IDENTIFY AND PREDICT ENVIRONMENTAL CHANGE EFFECTS ON TIGER MOSQUITOS, *AEDES POLYNESIENSIS*

AUTHORS

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SUMMARY

To control mosquito populations for managing vector-borne diseases, a critical need is to identify and predict their response to causal environmental variables. However, most existing attempts rely on linear approaches based on correlation, which cannot apply in complex, nonlinear natural systems, because correlation is neither a necessary nor sufficient condition for causation. Appling empirical dynamic modelling that acknowledges nonlinear dynamics on nine subpopulations of tiger mosquitos from three neighbouring reef islets of the Raiatea atoll, we identified temperature, precipitation, dew point, air pressure, and mean tide level as causal environmental variables. Interestingly, responses of subpopulations in close proximity (100-500 m) differed with respect to their causal environmental variables and the time delay of effect, highlighting complexity in mosquito-environment causality network. Moreover, we demonstrated how to explore the effects of changing environmental variables on number and strength of mosquito outbreaks, providing a new framework for pest control and disease vector ecology.

INTRODUCTION

Mosquito-borne diseases contribute significantly to human mortality causing more than 1 billion cases and over 1 million deaths every year (GBP 2016). A case in point are mosquitoes of the genus *Aedes* being the main vector of dengue fever. For predicting disease outbreaks, population dynamics of mosquitoes are important concern. A critical need is therefore to understand and predict how environmental change may impact on fluctuation of mosquito populations (Duncombe et al., 2013).

There are two main strains of research aiming to examine how mosquito populations respond to climate changes. On the one hand, lab experiments are conducted to detect causal environmental drivers of mosquito survival, larval developmental time and other fitness parameters (e.g., Galun and Fraenkel, 1961; Couret et al., 2014). However, whether the findings in lab experiments can be generalized to natural systems remain unclear. On the other hand, time series data of mosquito abundance in natural systems are analysed for statistical associations with climate variables. For example, population abundance of the yellow fever mosquito *Ae. aegypti* was found to correlate with temperature, humidity, and precipitation (Scott et al., 2000; Azil et al., 2010; Barrera et al., 2011; Bashar et al., 2014; Alencar et al., 2015, Chaves et al., 2012). However, majority of the studies linking natural mosquito abundances with climate variables employed linear correlation analyses. While, correlation does not necessarily imply causation, lack of correlation does not imply lack of causation either (Sugihara et al., 2012; Chang et al., 2017). Applying linear methods to time series data generated from nonlinear processes bears the risk to detect mirage correlations, which may lead to wrong and misleading conclusions (Sugihara et al., 2012; Deyle et al., 2013; Deyle et al., 2016).

Indeed, mirage correlation (or non-stationary relationship) between mosquito populations and climate variables were often found in nature (Chaves et al., 2012; Simoes et al., 2013). Mirage correlation is commonly found, as a result of a fundamental property of nonlinear dynamical systems, known as state dependency (Sugihara et al 2012; Ye et al., 2015). State dependency means that the relationships among interacting variables change with different states of the dynamical system (Ye et al., 2015). The following example illustrates state-dependence in mosquitoes. It was shown experimentally that *Ae. aegyptii* females have lower vector competence for dengue virus in lower temperature environments only when the temperature is fluctuating; in comparison, without such a specific temperature combination (i.e. mosquitoes living in the environment of high temperature or constant temperature), the females have higher vector competence (Lambrechts et al., 2011;

Carrington et al., 2013). For *Aedes* mosquitoes, evidence abounds suggesting that nonlinear

population dynamics are common. For example, experimental studies show that larval

survival depends on larval density (Barrera et al., 2006), temperature (Couret et al., 2014),

and the mosquito's genomic background and microbiome (Mains et al., 2013) in a complex,

nonlinear way. In addition, abundance data of lab and field populations are best described

by nonlinear mathematical models (Dye 1984b; Legros et al., 2009; Mains et al., 2013). All

these studies suggest that, in general, mosquito-environment interactions are complex,

nonlinear, and inter-dependent; as such, linear methods are ill-posed for these questions.

