# Dynamic modeling of abundance data in ecology 

Guillaume Franchi

ENSAI, Bruz

ENSAI $=こ=$ de lastatistique
de la statistique
de l'information


## ECODEP Conference <br> 12 February 2024

## Outlines

## I. Introduction

II. Modeling relative abundance
III. Modeling Absence/Presence of species

## Outlines

## I. Introduction

## II. Modeling relative abundance

## III. Modeling Absence/Presence of species

## Ecological definitions

Consider an ecosystem with a number $d$ of species.

## Ecological definitions

Consider an ecosystem with a number $d$ of species.

## Definition

Relative abundance: the vector of proportions of each species in the whole ecosystem.

## Ecological definitions

Consider an ecosystem with a number $d$ of species.

## Definition

Relative abundance: the vector of proportions of each species in the whole ecosystem.

It is an element of the simplex

$$
\mathcal{S}_{d-1}=\left\{\left(y_{1}, \ldots, y_{d}\right) \in\right] 0,+\infty\left[{ }^{d} / \sum_{i=1}^{d} y_{i}=1\right\}
$$

## Ecological definitions

Consider an ecosystem with a number $d$ of species.

## Definition

Relative abundance: the vector of proportions of each species in the whole ecosystem.

It is an element of the simplex

$$
\mathcal{S}_{d-1}=\left\{\left(y_{1}, \ldots, y_{d}\right) \in\right] 0,+\infty\left[{ }^{d} / \sum_{i=1}^{d} y_{i}=1\right\}
$$

One can also consider the Shannon entropy

$$
\forall y=\left(y_{1}, \ldots, y_{d}\right) \in \mathcal{S}_{d-1}, I_{S}(y)=-\sum_{i=1}^{d} y_{i} \log \left(y_{i}\right)
$$

## Objectives

(0) Make predictions about the relative abundance of an ecosystem over time.

## Objectives

(0) Make predictions about the relative abundance of an ecosystem over time.
(0) Understand the dynamics of this ecosystem:

## Objectives

(0) Make predictions about the relative abundance of an ecosystem over time.
(0) Understand the dynamics of this ecosystem:
$\rightarrow$ The dynamic of each species.

## Objectives

(0) Make predictions about the relative abundance of an ecosystem over time.
(0) Understand the dynamics of this ecosystem:
$\rightarrow$ The dynamic of each species.
$\rightarrow$ The interaction between species.

## Objectives

(0) Make predictions about the relative abundance of an ecosystem over time.
(0) Understand the dynamics of this ecosystem:
$\rightarrow$ The dynamic of each species.
$\rightarrow$ The interaction between species.
$\rightarrow$ The impact of exogenous variables.

## An example

We consider a population of insects studied in a sugar cane field in La Réunion, during the years 2022 and 2023.

## An example

We consider a population of insects studied in a sugar cane field in La Réunion, during the years 2022 and 2023.

We focus on three groups of species:

(a) Coleoptera

(b) Hymenoptera

(c) Diptera

## An example

We consider a population of insects studied in a sugar cane field in La Réunion, during the years 2022 and 2023.


## Outlines

## I. Introduction

II. Modeling relative abundance

## III. Modeling Absence/Presence of species

## Chain with complete connections (1/2)

We propose to model our abundance along time by a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ valued in the simplex $\mathcal{S}_{d-1}$.

## Chain with complete connections (1/2)

We propose to model our abundance along time by a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ valued in the simplex $\mathcal{S}_{d-1}$.

The idea is to define $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ as a chain with complete connections

$$
\mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

## Chain with complete connections (1/2)

We propose to model our abundance along time by a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ valued in the simplex $\mathcal{S}_{d-1}$.

The idea is to define $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ as a chain with complete connections

$$
\mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

where:

- $P$ is a transition kernel from source $\mathcal{S}_{d-1}^{\mathbb{N}}$ and target $\mathcal{S}_{d-1}$,


## Chain with complete connections (1/2)

We propose to model our abundance along time by a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ valued in the simplex $\mathcal{S}_{d-1}$.

The idea is to define $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ as a chain with complete connections

$$
\mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

where:

- $P$ is a transition kernel from source $\mathcal{S}_{d-1}^{\mathbb{N}}$ and target $\mathcal{S}_{d-1}$,
- $Y_{t}^{-}$denotes the entire past of the time series at time $t$ :

$$
Y_{t}^{-}=\left(Y_{t}, Y_{t-1}, \ldots\right)
$$

## Chain with complete connections (2/2)

## Remark

## Chain with complete connections (2/2)

## Remark

- The process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ has possibly an infinite memory.


## Chain with complete connections (2/2)

## Remark

- The process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ has possibly an infinite memory.
- If $P\left(A \mid Y_{t}^{-}\right)$depends only on the $p+1$ first values of $Y_{t}^{-}$

$$
P\left(A \mid Y_{t}^{-}\right)=\tilde{P}\left(A \mid Y_{t}, Y_{t-1}, \ldots, Y_{t-p}\right)
$$

we obtain a Markov chain.

