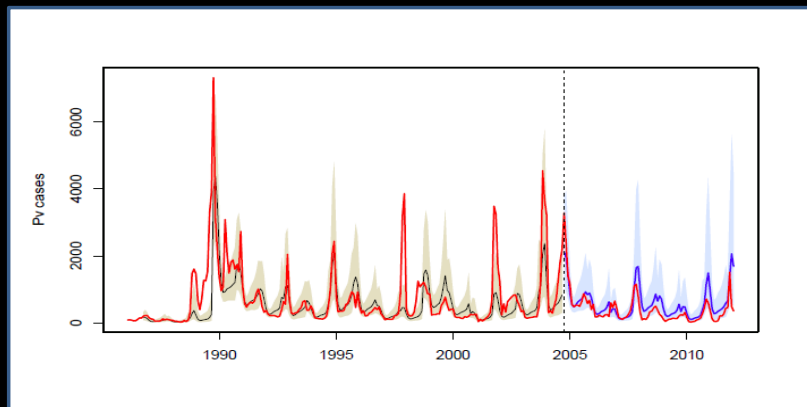
- w_i inverse of the prediction variance for the year i
- \hat{y}_i and y_i are the predicted and observed cases, accumulated over September to December for the year i
- μ : 20 year mean of the observed cases accumulated between September and December

skill \rightarrow 1 Good Prediction

skill \rightarrow 0 Bad Prediction

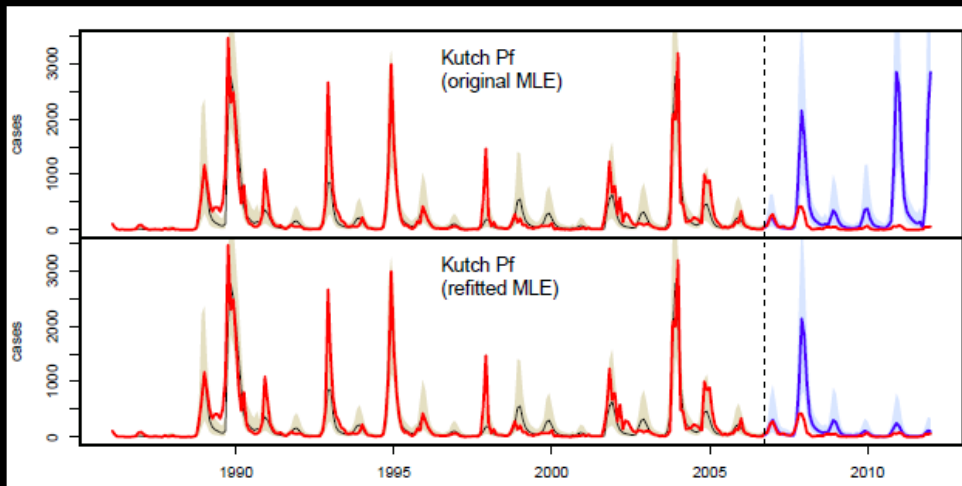
Predictability

- High prediction skill retrospectively (e.g. 0.9 for *P. falciparum* in Kutch)
- Also prospectively: illustrated here for *P. vivax*