Here, we aim at disentangling the causality network between climate variables and

mosquito populations and predict how environmental change may impact on the

populations. To overcome the limitation of linear approaches, we use the empirical

dynamical modelling (EDM) framework, which is specifically designed for analysing time

series data with underlying nonlinear dynamics (Ye et al., 2015; Deyle et al., 2016; Chang et

al., 2017). To demonstrate the efficacy of EDM framework for disease vectors, we analysed a

unique longitudinal data set of the Polynesian tiger mosquito, *Aedes polynesiensis*, which is

an important vector for lymphatic filariasis and dengue fever in the South Pacific (Lambdin

et al., 2009; Lau et al., 2016) (see supplementary material for more information on *Ae.*

polynesiensis). The data consist of biweekly (14-day interval) mosquito counts over two

years at nine sampling points on three motu islands (Mercer et al., 2012; Fig. S1, S2). The

motus have similar climate, but differ in vegetation and human activity as well as in larval

competition and average body size of adult mosquitoes. This exceptionally high temporal

and spatial resolution gives unique opportunities for studying population dynamics of

mosquitos. In particular, it allows investigating to what extent the local habitat drives

mosquito dynamics and how neighbouring population in a meta-population are

interconnected. Our objectives are (1) to identify the critical environmental variables that

causally affect mosquito populations using convergent cross mapping (CCM) (Methods), (2)

to test for causal interactions between mosquito subpopulations, and (3) to explore

scenarios how change in a causal environmental variable may affect mosquito abundance as

well as number and strength of mosquito outbreaks.

RESULTS

CAUSALITY BETWEEN ENVIRONMENT AND MOSQUITO ABUNDANCE

Comparing the results based on CCM versus correlation demonstrates the efficacy of using

EDM in our data. Indeed, correlation does not imply causation (Fig. S3), while lack of

correlation does not imply lack of causation (Fig. S4). Moreover, correlation can change sign,

depending on ecosystem context, known as mirage correlation (Figure S5). These examples

from our data highlight the need to employ correct methods for studying causality in

complex nonlinear systems.

When analysing the mosquito dynamics for each of the nine sampling sites separately using

CCM, we found that temperature, precipitation, dew point, air pressure, and mean tide level

are causal variables for the mosquito populations (Fig.1; supplementary Table S2).

Remarkably, the forcing environmental variables varied substantially among the nine

sampling sites (Fig. 1). For example, we found that sampling site S7 at Toamaro island is

affected by four climate variables (temperature, dew point, precipitation, mean tide level),

site S8 by only one (air pressure), and site S9 by none. Similarly strong differences between

sampling sites were found for the other two islands. However, when repeating the same

analysis for each of the three islands (sum of the three sites within the same island) and for

the whole meta-population (i.e., sum of the nine sampling sites), we found no climatic effect

on mosquito abundance from any variable that we analysed here. The spatial scale

dependence is remarkable, given the small sizes of the islands (100-500 m length) and short

distance between sites (a maximum distance between two sampling sites of less than 5 km).

Moreover, the causal forcing variables showed large variation in time lag, from zero day to

more than five weeks even for the same environmental variables. For example, the mean

tide level affects the mosquito dynamics at S6 with a time lag of 0-4 days, but the lag is 27-

30 days at S7; time lags for air pressure are 14-17 days at S1, but 34-39 days at S8. In other

words, the causal effect of climate on the mosquito dynamics has strong temporal scale

dependence.

A striking spatial pattern emerges when the three islands are compared with respect to the

number of causally forcing variables and the time lag of the causal effect. For Horea, we

found only the mean tide level and the air pressure to be causal drivers. The associated time

lags are 7-9 and 14-17 days, respectively (Fig. 1). For Tiano and Toamaro, on the other side,

we detected four and five causally forcing variables, respectively, and the associated time

lags range from 0 to 35 days for Tiano and 9 to 39 days for Toamaro (Fig. 1). These results

suggest that mosquito-environment interactions are more complex on Tiano and Toamaro

than on Horea.