## Chain with complete connections (2/2)

## Remark

- The process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ has possibly an infinite memory.
- If $P\left(A \mid Y_{t}^{-}\right)$depends only on the $p+1$ first values of $Y_{t}^{-}$

$$
P\left(A \mid Y_{t}^{-}\right)=\tilde{P}\left(A \mid Y_{t}, Y_{t-1}, \ldots, Y_{t-p}\right)
$$

we obtain a Markov chain.

- It is possible to add a process of exogenous variables $\left(X_{t}\right)_{t \in \mathbb{Z}}$ to the dynamic of the process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$

$$
\mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}, X_{t}^{-}=x_{t}^{-}\right)=P\left(A \mid y_{t}^{-}, x_{t}^{-}\right)
$$

## Existence of the process

## Existence of the process

## Theorem 1

Under assumptions A1 and A2 below, there exists a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ which is strictly stationary such that

$$
\forall t \in \mathbb{Z}, \mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

## Existence of the process

## Theorem 1

Under assumptions $\mathbf{A 1}$ and $\mathbf{A} 2$ below, there exists a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ which is strictly stationary such that

$$
\forall t \in \mathbb{Z}, \mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

Furthermore, its distribution is unique and it is ergodic.

## Existence of the process

## Theorem 1

Under assumptions $\mathbf{A 1}$ and $\mathbf{A} 2$ below, there exists a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ which is strictly stationary such that

$$
\forall t \in \mathbb{Z}, \mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

Furthermore, its distribution is unique and it is ergodic.

## Assumptions

A1 $b_{0}=\sup \left\{d_{T V}(P(\cdot \mid y), P(\cdot \mid z)) / y, z \in \mathcal{S}_{d-1}^{\mathbb{N}}\right\}<1$.

## Existence of the process

## Theorem 1

Under assumptions $\mathbf{A 1}$ and $\mathbf{A} 2$ below, there exists a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ which is strictly stationary such that

$$
\forall t \in \mathbb{Z}, \mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

Furthermore, its distribution is unique and it is ergodic.

## Assumptions

A1 $b_{0}=\sup \left\{d_{T V}(P(\cdot \mid y), P(\cdot \mid z)) / y, z \in \mathcal{S}_{d-1}^{\mathbb{N}}\right\}<1$.
A2 For $m \geqslant 1$, we denote

$$
b_{m}=\sup \left\{d_{T V}(P(\cdot \mid y), P(\cdot \mid z)) / y, z \in \mathcal{S}_{d-1}^{\mathbb{N}}, y \stackrel{m}{=} z\right\}
$$

## Existence of the process

## Theorem 1

Under assumptions $\mathbf{A 1}$ and $\mathbf{A} 2$ below, there exists a time series $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ which is strictly stationary such that

$$
\forall t \in \mathbb{Z}, \mathbb{P}\left(Y_{t+1} \in A \mid Y_{t}^{-}=y_{t}^{-}\right)=P\left(A \mid y_{t}^{-}\right)
$$

Furthermore, its distribution is unique and it is ergodic.

## Assumptions

A1 $b_{0}=\sup \left\{d_{T V}(P(\cdot \mid y), P(\cdot \mid z)) / y, z \in \mathcal{S}_{d-1}^{\mathbb{N}}\right\}<1$.
A2 For $m \geqslant 1$, we denote

$$
b_{m}=\sup \left\{d_{T V}(P(\cdot \mid y), P(\cdot \mid z)) / y, z \in \mathcal{S}_{d-1}^{\mathbb{N}}, y \stackrel{m}{=} z\right\}
$$

We have $\sum_{m \in \mathbb{N}} b_{m}<\infty$.

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

8 In the context of Ecology, it has been suggested in Marquet et al. (2017) that the relative abundance of a given species in large ecosystems is often compatible with a Beta distribution.

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

8 In the context of Ecology, it has been suggested in Marquet et al. (2017) that the relative abundance of a given species in large ecosystems is often compatible with a Beta distribution.

## Remark

The Dirichlet distribution is actually the generalization of the Beta distribution.

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

8 In the context of Ecology, it has been suggested in Marquet et al. (2017) that the relative abundance of a given species in large ecosystems is often compatible with a Beta distribution.

## Remark

The Dirichlet distribution is actually the generalization of the Beta distribution.

The Dirichlet distribution $\operatorname{Dir}(\lambda, \varphi)$, supported by $\mathcal{S}_{d-1}$ is characterized by

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

8 In the context of Ecology, it has been suggested in Marquet et al. (2017) that the relative abundance of a given species in large ecosystems is often compatible with a Beta distribution.

## Remark

The Dirichlet distribution is actually the generalization of the Beta distribution.

The Dirichlet distribution $\operatorname{Dir}(\lambda, \varphi)$, supported by $\mathcal{S}_{d-1}$ is characterized by
$\rightarrow$ its mean vector $\lambda=\left(\lambda_{1}, \ldots, \lambda_{d}\right)$;

## Dirichlet model ( $1 / 3$ )

A natural proposal for $P\left(\cdot \mid Y_{t}^{-}\right)$is a Dirichlet distribution

$$
P\left(\cdot \mid Y_{t}^{-}\right)=\operatorname{Dir}\left(\lambda_{t}, \varphi_{t}\right)
$$

8 In the context of Ecology, it has been suggested in Marquet et al. (2017) that the relative abundance of a given species in large ecosystems is often compatible with a Beta distribution.