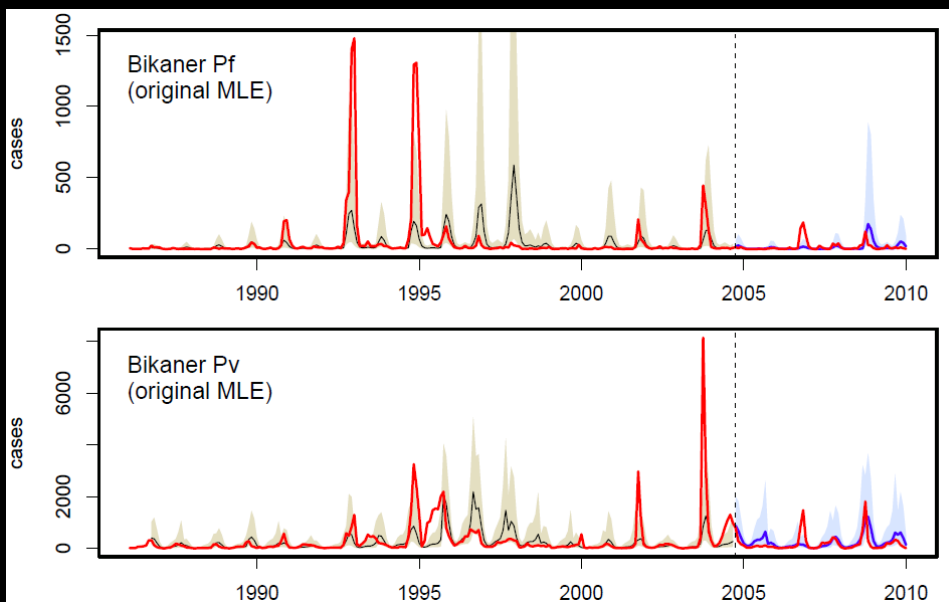


Roy *et al.*, *in review*

“Prediction” in the presence of non-stationarity



In this other district, we can see that the recent decrease in cases can completely be explained by the lack of rains



Ocean temperatures in the Tropical South Atlantic influence malaria epidemics in NW India

Sea Surface
Temperatures
(Atlantic)

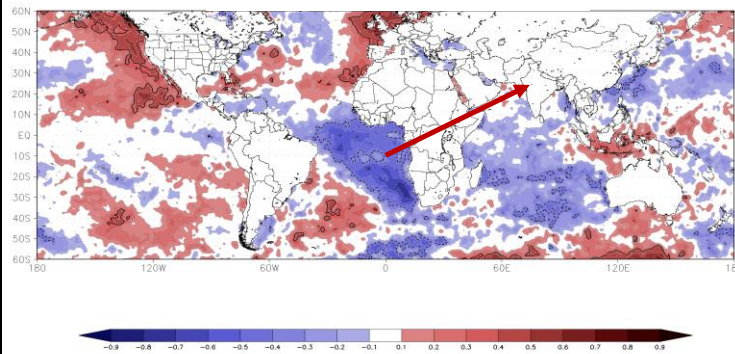


Rainfall
NW India

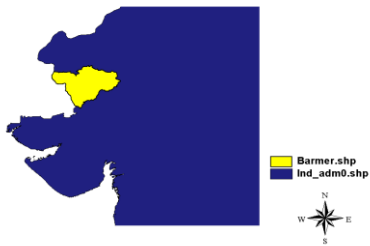
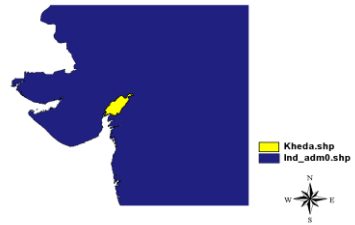


Malaria risk

Lag (ranked) correlation between
Kutch cases in **October** and
Sea Surface Temperatures in **June**