CAUSALITY BETWEEN MOSQUITO POPULATIONS

We tested for causal interactions between all pairs of the nine subpopulations, and found

bidirectional causality in one case (S8<->S9) and unidirectional causality in four cases (S1-

>S3, S2->S3, S2->S7, S7->S8) (Fig. 1, black arrows; supplementary Table S3). A clear spatial

pattern emerges. All causality links start and end at Horea or Toamaro, but none at Tiano.

Further, five of the six links appear within motu (either Horea or Toamaro), with no time lag.

Between-motu interaction among subpopulations was only found in one case, from Horea

(S2) to Toamaro (S7), with a time lag of 14 days. This case is surprising because Horea and

Toamaro are not neighbouring motus, but separated by Tiano (cf. Fig. 1). The lack of

connectivity between the subpopulations of Tiano and the other two motus may be

explained by the fact that Tiano has the lowest mosquito abundances due to intensive pest

control management (Mercer et al. 2012).

SCENARIO EXPLORATION

After identifying the critical forcing environmental variables, we then explored the scenarios

how changes of these causally forcing variables may affect mosquito abundance (Methods).

The key idea is to predict mosquito abundances for scenarios, in which a climate variable is

increased or decreased at each time point, and then compare the predicted abundances

with the original values. Take site S7 as an example, Figure 2 shows the observed

abundances of *Ae. polynesiensis* and the predicted abundances for a scenario with increased

mean tide level. We found that the predicted abundances are higher than the observed for

28 time points but lower for the other 19. Note that increasing tide level does not always

result in reduced or elevated mosquito abundance; rather, the response depends on the

state of population (e.g. the abundance of mosquitos in combination of other environmental

variables at a given sampling interval). We conducted a large-scale screen for all causality

links between climate variables and mosquito abundance. The results suggest that *state*

dependence is a common feature of the *Ae. polynesiensis* population dynamics

(supplementary Fig. S7).

Finally, using the same approach, we also explored how environmental change affects the

number and strength of mosquito outbreaks. Here, mosquito outbreak is defined as

mosquito abundance that is five standard deviations larger than the median abundance of a

given time series. This analysis focuses on mosquito peak abundances but not averages

because the former are of particular importance for pest control and disease prevention. For

example in S6, changes in the mean tide level may have substantial effect on mosquito

outbreaks (Fig. 3). Increased mean tide level is predicted to increase the number of

mosquito outbreaks from 12 up to 17 (Fig. 3A, C) and the outbreak strength by up to a factor

of 1.5 (Fig. 3A, C, D). However, the predicted effect depends on the specific time lag under

consideration. For instance, a moderate reduction in MTL is predicted to reduce the number

of outbreak for d=0 (Fig. 3A), but to increase outbreak for d=1 (Fig. 3B). General predictions

for the mean tide level are difficult to make because our methodology does not allow

evaluating the joined effect of all time lags. At S7, changes in three climate variables were

found to significantly alter number and/or strength of mosquito outbreaks (Fig. 3E and F),

including the dew point with a lag of 22 days, the mean tide level with a lag of 28 days, and

precipitation with a lag of 9 days.

169   DISCUSSION

170 CAUSALITY BETWEEN ENVIRONMENT AND MOSQUITO ABUNDANCE

We applied for the first time the causality test using CCM to a longitudinal mosquito data set

and found that temperature, precipitation, dew point, air pressure, and mean tide level are

forcing climate variables. Our analyses confirm the common belief that temperature and

precipitation causally affect mosquito abundance; in particular, container breeder

mosquitoes, such as *Ae. polynesiensis* and *Ae. aegypti*, critically rely on rainfall and

moderate temperatures for their larval development (Focks et al., 1993). However, previous

rigorous tests for causal effects of environmental factors on mosquitos stem only from lab

experiments (e.g., Couret et al., 2014), whereas studies using longitudinal field data

employed mostly linear statistical methods such as correlation analysis (Scott et al., 2000;

Azil et al., 2010; Alencar et al., 2015) that are ill-posed for the nonlinear mosquito dynamics.

Here, our causality analysis confirms the experimental predictions that, however, hitherto

have not been confirmed at the population level in the natural environment.