## Remark

The Dirichlet distribution is actually the generalization of the Beta distribution.

The Dirichlet distribution $\operatorname{Dir}(\lambda, \varphi)$, supported by $\mathcal{S}_{d-1}$ is characterized by
$\rightarrow$ its mean vector $\lambda=\left(\lambda_{1}, \ldots, \lambda_{d}\right)$;
$\rightarrow$ a dispersion parameter $\varphi>0$.

## Dirichlet model (2/3)

$\rightarrow$ In the spirit of the logistic regression, we propose that for all $t \in \mathbb{Z}$

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\sum_{k \geqslant 1} \eta_{k} \bar{Y}_{t-k}+\sum_{k \geqslant 1} \zeta_{k} X_{t-k}
$$

where alr is the mapping

$$
\begin{aligned}
\text { alr }: & \mathcal{S}_{d-1} \longrightarrow \mathbb{R}^{d-1} \\
& y=\left(y_{1}, \ldots, y_{d}\right) \longmapsto\left(\log \left(\frac{y_{1}}{y_{d}}\right), \ldots, \log \left(\frac{y_{d-1}}{y_{d}}\right)\right),
\end{aligned}
$$

$\bar{Y}=\left(Y_{1}, \ldots, Y_{d-1}\right)$, and the $\eta$ 's and $\zeta$ 's are matrices.

## Dirichlet model ( $2 / 3$ )

$\rightarrow$ In the spirit of the logistic regression, we propose that for all $t \in \mathbb{Z}$

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\sum_{k \geqslant 1} \eta_{k} \bar{Y}_{t-k}+\sum_{k \geqslant 1} \zeta_{k} X_{t-k}
$$

where alr is the mapping

$$
\begin{aligned}
\text { alr }: & \mathcal{S}_{d-1} \longrightarrow \mathbb{R}^{d-1} \\
& y=\left(y_{1}, \ldots, y_{d}\right) \longmapsto\left(\log \left(\frac{y_{1}}{y_{d}}\right), \ldots, \log \left(\frac{y_{d-1}}{y_{d}}\right)\right),
\end{aligned}
$$

$\bar{Y}=\left(Y_{1}, \ldots, Y_{d-1}\right)$, and the $\eta$ 's and $\zeta$ 's are matrices.
$\rightarrow$ We also propose that for all $t \in \mathbb{Z}$

$$
\varphi_{t}=\exp \left(\theta_{0}+\sum_{k \geqslant 1} \theta_{k} I_{S}\left(Y_{t-k+1}\right)\right)
$$

where the $\theta_{k}$ 's are real numbers.

## Dirichlet model (3/3)

8 The matrices $\eta$ 's give us precise information about the interactions between species.

## Dirichlet model (3/3)

\& The matrices $\eta$ 's give us precise information about the interactions between species.

8 The matrices $\zeta$ 's give us precise information about the impact of exogenous variables on the abundance of species.

## Dirichlet model (3/3)

8 The matrices $\eta$ 's give us precise information about the interactions between species.

8 The matrices $\zeta$ 's give us precise information about the impact of exogenous variables on the abundance of species.

8 The $\theta$ 's give us information about the volatility of the abundance. The idea is to connect the biodiversity of the ecosystem and its variability.

## Return to our example ( $1 / 2$ )

We fit the following Dirichlet model to the population of insects in La Réunion

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\eta_{1} Y_{t}+\zeta_{1} X_{t}
$$

## Return to our example ( $1 / 2$ )

We fit the following Dirichlet model to the population of insects in La Réunion

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\eta_{1} Y_{t}+\zeta_{1} X_{t}
$$

and

$$
\varphi_{t}=\exp \left(\theta_{0}+\theta_{1} I_{S}\left(Y_{t}\right)\right)
$$

## Return to our example (1/2)

We fit the following Dirichlet model to the population of insects in La Réunion

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\eta_{1} Y_{t}+\zeta_{1} X_{t}
$$

and

$$
\varphi_{t}=\exp \left(\theta_{0}+\theta_{1} I_{S}\left(Y_{t}\right)\right)
$$

The vector of covariates $X_{t}$ is composed by climatic variables such as the total rainfall amount, the temperature, the ground radiation and evapotranspiration.

## Return to our example (1/2)

We fit the following Dirichlet model to the population of insects in La Réunion

$$
\operatorname{alr}\left(\lambda_{t}\right)=\eta_{0}+\eta_{1} Y_{t}+\zeta_{1} X_{t}
$$

and

$$
\varphi_{t}=\exp \left(\theta_{0}+\theta_{1} I_{S}\left(Y_{t}\right)\right)
$$

The vector of covariates $X_{t}$ is composed by climatic variables such as the total rainfall amount, the temperature, the ground radiation and evapotranspiration.