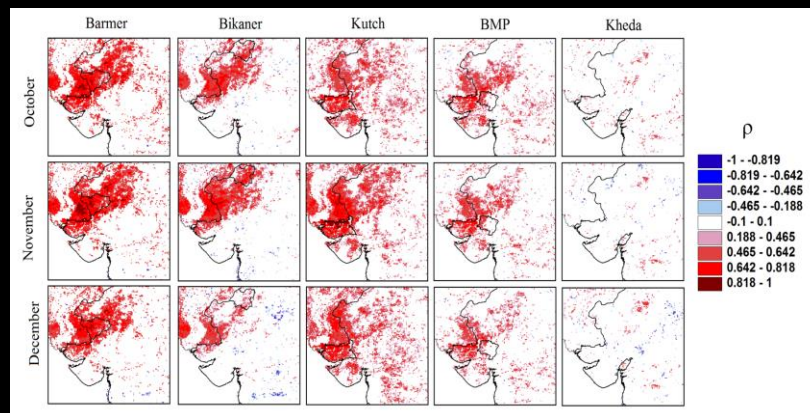


Cash *et al.* Nature Climate Change 2013



Association with climate breaks down along an irrigation gradient

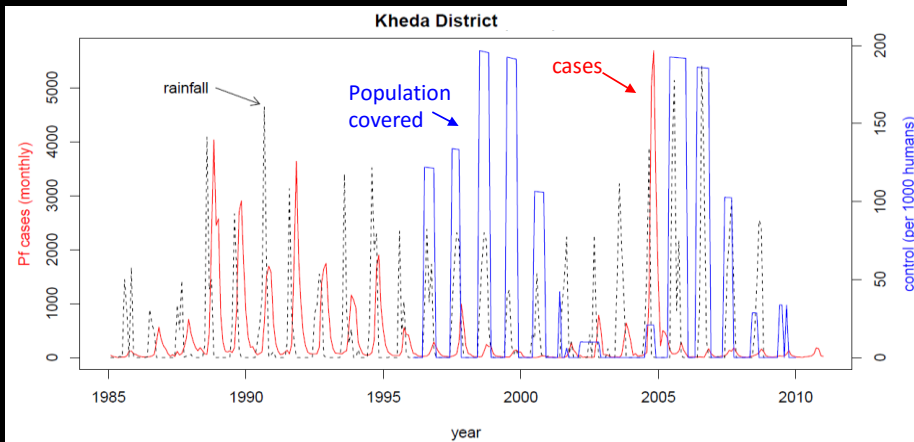
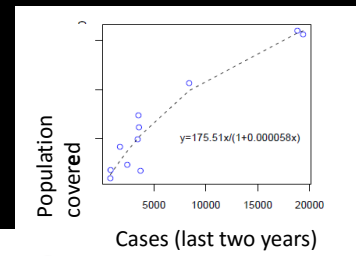
More irrigated land (more mosquito habitat / more wealth)



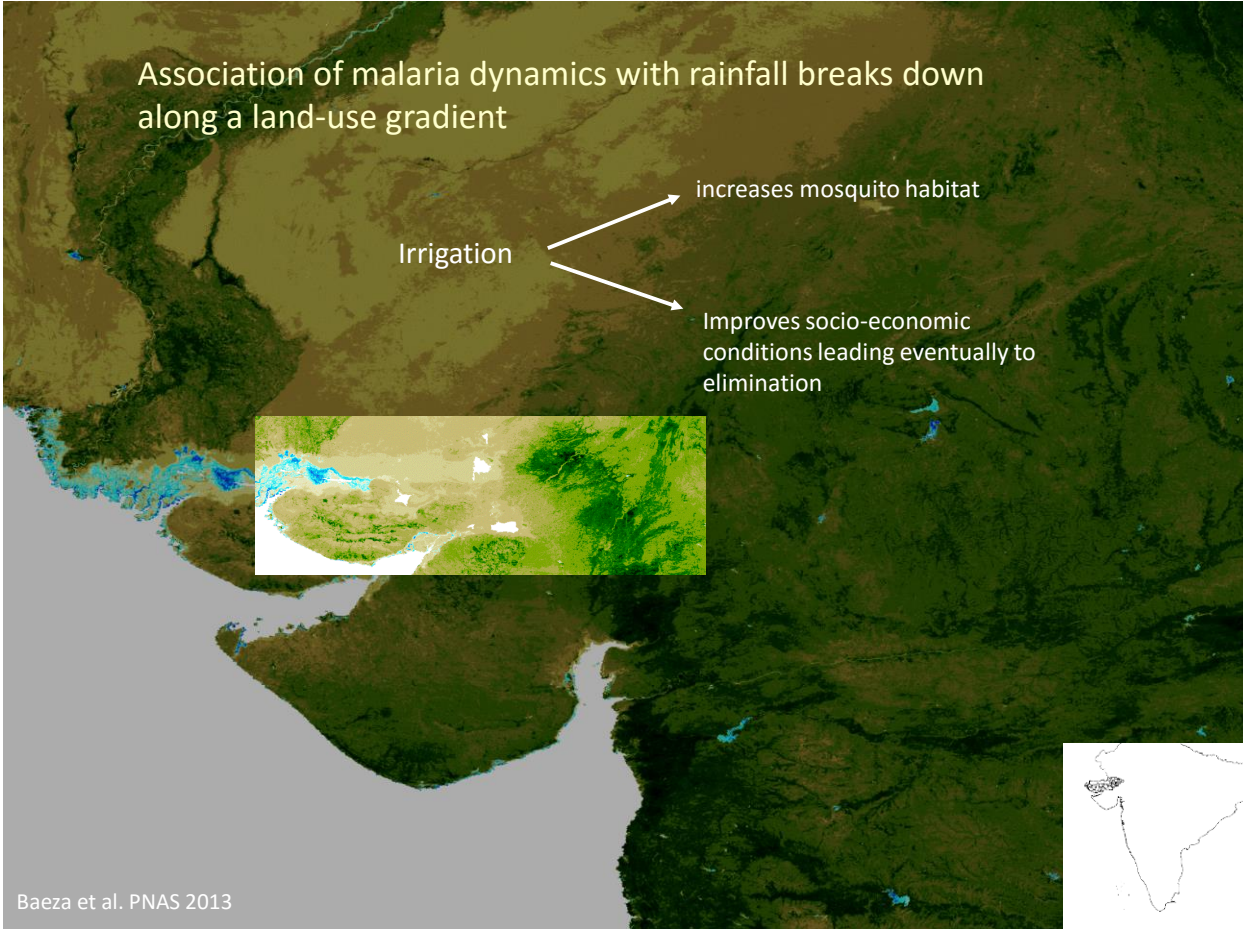
Rank correlation maps with vegetation index from remote sensing