Surprisingly, we found that the mean tide level, an environmental factor that was rarely

discussed in the literature, is the most influential environmental variable (quantified by the

number of significant tests) in affecting mosquito populations in French Polynesia (Fig. 1).

We suggest the effect of tide may be related with the special life style of *Ae. polynesiensis*.

Specifically, larvae of *Ae. polynesiensis* can develop in breeding containers filled with

brackish water due to their high tolerance to salinity (Ingram 1954; Chang 2010).

Considering that tide is likely to affect both the water level and the degree of salinity in the

breeding containers, tide potentially has an impact on the number and quality of breeding

sites, which may cause changes of oviposition, larval density, larval survival and

subsequently mosquito population dynamics. Our finding may have important management

implications because the tide level is mainly determined by lunar cycles and therefore bears

a high long-term predictability. This is in stark contrast to the other climate variables, which

have comparatively low long-term predictability. An interesting question for future research

is whether the long-term predictability of the tides may be used for improving forecast of

Ae. polynesiensis outbreaks.

We additionally found that the air pressure is a forcing variable of *Ae. polynesiensis*. While

the effect of air pressure on mosquito survival has been suggested from lab experiments,

although for a different species (*Ae. aegypti*; Galun and Fraenkel, 1961), the empirical

evidence for natural mosquito populations has never been found. A possible explanation is

that mosquitoes use decreasing air pressure as an indicator for adverse weather conditions

and modify their behaviour (e.g. mating) to reduce the risk of injury or death. Such changes

in behaviour were previously reported for *Drosophila melanogaster* (Austin et al., 2014) as

well as for beetles, moths, and aphids (Pellegrino et al., 2013).

Our results on the dew point suggest that evaporation and condensation of water are causal

drivers of the mosquito dynamics. Evaporation reduces water level and increases salinity in

breeding containers, whereas condensation has the opposite effect. A previous study

showed a positive correlation between the dew point and the percentage of containers

holding water versus number of *Ae. aegypti* larvae (Favier et al., 2006).

TEMPORAL SCALE DEPENDENCE OF MOSQUITO-CLIMATE INTERACTIONS

The mosquito-climate interactions exhibited temporal scale dependence. This may be

associated with the mosquito life cycle that consists of 5-10 days larval development

including pupation, and a total life span of up to four weeks. Our causality analysis revealed

short time-lag effects of 0-9 days for temperature, precipitation, dew point, and mean tide

level, which may reflect a causal effect of the climate variable on adult mortality, larval

developmental time or mortality at late larval stages. Intermediate time-lag effects were

found for temperature, precipitation, mean tide level, and air pressure, which may reflect an

effect on larval mortality, larval competition or larval developmental time. Long time-lag

effects were found for temperature, dew point, and mean tide level, which may indicate an

effect of the climate variables on female fecundity or host availability (e.g., human, bird, rat,

dog). Very long time-lag effects of 30-39 days were found for dew point, mean tide level,

and air pressure, which may reflect an effect on the availability of breeding habitats such as

broken coconuts or crab holes. We remark that the causal effect may be direct, e.g.,

temperature may directly affect larval survival due to stress, or indirect, e.g., temperature

may alter behaviour of larval predators. In addition, indirect causality may involve human

activity, e.g., temperature dependent behaviour of pesticide usage. Importantly, the same

climate variable may affect the mosquito at different life stages with different time lags. This

may even occur at the same population; e.g., S4 is affected by temperature with delays of 8

days and 20-23 days.

SPATIAL VARIATION IN MOSQUITO-CLIMATE INTERACTIONS

The causality network displays strong spatial scale dependence with respect to mosquito-

environment interactions. Causal interactions were only detected at the smallest spatial

scale of a single sampling site, but neither at the level of a motu nor at the whole meta-

population. Similar patterns were previously reported for the dynamics of measles, which

showed signatures of nonlinearity and predictability only at the city-by-city scale, but not at

the larger country-wide scale (Sugihara et al. 1992). In both cases, signatures for nonlinearity

are lost by summation of subpopulation dynamics. This may seem counter-intuitive, but is

actually to be expected for a nonlinear dynamical system, in which subpopulations are

affected by different forcing variables or the forcing is non-synchronized (Sugihara and May

1990), as was found in this study (Fig. 1).