An optimization of the conditional likelihood is performed to obtain an estimation of the parameters.

## Return to our example $(2 / 2)$

We kept the 12 last weeks of our data apart from our estimation sample, they are indeed used to compare our predictions with the values observed in the reality.

## Return to our example $(2 / 2)$

We kept the 12 last weeks of our data apart from our estimation sample, they are indeed used to compare our predictions with the values observed in the reality.

The predictions are made by simulating a thousand trajectories of the abundance for the last 12 weeks, and taking the mean of these trajectories gives us our predicted values.

## Return to our example ( $2 / 2$ )

We kept the 12 last weeks of our data apart from our estimation sample, they are indeed used to compare our predictions with the values observed in the reality.

The predictions are made by simulating a thousand trajectories of the abundance for the last 12 weeks, and taking the mean of these trajectories gives us our predicted values.


## Return to our example ( $2 / 2$ )

We kept the 12 last weeks of our data apart from our estimation sample, they are indeed used to compare our predictions with the values observed in the reality.

The predictions are made by simulating a thousand trajectories of the abundance for the last 12 weeks, and taking the mean of these trajectories gives us our predicted values.


## Limits of the model

## Limits of the model

X In practice, ecological data do not have a lot of observations along time.

## Limits of the model

$X$ In practice, ecological data do not have a lot of observations along time.

X In practice, there are a lot of zero values in the abundances observed, which will lead to an error when computing our estimators.

## Limits of the model

X In practice, ecological data do not have a lot of observations along time.
$x$ In practice, there are a lot of zero values in the abundances observed, which will lead to an error when computing our estimators.

8 Use panel data.

## Limits of the model

X In practice, ecological data do not have a lot of observations along time.

X In practice, there are a lot of zero values in the abundances observed, which will lead to an error when computing our estimators.

8 Use panel data.
\& Model the absence/presence of species in the ecosystem.

## Outlines

## I. Introduction

## II. Modeling relative abundance

## III. Modeling Absence/Presence of species

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

We assume that for all $i \in\{1, \ldots, d\}$

$$
Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

We assume that for all $i \in\{1, \ldots, d\}$

$$
Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

where

$$
\rightarrow \lambda_{t}=\sum_{l=1}^{p} A_{l} \cdot Y_{t-l}+B \cdot X_{t-1}
$$

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

We assume that for all $i \in\{1, \ldots, d\}$

$$
Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

where
$\rightarrow \lambda_{t}=\sum_{l=1}^{p} A_{l} \cdot Y_{t-l}+B \cdot X_{t-1}$;
$\rightarrow\left(X_{t}\right)_{t \in \mathbb{Z}}$ is a process of covariates;

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

We assume that for all $i \in\{1, \ldots, d\}$

$$
Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

where
$\rightarrow \lambda_{t}=\sum_{l=1}^{p} A_{l} \cdot Y_{t-l}+B \cdot X_{t-1}$;
$\rightarrow\left(X_{t}\right)_{t \in \mathbb{Z}}$ is a process of covariates;
$\rightarrow\left(\varepsilon_{t}\right)_{t \in \mathbb{Z}}$ is a sequence of i.i.d random variables with distribution $\mathcal{N}_{\mathbb{R}^{d}}(0, R)$.

## Dynamic probit regression

We model here the absence/presence of $d$ species in an ecosystem at time $t$ by

$$
Y_{t}=\left(Y_{1, t}, \ldots, Y_{d, t}\right) \in\{0,1\}^{d}
$$

We assume that for all $i \in\{1, \ldots, d\}$

$$
Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

where
$\rightarrow \lambda_{t}=\sum_{l=1}^{p} A_{l} \cdot Y_{t-l}+B \cdot X_{t-1}$;
$\rightarrow\left(X_{t}\right)_{t \in \mathbb{Z}}$ is a process of covariates;
$\rightarrow\left(\varepsilon_{t}\right)_{t \in \mathbb{Z}}$ is a sequence of i.i.d random variables with distribution $\mathcal{N}_{\mathbb{R}^{d}}(0, R)$.

8 It is actually a dynamic version of a multivariate probit regression.

## Existence of the process

## Existence of the process

## Theorem 2

Assume that the process $\left(\zeta_{t}\right)_{t \in \mathbb{Z}}$ defined by

$$
\zeta_{t}=\left(X_{t-1}, \varepsilon_{t}\right)
$$

is strongly stationary.

## Existence of the process

## Theorem 2

Assume that the process $\left(\zeta_{t}\right)_{t \in \mathbb{Z}}$ defined by

$$
\zeta_{t}=\left(X_{t-1}, \varepsilon_{t}\right)
$$

is strongly stationary.
There exists a strongly stationary process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ satisfying

$$
\forall i \in\{1, \ldots, d\}, Y_{i, t}=\mathbf{1}_{] 0,+\infty[ }\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

## Existence of the process

## Theorem 2

Assume that the process $\left(\zeta_{t}\right)_{t \in \mathbb{Z}}$ defined by

$$
\zeta_{t}=\left(X_{t-1}, \varepsilon_{t}\right)
$$

is strongly stationary.
There exists a strongly stationary process $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ satisfying

$$
\forall i \in\{1, \ldots, d\}, Y_{i, t}=\mathbf{1}_{] 0,+\infty}\left(\lambda_{i, t}+\varepsilon_{i, t}\right)
$$

In addition, its distribution is unique.