Baeza *et al.*, Malaria Journal 2011

“Reactive” control policy
generates cycles and unexpected epidemics,
precluding elimination



Baeza *et al.* Acta Tropica 2013
Baeza *et al.*, PNAS 2013

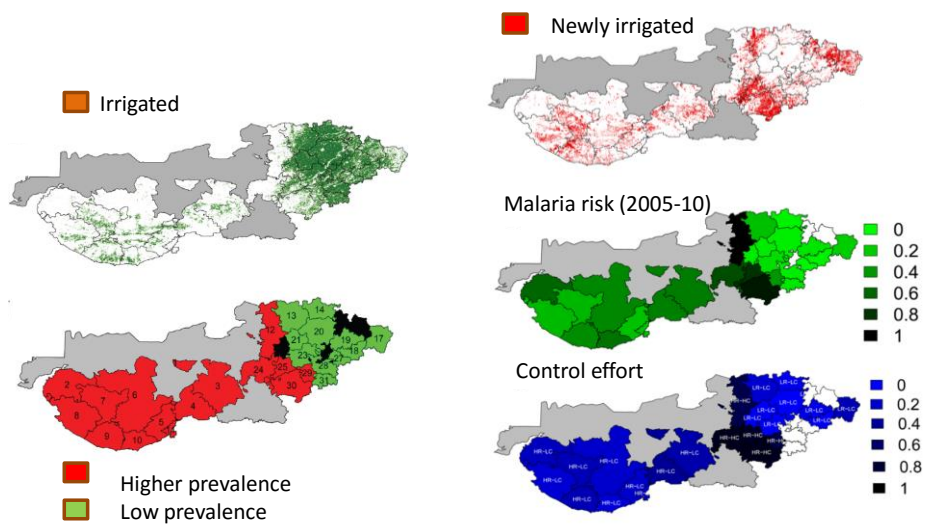


22 Talukas (sub-districts) from Gujarat State

- Confirmed monthly cases of *Plasmodium falciparum* and *P. vivax* [1997-2011]
- IRS (Indoor Residual Spray) application (population covered) [2000-2010]

