# Dynamic modeling of abundance data in ecology

### Guillaume Franchi

#### ENSAI, Bruz





### ECODEP Conference 12 February 2024

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### I. Introduction

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

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One can also consider the Shannon entropy

$$\forall y = (y_1, \dots, y_d) \in \mathcal{S}_{d-1}, \ I_S(y) = -\sum_{i=1}^d y_i \log(y_i).$$

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- Make predictions about the relative abundance of an ecosystem over time.
- Output Understand the dynamics of this ecosystem:
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  - → 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.

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We focus on three groups of species:



(a) Coleoptera

(b) Hymenoptera

(c) Diptera

Image: A matrix and a matrix

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

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where:

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

$$Y_t^- = (Y_t, Y_{t-1}, \ldots) \, .$$

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Introduction

# Chain with complete connections (2/2)

#### Remark

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we obtain a Markov chain.

• It is possible to add a process of exogenous variables  $(X_t)_{t\in\mathbb{Z}}$  to the dynamic of the process  $(Y_t)_{t\in\mathbb{Z}}$ 

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

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#### Theorem 1

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

$$\forall t \in \mathbb{Z}, \ \mathbb{P}(Y_{t+1} \in A \mid Y_t^- = y_t^-) = P(A \mid y_t^-).$$

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#### Assumptions

**A1** 
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We have  $\sum_{m \in \mathbb{N}} b_m < \infty$ .

## A natural proposal for $P(\cdot \mid Y_t^-)$ is a Dirichlet distribution

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

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- → its mean vector  $\lambda = (\lambda_1, \dots, \lambda_d)$ ;
- → a dispersion parameter  $\varphi > 0$ .

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# Dirichlet model (2/3)

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

$$\operatorname{alr}(\lambda_t) = \eta_0 + \sum_{k \ge 1} \eta_k \overline{Y}_{t-k} + \sum_{k \ge 1} \zeta_k X_{t-k},$$

where  $\operatorname{alr}$  is the mapping

alr : 
$$\mathcal{S}_{d-1} \longrightarrow \mathbb{R}^{d-1}$$
  
 $y = (y_1, \dots, y_d) \longmapsto \left( \log \left( \frac{y_1}{y_d} \right), \dots, \log \left( \frac{y_{d-1}}{y_d} \right) \right),$ 

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→ We also propose that for all  $t \in \mathbb{Z}$ 

$$\varphi_t = \exp\left(\theta_0 + \sum_{k \ge 1} \theta_k I_S(Y_{t-k+1})\right),$$

where the  $\theta_k$ 's are real numbers.

Dirichlet model (3/3)

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 $\Im$  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.

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

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- $\bigcirc$  Model the absence/presence of species in the ecosystem.



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 $\mathbb{P}$  It is actually a dynamic version of a multivariate probit regression.

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#### Theorem 2

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There exists a strongly stationary process  $(Y_t)_{t\in\mathbb{Z}}$  satisfying

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In addition, its distribution is unique.

#### Remark

Furthermore, if  $(\zeta_t)_{t\in\mathbb{Z}}$  is ergodic,  $(Y_t)_{t\in\mathbb{Z}}$  is also ergodic.

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## Estimation results (1/3)

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

$$\theta = (A_1, \ldots, A_p, B, R).$$

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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

$$I_{Y_{i,t}} = \begin{cases} \ ]0, +\infty[ & \text{if } Y_{i,t} = 1 \\ \ ]-\infty, 0] & \text{if } Y_{i,t} = 0 \end{cases}$$

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X Difficult function to optimize...

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

→ We then maximize all pairwise conditional likelihoods

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

## 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

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

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

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

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Usually, if we consider the estimator

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

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

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and its asymptotic normality on

$$\mathbb{E}\left(\|\dot{m}_{\theta_0}(Z_0)\|^2\right) < +\infty.$$

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

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

Image: A matrix and a matrix

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by modifying the "order conditions"

$$\mathbb{E}\left(\sup_{\theta} |m_{\theta}(Z_{0,0})|^{1+\delta}\right) < +\infty \quad \text{and} \quad \mathbb{E}\left(\|\dot{m}_{\theta_0}(Z_{0,0})\|^{2(1+\delta)}\right) < +\infty;$$

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→ and by adding the following "order condition"

$$\mathbb{E}\left(\|\ddot{m}_{\theta_0}(Z_{0,0})\|^{1+\delta}\right) < +\infty$$

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

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

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

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and

$$\widehat{R}(i_1, i_2) = \operatorname{argmax} \sum_{j=1}^n \sum_{t=p+1}^T \log \int_{I_{Y_{i_1,j,t}} - \hat{\lambda}_{i_1,j,t}} \Phi\left( (2Y_{i_2,j,t} - 1) \frac{\hat{\lambda}_{i_2,j,t} + rx_{i_1}}{\sqrt{1 - r^2}} \right) \varphi(x_{i_1}) \mathrm{d}x_{i_1}$$

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

#### Proposition 2

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

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

$$\sqrt{n(T-p)}\left(\hat{\theta}-\theta_0\right) \quad \text{and} \quad \sqrt{n(T-p)}\left(\hat{\gamma}-\gamma_0, \widehat{R}-R_0\right).$$

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

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

We obtain the following estimations results

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$$A = \begin{pmatrix} 0.2 & 0.1 & -0.2 \\ 0.5 & 0.1 & -0.2 \\ -0.5 & 0.3 & 0.2 \end{pmatrix} \quad \text{and} \quad \hat{A} = \begin{pmatrix} 0.296 & -0.499 & -0.590 \\ 0.444 & 0.320 & -0.138 \\ -0.183 & 0.385 & 0.198 \end{pmatrix},$$

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$$B = \begin{pmatrix} 0.5 & -0.1 \\ 0.2 & -0.1 \\ -0.3 & 0.1 \end{pmatrix} \text{ and } \hat{B} = \begin{pmatrix} 0.582 & -0.118 \\ 0.232 & -0.110 \\ -0.317 & 0.096 \end{pmatrix},$$

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and

$$R = \begin{pmatrix} 1 & 0.2 & -0.5 \\ 0.2 & 1 & -0.3 \\ -0.5 & -0.3 & 1 \end{pmatrix} \text{ and } \hat{R} = \begin{pmatrix} 1 & 0.204 & -0.436 \\ 0.204 & 1 & -0.303 \\ -0.436 & 0.204 & 1 \end{pmatrix}$$

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

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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%

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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.

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## Real data (2/2)

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

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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%

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	Alexandrium	Dinophysis
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Observations
Previsions



# Thank you !

Guillaume Franchi

Dynamic modeling of abundance data in ecology 33/33

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