## Remark

Furthermore, if $\left(\zeta_{t}\right)_{t \in \mathbb{Z}}$ is ergodic, $\left(Y_{t}\right)_{t \in \mathbb{Z}}$ is also ergodic.

## Estimation results (1/3)

We consider first a single trajectory of an absence/presence process $\left(Y_{t}\right)_{1 \leqslant t \leqslant T}$, and we are interested in the estimation of

$$
\theta=\left(A_{1}, \ldots, A_{p}, B, R\right)
$$

## Estimation results ( $1 / 3$ )

We consider first a single trajectory of an absence/presence process $\left(Y_{t}\right)_{1 \leqslant t \leqslant T}$, and we are interested in the estimation of

$$
\theta=\left(A_{1}, \ldots, A_{p}, B, R\right)
$$

$\rightarrow$ Optimizing the pseudo conditional log-likelihood

$$
\hat{\theta}=\operatorname{argmax} \sum_{t=p+1}^{T} \log \left(\int_{\mathbb{R}^{k}} \prod_{i=1}^{k} \mathbb{1}_{I_{Y_{i, t}}}\left(\lambda_{i, t}+x_{i}\right) \varphi_{R}(x) \mathrm{d} x\right)
$$

where $\varphi_{R}$ is the density of the distribution $\mathcal{N}(0, R)$ and

$$
I_{Y_{i, t}}=\left\{\begin{array}{l}
] 0,+\infty\left[\quad \text { if } Y_{i, t}=1\right. \\
]-\infty, 0] \quad \text { if } Y_{i, t}=0
\end{array} .\right.
$$

## Estimation results ( $1 / 3$ )

We consider first a single trajectory of an absence/presence process $\left(Y_{t}\right)_{1 \leqslant t \leqslant T}$, and we are interested in the estimation of

$$
\theta=\left(A_{1}, \ldots, A_{p}, B, R\right)
$$

$\rightarrow$ Optimizing the pseudo conditional log-likelihood

$$
\hat{\theta}=\operatorname{argmax} \sum_{t=p+1}^{T} \log \left(\int_{\mathbb{R}^{k}} \prod_{i=1}^{k} \mathbb{1}_{I_{Y_{i, t}}}\left(\lambda_{i, t}+x_{i}\right) \varphi_{R}(x) \mathrm{d} x\right)
$$

where $\varphi_{R}$ is the density of the distribution $\mathcal{N}(0, R)$ and

$$
I_{Y_{i, t}}=\left\{\begin{array}{l}
] 0,+\infty\left[\quad \text { if } Y_{i, t}=1\right. \\
]-\infty, 0] \quad \text { if } Y_{i, t}=0
\end{array} .\right.
$$

$\times$ Difficult function to optimize...

## Estimation results (2/3)

We thus propose a two-step method.

## Estimation results (2/3)

We thus propose a two-step method.
$\rightarrow$ We first optimize with respect to $\gamma=\left(A_{1}, \ldots, A_{p}, B\right)$

$$
\hat{\gamma}=\operatorname{argmax} \sum_{t=p+1}^{T} \sum_{i=1}^{k} Y_{i, t} \log \left(\Phi\left(\lambda_{i, t}\right)\right)+\left(1-Y_{i, t}\right) \log \left(\Phi\left(-\lambda_{i, t}\right)\right.
$$

where $\Phi$ denotes the cdf of the gaussian distribution.

## Estimation results (2/3)

We thus propose a two-step method.
$\rightarrow$ We first optimize with respect to $\gamma=\left(A_{1}, \ldots, A_{p}, B\right)$

$$
\hat{\gamma}=\operatorname{argmax} \sum_{t=p+1}^{T} \sum_{i=1}^{k} Y_{i, t} \log \left(\Phi\left(\lambda_{i, t}\right)\right)+\left(1-Y_{i, t}\right) \log \left(\Phi\left(-\lambda_{i, t}\right)\right.
$$

where $\Phi$ denotes the cdf of the gaussian distribution.
$\rightarrow$ We then maximize all pairwise conditional likelihoods

$$
\widehat{R}(i, j)=\underset{r \in]-1,1[ }{\operatorname{argmax}} \sum_{t=p+1}^{T} \log \left\{\int_{I_{Y_{i, t}}-\hat{\lambda}_{i, t}} \Phi\left(\left(2 Y_{j, t}-1\right) \frac{\hat{\lambda}_{j, t}+r x_{i}}{\sqrt{1-r^{2}}}\right) \varphi\left(x_{i}\right) \mathrm{d} x_{i}\right\}
$$

## Estimation results (3/3)

## Proposition 1

Assume the process $\zeta_{t}$ is ergodic. Under some reasonable assumptions on the covariates:

1) All estimators $\hat{\theta}, \hat{\gamma}$ and $\hat{R}$ are strongly consistent.
2) Moreover, we have the asymptotic normality of

$$
\sqrt{T-p}\left(\hat{\theta}-\theta_{0}\right) \quad \text { and } \quad \sqrt{T-p}\left(\hat{\gamma}-\gamma_{0}, \widehat{R}-R_{0}\right)
$$

## The case of panel data

We now consider a number of $n$ trajectories of an absence/presence process $\left(Y_{j, t}\right)_{1 \leqslant j \leqslant n, 1 \leqslant t \leqslant T}$, and are still interested in the estimation of $\theta$.