The causality analysis revealed substantial heterogeneity in terms of causal environmental

variables and time-lagged effects between neighbouring populations even on the same

motu (Fig. 1). How are such differences at a very small spatial scale determined? A possible

explanation is that populations in close proximity respond differently to the climate even if

the distance is less then 100 m. When interpreting the results, however, it should be kept in

mind that the climate data were not directly collected at the mosquito sampling points

(Methods). Climate data with spatial resolution are necessary to evaluate whether *Ae.*

polynesiensis is really affected by its microhabitat.

An interesting finding was that mosquito-environment interactions are more complex on

Toamaro and Tiano than on Horea (Fig. 1). This difference in complexity may, to some

extent, be explained by differences in mosquito body size. In a previous study on *Ae.*

polynesiensis using the same mosquito samples, Mercer et al. (2012) reported that female

body size is significantly smaller on Toamaro motu than on Horea. The authors explain this

finding by higher larval densities and longer larval developmental times on Toamaro (Mercer

et al. 2012). We hypothesize that either (i) the longer developmental times and the smaller

adult body sizes make mosquitoes on Toamaro more sensitive to environmental

fluctuations, and that this causes the complex mosquito-climate interactions found for this

motu or (ii) the climatic variables influence the larval stage which leads to longer

development time, smaller adult body size, and the complex dynamics. The hypotheses

could be tested using longitudinal data that include both adult and larval stages.

While the three motus were chosen as study sites because they have similar climate, they

actually differ with respect to vegetation and human impacts. In Tiano, pest control is

intense and canopy cover and mosquito abundances are low; in Horea, canopy cover is high,

pest control happens on irregular basis and mosquito abundances are intermediate; in

Toamaro, canopy cover is high, pest control is mostly absent, and mosquito abundance is

high. Among the three motus, Toamaro suffered the least human effects, while the

mosquito population on this motu are characterized by the most complex interactions

between the climate variables and mosquito abundances (Fig. 1). This suggests that pest

control strategies may not only reduce mosquito abundance but also alter the complexity of

mosquito-climate interdependencies. This may be a relevant aspect in pest control.

CAUSALITY BETWEEN MOSQUITO POPULATIONS

An interesting finding was the detection of causality between subpopulations (Fig. 1). There

is a clear spatial pattern with five out of six links connecting neighbouring populations, and

only one link connecting two motus. A possible explanation is that female mosquitoes

disperse by prevailing winds.

Our results demonstrate that CCM allows identifying interactions within a meta-population.

Meta-population dynamics and structure are important determinants of ecological and

evolutionary processes (Hanski and Gilpin, 1997; Hanski, 1998). Meta-population dynamics

and structure were typically studied by directly measuring migration and dispersal, e.g., by

mark-release-recapture experiments (Guerra et al 2014), or indirectly reconstructed by

estimating gene flow between populations, e.g. using microsatellite data (Brelsfoard and

Dobson, 2012). Here for the first time, we demonstrated that CCM may provide an

alternative solution, which may be of wide applicability. For example, one may study how

climate variables, community composition, and population density causally affect the

migration behaviour of a focal species. This analysis provides information of the

interconnectedness in a meta-population. CCM and scenario exploration allow further

investigating whether the meta-population structure itself is context-dependent and how

changes of climate variables alter the interconnectedness in the meta-population. Such

research questions are difficult to study with the existing methods.

SCENARIO EXPLORATIONS

We used the scenario exploration technique (Deyle et al., 2013) to predict how

environmental change affects mosquito abundance. Our analysis revealed strong state

dependent responses (Fig. 2, supplementary Fig. S7). This result is expected given the

various empirical evidences for complex and nonlinear mosquito-environment interactions.

Here, we tailored scenario exploration to the specific needs of disease vector control, and

investigated how environmental change affects mosquito outbreaks. This analysis suggests

that climate change may substantially alter the number and strength of outbreaks, but that

the outcome critically depends on the local environmental conditions and may differ even

between populations in short distance of less than 200 m (Fig. 3).