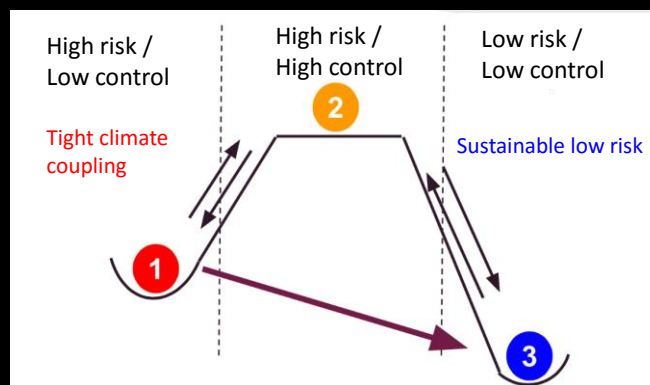


Transition between epidemic malaria and elimination can be long-lasting
(more than a decade) despite forceful control efforts



Baeza *et al.* PNAS 2013

Three distinct regimes: the transition regime can be long lasting (over a decade)

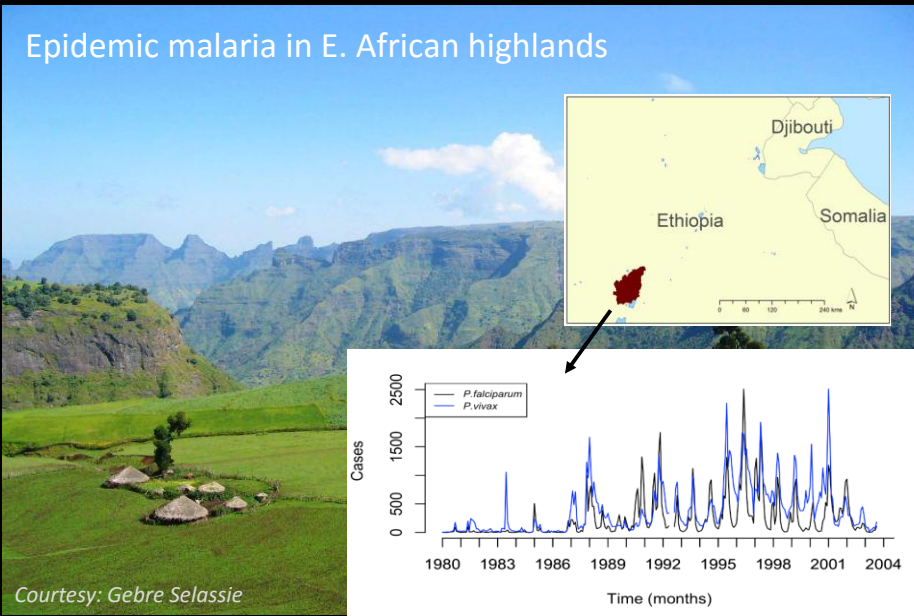


Baeza *et al.* PNAS 2013

So far:

- Clear signature of climate forcing in epidemic regions.
- Nonlinear responses are not seen in terms of cycles. The depletion of the resource and therefore the strength of 'competition' for hosts is too low.
- Consideration of population dynamics (including immunity) remains important, especially for persistence during inter-epidemic periods.
- Interannual cycles can be generated when the epidemiological system includes intervention feedbacks, and these cycles can interact with climate anomalies to delay or impede elimination.

Epidemic malaria in E. African highlands



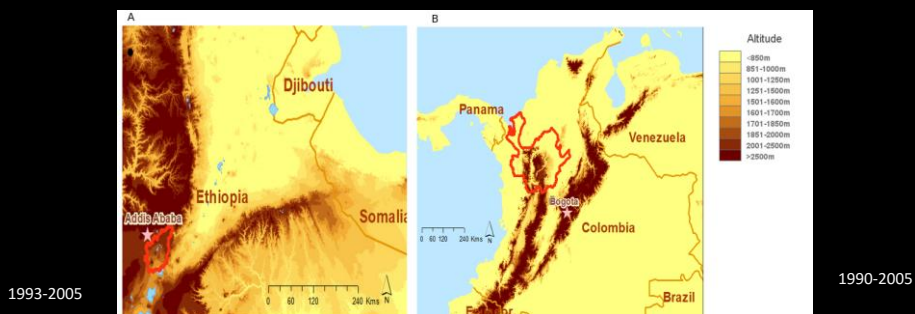
Courtesy: Gebre Selassie



Climate change vs.