## The case of panel data

We now consider a number of $n$ trajectories of an absence/presence process $\left(Y_{j, t}\right)_{1 \leqslant j \leqslant n, 1 \leqslant t \leqslant T}$, and are still interested in the estimation of $\theta$.

8 The aim is to improve the speed of convergence of our estimators with the number of sites.

## The case of panel data

We now consider a number of $n$ trajectories of an absence/presence process $\left(Y_{j, t}\right)_{1 \leqslant j \leqslant n, 1 \leqslant t \leqslant T}$, and are still interested in the estimation of $\theta$.

8 The aim is to improve the speed of convergence of our estimators with the number of sites.

敖 Obtain a general version of Birkhoff's ergodic theorem (Giap \& Van Quang, 2016).

## The case of panel data

We now consider a number of $n$ trajectories of an absence/presence process $\left(Y_{j, t}\right)_{1 \leqslant j \leqslant n, 1 \leqslant t \leqslant T}$, and are still interested in the estimation of $\theta$.

8 The aim is to improve the speed of convergence of our estimators with the number of sites.

敖 Obtain a general version of Birkhoff's ergodic theorem (Giap \& Van Quang, 2016).

Generalize the results about consistency and central limit theorems for $M$-estimators.

## Results about $M$-estimators (1/2)

## Results about $M$-estimators (1/2)

Usually, if we consider the estimator

$$
\hat{\theta}=\operatorname{argmax} \sum_{t=1}^{T} m_{\theta}\left(Z_{t}\right)
$$

where $m_{\theta}$ is a measurable mapping and $\left(Z_{t}\right)_{t \in \mathbb{Z}}$ an ergodic process,

## Results about $M$-estimators (1/2)

Usually, if we consider the estimator

$$
\hat{\theta}=\operatorname{argmax} \sum_{t=1}^{T} m_{\theta}\left(Z_{t}\right)
$$

where $m_{\theta}$ is a measurable mapping and $\left(Z_{t}\right)_{t \in \mathbb{Z}}$ an ergodic process, the consistency of $\hat{\theta}$ relies in particular on

$$
\mathbb{E}\left(\sup _{\theta}\left|m_{\theta}\left(Z_{0}\right)\right|\right)<+\infty
$$

## Results about $M$-estimators (1/2)

Usually, if we consider the estimator

$$
\hat{\theta}=\operatorname{argmax} \sum_{t=1}^{T} m_{\theta}\left(Z_{t}\right)
$$

where $m_{\theta}$ is a measurable mapping and $\left(Z_{t}\right)_{t \in \mathbb{Z}}$ an ergodic process, the consistency of $\hat{\theta}$ relies in particular on

$$
\mathbb{E}\left(\sup _{\theta}\left|m_{\theta}\left(Z_{0}\right)\right|\right)<+\infty
$$

and its asymptotic normality on

$$
\mathbb{E}\left(\left\|\dot{m}_{\theta_{0}}\left(Z_{0}\right)\right\|^{2}\right)<+\infty
$$

## Results about $M$-estimators (2/2)

In the case of panel data, the same results can be obtained with

$$
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=1}^{T} m_{\theta}\left(Z_{j, t}\right)
$$

## Results about $M$-estimators (2/2)

In the case of panel data, the same results can be obtained with

$$
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=1}^{T} m_{\theta}\left(Z_{j, t}\right)
$$

$\rightarrow$ by assuming that all processes $\left(Z_{1, t}\right)_{t \in \mathbb{Z}}, \ldots,\left(Z_{n, t}\right)_{t \in \mathbb{Z}}$ are mutually independent, and their distributions are identical;

## Results about $M$-estimators (2/2)

In the case of panel data, the same results can be obtained with

$$
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=1}^{T} m_{\theta}\left(Z_{j, t}\right),
$$

$\rightarrow$ by assuming that all processes $\left(Z_{1, t}\right)_{t \in \mathbb{Z}}, \ldots,\left(Z_{n, t}\right)_{t \in \mathbb{Z}}$ are mutually independent, and their distributions are identical;
$\rightarrow$ by modifying the "order conditions"

$$
\mathbb{E}\left(\sup _{\theta}\left|m_{\theta}\left(Z_{0,0}\right)\right|^{1+\delta}\right)<+\infty \quad \text { and } \quad \mathbb{E}\left(\left\|\dot{m}_{\theta_{0}}\left(Z_{0,0}\right)\right\|^{2(1+\delta)}\right)<+\infty ;
$$