We propose using scenario exploration as a new framework to predict environmental

change effects on mosquito abundances and disease outbreaks. The main difference to

previous forecasting methods is that realistic, nonlinear mosquito-climate

interdependencies are fully acknowledged. We believe that this offers new opportunities for

mosquito forecasting. As the skill of EDM forecasting depends crucially on the quality of

data, there is a critical need for long-term mosquito monitoring. We suggest to collect time

series data of mosquito abundances on a regular base of 1-2 weeks, and combine them with

climate information from data logger and case reports of transmitted diseases at local

hospitals. Scenario exploration may then be used to (1) predict when outbreaks of

mosquitoes or mosquito-borne diseases happen, and to (2) simulate different pest control

strategies and explore how they depend on climate conditions. In that sense, we hope that

this study sets a first step towards a new direction of pest control and disease prevention

strategies.

In conclusion, we deciphered the causality network of a mosquito meta-population with

climate variables (temperature, precipitation, dew point, air pressure, mean tide level) using

EDM (Fig. 1) that was otherwise difficult to discern using linear methods. The findings that

the responses of subpopulations in close proximity (100-500 m) differed substantially with

respect to their causal environmental drivers and that no causal climate variables were

found for the motu scale and whole meta-population scale highlights the importance of

spatial scale dependency in complex systems. Moreover, we found strong temporal scale

dependence of the causality with time lags ranging from zero up to five weeks, suggesting

that climate acts on the different life stages of the mosquito and that the interdependencies

of mosquito-climate interactions have a high degree of complexity. More importantly, albeit

with the complexity, we presented a new framework based on EDM to predict how

environmental change may affect the number and strength of mosquito outbreaks. These

results bear profound management implications for disease vector ecology and pest control.

METHODS

MOSQUITO FIELD SITE

Adult *Aedes polynesiensis* mosquitoes were collected on three neighboring motu islands,

Horea, Tiano, and Toamaro, on the western coast of Raiatea (Mercer et al., 2012, see also

figure 1). Mosquito data are shown in figure S1.

The linear distance between the shoreline of Raiatea and the motus is approximately 500-

700 m. Horea, the smallest among the three, is a publically held circular island of an area of

1.41 ha, with vegetation consisting of 60-80% canopy cover. This motu is semi permanently

inhabited, and insecticides and wood smoke are used for mosquito control. Mosquito larvae

on this motu develop in artificial containers, coconuts, spathes, phytotemata, and burrows

made by the terrestrial crab *Cardisoma carnifex*. Blood meals of mosquitos are from

humans, dogs, rats, and birds including chicken.

Tiano, the largest among the three, is privately owned and has a size of 6.34 ha. The

vegetation has an open canopy in the center (<20% cover) and 80-90% canopy around the

edge. Mosquitoes are controlled by pesticide usage and breeding site reduction. Mosquito

larvae on this motu develop in cryptic tree holes, crab burrows, and a pool that is

temporarily filled with water. Blood meals are from dogs, rats, birds, and humans.

Toamaro is a privately owned motu with an area of 4.06 ha. The vegetation consists of 70-

95% canopy and has greater stratification with herbaceous and shrub layers than Horea and

Tiano. This island is occasionally managed by vegetation clearing. Insecticides are used for

mosquito control during periods of coconut collection. Mosquito larvae on this motu

develop in artificial containers, broken coconuts, phytotemata, and crab burrows. Blood

meals are from rats, birds, and occasionally dogs, cats, and human.

TIME SERIES DATA

Ae. polynesiensis population data were previously presented by Mercer et al. (2012). Adult

mosquitoes were collected on three neighboring motu islands, Horea, Tiano, and Toamaro,

on the western coast of Raiatea (Society Archipelago, French Polynesia; 16°49.4'S,

159°29.2'W) from 1 February 2008 to 3 December 2009 (Fig. 1). On each motu, there were

three sampling sites, and mosquitoes were collected at a 14-day interval using Biogent

Sentinel traps (data shown in supplementary Fig. S1). Sampling sites are numbered serially

from north to south and denoted by S1 to S9. All collected mosquitoes were species
identified, sex determined, and counted for each time interval. In the present study, we
focused on female *Ae. polynesiensis*, which comprised about 99% of the caught mosquitoes.
For further details see Mercer et al. (2012).