- Evolution of drug resistance (Shanks et al. EID 2005)
- More frequent exposure of non-immune populations
- Emergence of HIV/AIDS
- Land-use change
- Breakdown of public health systems

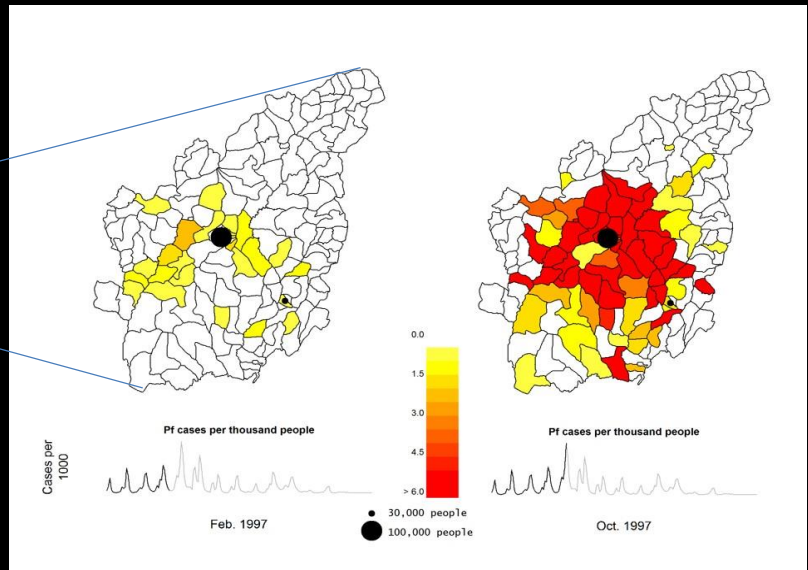
Taking advantage of high-resolution spatio-temporal data to address climate change