## Results about $M$-estimators (2/2)

In the case of panel data, the same results can be obtained with

$$
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=1}^{T} m_{\theta}\left(Z_{j, t}\right),
$$

$\rightarrow$ by assuming that all processes $\left(Z_{1, t}\right)_{t \in \mathbb{Z}}, \ldots,\left(Z_{n, t}\right)_{t \in \mathbb{Z}}$ are mutually independent, and their distributions are identical;
$\rightarrow$ by modifying the "order conditions"

$$
\mathbb{E}\left(\sup _{\theta}\left|m_{\theta}\left(Z_{0,0}\right)\right|^{1+\delta}\right)<+\infty \quad \text { and } \quad \mathbb{E}\left(\left\|\dot{m}_{\theta_{0}}\left(Z_{0,0}\right)\right\|^{2(1+\delta)}\right)<+\infty ;
$$

$\rightarrow$ and by adding the following "order condition"

$$
\mathbb{E}\left(\left\|\ddot{m}_{\theta_{0}}\left(Z_{0,0}\right)\right\|^{1+\delta}\right)<+\infty
$$

for some $\delta>0$.

## Estimation Results for panel data (1/2)

In the case of panel data, we can consider similar estimators as the ones mentioned previously

$$
\begin{gathered}
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=p+1}^{T} \log \left(\int_{\mathbb{R}^{k}} \prod_{i=1}^{k} \mathbb{1}_{I_{Y_{i, j, t}}}\left(\lambda_{i, j, t}+x_{i}\right) \varphi_{R}(x) \mathrm{d} x\right), \\
\hat{\gamma}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=p+1}^{T} \sum_{i=1}^{k} Y_{i, j, t} \log \left(\Phi\left(\lambda_{i, j, t}\right)\right)+\left(1-Y_{i, j, t}\right) \log \left(\Phi\left(-\lambda_{i, j, t}\right)\right)
\end{gathered}
$$

## Estimation Results for panel data (1/2)

In the case of panel data, we can consider similar estimators as the ones mentioned previously

$$
\begin{gathered}
\hat{\theta}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=p+1}^{T} \log \left(\int_{\mathbb{R}^{k}} \prod_{i=1}^{k} \mathbb{1}_{I_{Y_{i, j, t}}}\left(\lambda_{i, j, t}+x_{i}\right) \varphi_{R}(x) \mathrm{d} x\right), \\
\hat{\gamma}=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=p+1}^{T} \sum_{i=1}^{k} Y_{i, j, t} \log \left(\Phi\left(\lambda_{i, j, t}\right)\right)+\left(1-Y_{i, j, t}\right) \log \left(\Phi\left(-\lambda_{i, j, t}\right)\right)
\end{gathered}
$$

and
$\widehat{R}\left(i_{1}, i_{2}\right)=\operatorname{argmax} \sum_{j=1}^{n} \sum_{t=p+1}^{T} \log \int_{I_{Y_{i_{1}}, j, t}-\hat{\lambda}_{i_{1}, j, t}} \Phi\left(\left(2 Y_{i_{2}, j, t}-1\right) \frac{\hat{\lambda}_{i_{2}, j, t}+r x_{i_{1}}}{\sqrt{1-r^{2}}}\right) \varphi\left(x_{i_{1}}\right) \mathrm{d} x_{i_{1}}$.

## Estimation Results for panel data (2/2)

## Proposition 2

Under some reasonable assumptions on the processes $\left(\zeta_{j, t}\right)_{t \in \mathbb{Z}}$ 's:

1) All estimators $\hat{\theta}, \hat{\gamma}$ and $\widehat{R}$ are strongly consistent.
2) Moreover, we have the asymptotic normality of

$$
\sqrt{n(T-p)}\left(\hat{\theta}-\theta_{0}\right) \quad \text { and } \quad \sqrt{n(T-p)}\left(\hat{\gamma}-\gamma_{0}, \widehat{R}-R_{0}\right)
$$

## Simulations ( $1 / 3$ )

We simulated the absence/presence of 3 fish species, depending on the temperature and salinity of the water, over 5 sites.

## Simulations ( $1 / 3$ )

We simulated the absence/presence of 3 fish species, depending on the temperature and salinity of the water, over 5 sites.




Then, four of these sites are used for estimation, the last one for testing.

## Simulations ( $1 / 3$ )

We simulated the absence/presence of 3 fish species, depending on the temperature and salinity of the water, over 5 sites.




Then, four of these sites are used for estimation, the last one for testing. Here, we have

$$
\lambda_{t}=A \cdot Y_{t-1}+B \cdot X_{t-1},
$$

where $\left(X_{t}\right)_{t \in \mathbb{Z}}$ is the process composed by the temperature and salinity.