Climate data were downloaded from the *National Oceanic and Atmospheric Administration*
(http://www.noaa.gov/). From Bora Bora airport (16°26.4'S, 151°45.1'W; approx. 50km
distance to the sampling sites at Raiatea), we obtained daily records of mean temperature
(*T*), dew point (*DEW*), precipitation (*PREC*), and air pressure (mean atmospheric sea level
pressure, *SLP*). From Pago Pago (American Samoa; 14°16.6'S, 170°41.3'W), we obtained
hourly tide data, from which we calculated the daily average mean tide levels (*MTL*). Pago
Pago has a distance of approximately 2000 km to Raiatea and was the nearest weather
station providing tide data. Note that Raiatea and Pago Pago have similar latitude. All
climate data were recorded daily for the period from 7 December 2007 to 3 December 2009
(supplementary Fig. S2). This period is eight weeks longer than mosquito sampling and thus
allows investigating possible time-lag effects of climate variables.

DATA ANALYSIS

Convergent cross mapping Causal effects between mosquito populations, and from
environmental variables to mosquito populations were tested using *convergent cross*
mapping (CCM) (Sugihara et al., 2012). CCM tests whether two variables, each given as a
time series, are coupled in the same dynamical system. Variable *X* is said to be causally
affected by *Y* if the underlying dynamic of *X* is a function of *Y*. This method, based on Takens'
theory for dynamical systems (Takens, 1981), tests for causation by measuring the extent to
which the causal variable has left an imprint in the time series of the affected (Sugihara et al
2012). The method is rooted in state space reconstruction of shadow manifold by lagged
coordinate embedding, and the essential ideas are summarized in the following animations:
tinyurl.com/EDM-intro. In this study, the embedding dimension (E_b) for each causal link (e.g.
from *Y*(*t*) to *X*(*t*)) in CCM analysis was determined by testing values of *E* from 2 to 10
dimensions that optimizes the cross-mapping ρ in which *X*(*t*) is used to predict *Y*(*t*).

The causation, *Y* causes *X*, can be confirmed if and only if $SSR_{X(t)}$ cross-mapping $SSR_{Y(t)}$
converges, meaning that the cross-mapping skill $\rho(L)$ improves with increasing library lengths
(*L*) of *X* (Sugihara et al 2012). The direction of cross-mapping (*X* maps *Y*) is opposite to the
direction of causation (*Y* causes *X*) because only *Y* can leave footprints on *X* and makes the
backward cross-mapping from *X* possible. Note that in this study, we also explored the

causations with lag response (van Nes et al., 2015). Specifically, for investigating effects of

climate variables on mosquitos, we tested for time lags ranging from $d = 0$ to 42 days, while

14-day arithmetic means of the climate variables were used such that the climate time

series match the mosquito-sampling interval. To investigate causal effect among mosquito

populations, we conduced CCM between paired populations. Potential lag effects were

evaluated for $d = 0$, 14 and 28 days.

To determine the convergence in cross-mapping, three statistical criteria were applied. First,

Kendall's τ test was used to examine whether the cross-map skill ρ increases monotonously

as a function of library size L. Results are considered as significant if $\tau > 0$ and $P < 0.01$.

Second, Fisher's Z test was used to examine whether the predictability under maximum

library size, $\rho(L_{max})$, is significantly higher than the predictability under minimum library

size, $\rho(L_{min})$. The threshold was set to $P < 0.05$. Third, we tested whether $\rho(L_{max})$ of the

observed time series is significant different from a null model, using the surrogate time

series test. 100 surrogates were generated, following Ebisuzaki (1997), by randomly

reshuffling phases, while preserving the mean, variation and power spectrum. The results

are deemed significant if $\rho(L_{max})$ of the original data is larger than the 95% upper bound of

the surrogates ($P < 0.05$) (Deyle et al., 2013; van Nes et al., 2015). To account for multiple

testing, however, we reduced the level of significance to $P < 0.01$ when testing for causal

climate effects.