Confirmed monthly cases before major interventions of last decade

Siraj, Santos *et al.*, Science 2014

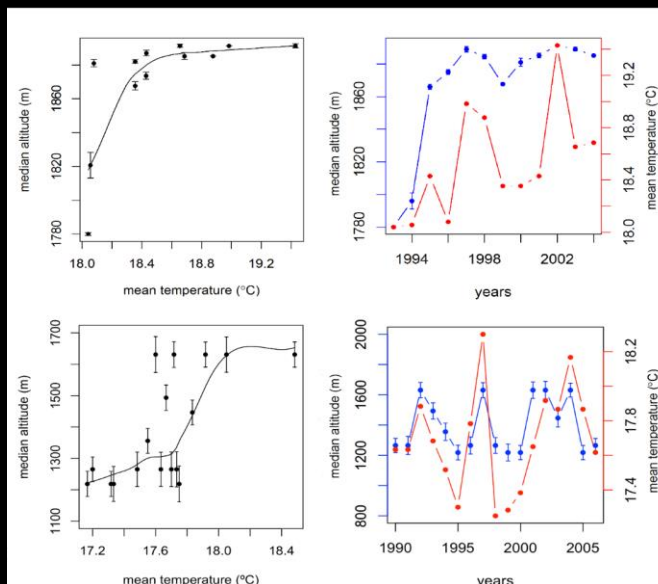
Expansion of the spatial distribution



Siraj, Santos-Vega *et al.*, *Science* 2014

The spatial distribution of the disease expands upwards in warmer years

Ethiopia



Colombia

Siraj, Santos-Vega *et al.*, *Science* 2014

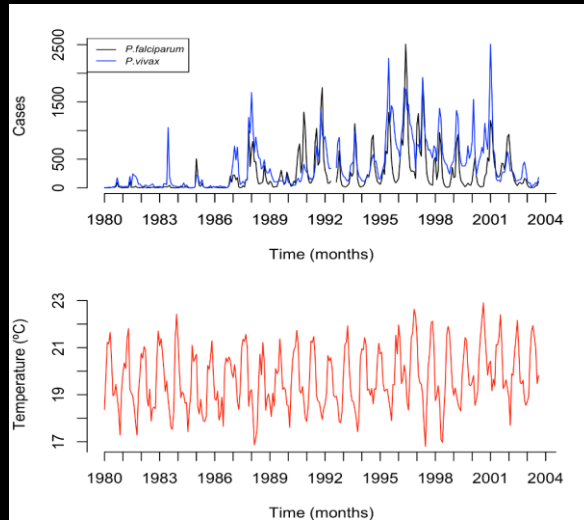
Is the long-term trend consistent with the magnitude of the altitudinal expansion?

From movement in altitudinal distribution

⇒ ~ 1980 cases / degree C

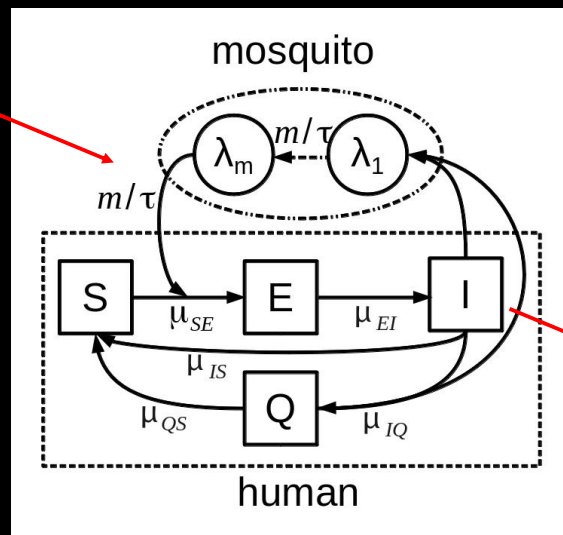
From longer temporal trend

⇒ ~2166 cases / degree C

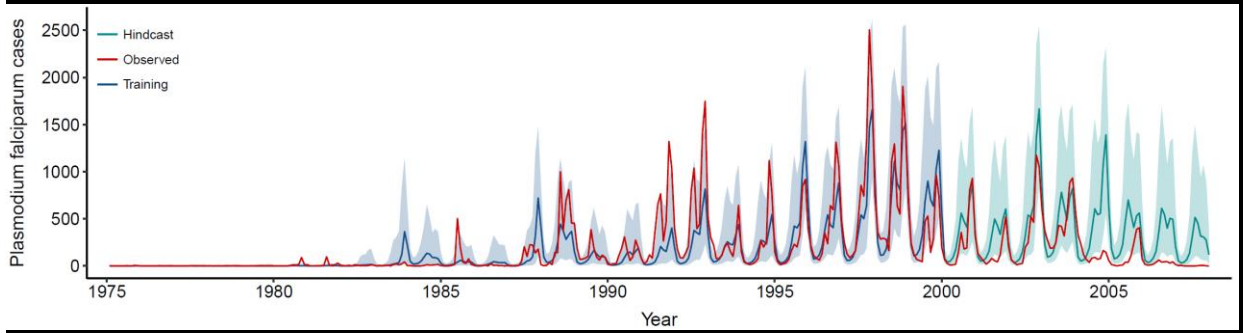


Transmission model

Force of Infection
(depends on
temperature,
season,
infection levels
and noise)



Likelihood maximization by iterated filtering



Pascual *et al.*, *in prep.*

Gracias



Andres Baeza



Anindya Bhadra

Menno Bouma LSHTM



Ben Cash (COLA; IGES); Xavier Rodo (IC3); and Manojit Roy (UM)



Graham Environmental Sustainability Institute (GESI, UM)
NOAA, Oceans and Health