## Simulations (2/3)

## We obtain the following estimations results

## Simulations (2/3)

We obtain the following estimations results

$$
A=\left(\begin{array}{ccc}
0.2 & 0.1 & -0.2 \\
0.5 & 0.1 & -0.2 \\
-0.5 & 0.3 & 0.2
\end{array}\right) \quad \text { and } \quad \hat{A}=\left(\begin{array}{ccc}
0.296 & -0.499 & -0.590 \\
0.444 & 0.320 & -0.138 \\
-0.183 & 0.385 & 0.198
\end{array}\right)
$$

## Simulations (2/3)

We obtain the following estimations results

$$
\begin{gathered}
A=\left(\begin{array}{ccc}
0.2 & 0.1 & -0.2 \\
0.5 & 0.1 & -0.2 \\
-0.5 & 0.3 & 0.2
\end{array}\right) \quad \text { and } \quad \hat{A}=\left(\begin{array}{ccc}
0.296 & -0.499 & -0.590 \\
0.444 & 0.320 & -0.138 \\
-0.183 & 0.385 & 0.198
\end{array}\right) \\
B=\left(\begin{array}{cc}
0.5 & -0.1 \\
0.2 & -0.1 \\
-0.3 & 0.1
\end{array}\right) \quad \text { and } \quad \hat{B}=\left(\begin{array}{cc}
0.582 & -0.118 \\
0.232 & -0.110 \\
-0.317 & 0.096
\end{array}\right)
\end{gathered}
$$

## Simulations (2/3)

We obtain the following estimations results

$$
A=\left(\begin{array}{ccc}
0.2 & 0.1 & -0.2 \\
0.5 & 0.1 & -0.2 \\
-0.5 & 0.3 & 0.2
\end{array}\right) \quad \text { and } \quad \hat{A}=\left(\begin{array}{ccc}
0.296 & -0.499 & -0.590 \\
0.444 & 0.320 & -0.138 \\
-0.183 & 0.385 & 0.198
\end{array}\right)
$$

$$
B=\left(\begin{array}{cc}
0.5 & -0.1 \\
0.2 & -0.1 \\
-0.3 & 0.1
\end{array}\right) \quad \text { and } \quad \hat{B}=\left(\begin{array}{cc}
0.582 & -0.118 \\
0.232 & -0.110 \\
-0.317 & 0.096
\end{array}\right)
$$

and
$R=\left(\begin{array}{ccc}1 & 0.2 & -0.5 \\ 0.2 & 1 & -0.3 \\ -0.5 & -0.3 & 1\end{array}\right) \quad$ and $\quad \hat{R}=\left(\begin{array}{ccc}1 & 0.204 & -0.436 \\ 0.204 & 1 & -0.303 \\ -0.436 & 0.204 & 1\end{array}\right)$.

## Simulations (3/3)

We then make previsions at horizon 1 for the testing site, and obtain the following accuracy.

## Simulations (3/3)

We then make previsions at horizon 1 for the testing site, and obtain the following accuracy.

|  | Species 1 | Species 2 | Species 3 |
| :---: | :---: | :---: | :---: |
| Accuracy | $79.3 \%$ | $83.7 \%$ | $78.3 \%$ |
| Mean Presence | $73.5 \%$ | $15.7 \%$ | $57.8 \%$ |

## Simulations (3/3)

We then make previsions at horizon 1 for the testing site, and obtain the following accuracy.

|  | Species 1 | Species 2 | Species 3 |
| :---: | :---: | :---: | :---: |
| Accuracy | $79.3 \%$ | $83.7 \%$ | $78.3 \%$ |
| Mean Presence | $73.5 \%$ | $15.7 \%$ | $57.8 \%$ |



## Real data (1/2)

The previous simulation is based upon a real dataset collected by the government of Scotland: https://data.marine.gov.scot/.

## Real data (1/2)

The previous simulation is based upon a real dataset collected by the government of Scotland: https://data.marine.gov.scot/.

We study here the absence/presence of two aquatic micro-organisms: Alexandrium and Dinophysis.

## Real data (1/2)

The previous simulation is based upon a real dataset collected by the government of Scotland: https://data.marine.gov.scot/.

## $\longrightarrow \mid$ Scottish Government Riaghaltas na h-Alba <br> gov.scot <br> marinescotland

We study here the absence/presence of two aquatic micro-organisms:
Alexandrium and Dinophysis.
The data were collected monthly from 1997 to 2013 on 5 different locations in Scotland, and we have access to the covariates: Temperature, Salinity and Oxidised Nitrogen.

## Real data (2/2)

Once again, 4 sites were used for estimation, and we use the last site to perform previsions at horizon 1.

## Real data (2/2)

Once again, 4 sites were used for estimation, and we use the last site to perform previsions at horizon 1.

We obtain the following accuracy.

|  | Alexandrium | Dinophysis |
| :---: | :---: | :---: |
| Accuracy | $72.7 \%$ | $75.0 \%$ |
| Mean Presence | $65.2 \%$ | $64.0 \%$ |

## Real data (2/2)

Once again, 4 sites were used for estimation, and we use the last site to perform previsions at horizon 1.
We obtain the following accuracy.

|  | Alexandrium | Dinophysis |
| :---: | :---: | :---: |
| Accuracy | $72.7 \%$ | $75.0 \%$ |
| Mean Presence | $65.2 \%$ | $64.0 \%$ |



## Thank you !