Scenario exploration To evaluate how change in a climate variable (the causal variable

identified by CCM) affect mosquito abundances of a given subpopulation, we used

multivariate simplex projection following (Deyle et al., 2013). Specifically, we conducted

scenario explorations with an extended shadow manifold that has E_b+1 dimensions (Deyle et

al., 2013; supplementary Fig. S6). The first E_b dimensions contain the time-delayed

embeddings of the mosquito abundances, as described above. Then, to explore potential

effects of changing environment, an additional coordinate axis of either the time series of a

climate variable or another mosquito population was appended, resulting in E_b+1

dimensions in state space. It is this extra variable that we aim to "manipulate" to achieve

scenario exploration. Specifically, we purposely increase or decrease that forcing variable by

multiples of its standard deviation and then forecast the resultant mosquito abundance one

step (in our case, 14 days) into the future for every time point.

The scenario exploration approach allows us to investigate how change in a climate variable

affects the number and strength of mosquito outbreaks for each mosquito time series. Here,

mosquito outbreaks are defined as mosquito abundances that are five standard deviations

larger than the median abundance. The outbreak strength is defined as the arithmetic mean
of the five highest abundances. For each mosquito time series, we compared the number
and strength of mosquito outbreaks as predicted under a "scenario" versus those in the
original data. To quantify the effect of a scenario of changing a climate variable, we
calculated the relative outbreak strength, which is defined as the arithmetic mean of the five
highest population abundances under scenario exploration divided by that of the original
data.

Further details of data analysis Time series analysis was conducted using the rEDM package
(version 0.2.4) of the programming language *R* (Ye at al., 2016). To minimize autocorrelation
and heteroscedasticity, all time series were first differenced and normalized to zero mean
and unit variance (Sugihara, 1994; Telschow et al., 2017). Due to the paucity of observation
points, we performed leave-one-out cross-validation (Glaser et al., 2014).

AUTHORS CONTRIBUTIONS

Design of study: FG, CH, AT; data analysis: FG; interpretation of results: FG, JS, CH, AT;
writing of manuscript: FG, JS, CH, AT. All authors gave final approval for publication.

ADDITIONAL INFORMATION
The authors declare no competing financial interests.

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 FIGURES

Figure 1. Causality network of the Polynesian tiger mosquito (*Ae. polynesiensis*). *Upper*

graph: Field site in Raiatea (French Polynesia). *Lower graph:* Causal interactions within the

mosquito meta-population (black arrows) and between climate variables and mosquito

abundances (coloured arrows). Causality was evaluated using convergent cross mapping.

Only significant results are shown (see Methods, supplementary Tables S2, S3 for details).

Numbers next to arrows indicate the time lag d (in days), for which causality was detected.

Symbols S1 to S9 denote the different sampling sites.

Figure 2. Predicted state-dependent effect of environmental change on mosquito

dynamics using scenario exploration. Shown are the observed mosquito abundances at

sampling site S7 (black), and the predicted abundances for the same sampling site under an

increase of the mean tide level by half of its standard deviation (red). Predictions were

generated using multivariate simplex projections and a time lag of d=28 days. Note that

there is a causal effect of the MTL on the S7 population for this time lag (see Fig. 1). The

analysis predicts that an increase in the MTL results in increased or decreased mosquito

abundance depending on the state of the system.

Figure 3. Effect of climate change on number and strength of mosquito outbreaks. Two types of scenario explorations were conducted. The blue line shows how climate change affects the number of mosquito outbreaks, defined as mosquito abundances that are five standard deviations larger than the median of the original data. The red line indicates the relative strength of mosquito outbreaks, defined as the arithmetic mean of the highest five population abundances under climate change divided by that of the original data. The light red zone indicates the range between the highest and the fifth highest mosquito abundance of the original time series. The figure shows only the cases, in which the outbreak strength is at least once outside of the light red zone. These are considered as scenarios with a substantial effect of climate change on mosquito abundance. However, the analysis was conducted for all causality links between climate and mosquito abundance (see supplementary Table S2).