

# Finite mixture of size projection matrix models for highly diverse rainforests in a variable environment

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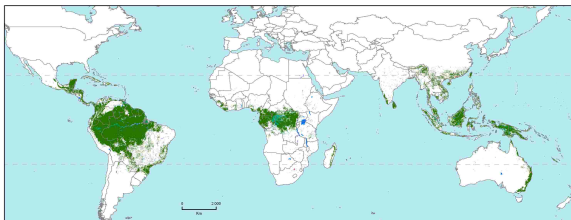
# Outline

- 1 Ecological Context
- 2 Forest dynamics models
  - The Usher model
  - Environmental variability
  - High biodiversity
- 3 The M'Baïki experimental site: unique data in Central Africa
  - Mixture models outputs
  - Dynamics species groups
- 4 Conclusions

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# Tropical forests I



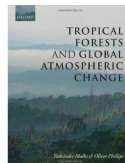
## Generality and Specificity

- ❶ 6% of the world's surface area
- ❷ 50% of all living organisms on Earth
- ❸ more than 1.5b people depending on the forest

# Tropical forests II

## Multiple uses and actors

- local: resource (wood, NWFP)
- national: foreign exchange (timber)
- global scale: CO<sub>2</sub> concentration regulation through carbon sequestration  
(Millennium Ecosystem Assessment, 2005)



# Tropical forests III

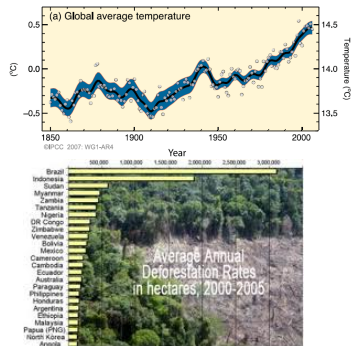
## Conservation and management

### increasing atmospheric CO<sub>2</sub> concentration:

- increasing global temperatures
- climate change Solomon et al. (2007)

### increasing land uses

- Agricultural, fuel
- ... FAO (2010)

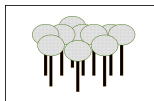


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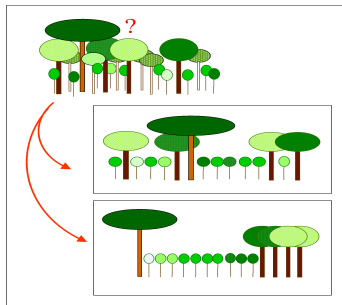
# Different type of models I

Stand models: all trees are equivalents

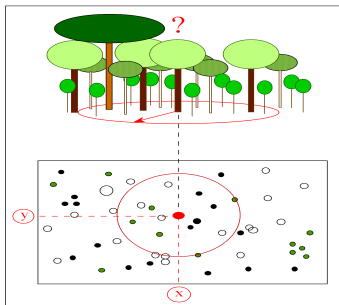


$$(V, G, N) = f(\text{age, species, site, density})$$

Individual tree models



distance independant

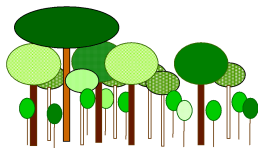


distance dependent

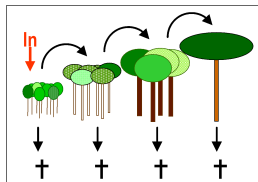


# Different type of models II

## Size projection matrix



Discrete distributions: tree equivalents inside class



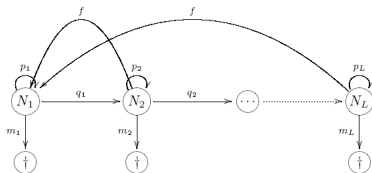
# The Usher matrix I

## The Usher model (Usher, 1966, 1969)

- matrix population model for size-structured populations
- describes the evolution of the population by a vector  $N(t)$
- $N(t)$  vector of the number of individuals in  $L$  ordered state class at discrete time  $t$ .
- This matrix population model relies on the four following hypotheses:
  - Hypothesis of independence: evolution of individuals is independent.
  - Markov hypothesis: evolution of an individual between two time steps  $t$  and  $t + 1$  only depends on its state at  $t$ .
  - Usher hypothesis: during each time step, an individual can stay in the same class, move up a class, or die; each individual may give birth to a number of offspring.
  - Hypothesis of stationarity: evolution of individuals between two time steps is independent of time.

# The Usher matrix II

graphical representation of the Usher model



$$\mathbb{E}(\mathbf{N}_{t+1} | \mathbf{N}_t) = \mathbf{P} \mathbf{S} \mathbb{E} \mathbf{N}_t + \mathbf{R}$$

$$\mathbf{P} = \begin{pmatrix} 1 - q_2^\bullet & 0 & 0 \\ q_2^\bullet & \ddots & \vdots \\ 0 & \ddots & 1 - q_L^\bullet & 0 \\ & & q_L^\bullet & 1 \end{pmatrix} \quad \mathbf{S} = \begin{pmatrix} 1 - m_1 & 0 \\ & \ddots \\ 0 & 1 - m_L \end{pmatrix} \quad \mathbf{R} = \begin{pmatrix} r \\ 0 \\ \vdots \\ 0 \end{pmatrix}$$

- $q_{i+1}^\bullet$  conditional upgrowth rate
- $m_i$  probability to die
- $r$  number of recruited trees

# Environment and Usher model I

Temporal changes of the diameter distribution of a tree population

$$\mathbf{N}(t+1) = \mathbf{P}(X_t)\mathbf{S}(X_t)\mathbf{N}(t) + \mathbf{R}(X_t)$$

$$\mathbf{P}(X_t) = \begin{pmatrix} 1 - q_2^\bullet(X_t) & 0 & 0 \\ q_2^\bullet(X_t) & \ddots & \vdots \\ 0 & \ddots & 1 - q_l^\bullet(X_t) & 0 \\ & & q_l^\bullet(X_t) & 1 \end{pmatrix}$$

$$\mathbf{S}(X_t) = \begin{pmatrix} 1 - m_1(X_t) & 0 \\ & \ddots \\ 0 & 1 - m_l(X_t) \end{pmatrix}$$

$$\mathbf{R}(X_t) = \begin{pmatrix} r(X_t) \\ 0 \\ \vdots \\ 0 \end{pmatrix}$$

- $q_{i+1}^\bullet(X_t)$   
conditional upgrowth rate
- $m_i(X_t)$  probability to die
- $r(X_t)$  number of recruited trees

## Regression estimator

(Rogers-Bennett and Rogers, 2006; Picard et al., 2008; Zetlaoui, 2006)

$$q_{i+1}^\bullet(X_t) = \frac{\Delta D_i(X_t)}{d_i}$$

$\Delta D_i(X_t)$  "typical" dbh growth rate  
 $d_i$  width of diameter class  $i$

# Environment and Usher model II

## Modeling growth, mortality and recruitment

- Let  $\Delta D$  be the Diameter increment,

$$\Delta D_{sti} = \mathbf{x}_{ti}^D \beta_s + \varepsilon_s$$

where  $\beta_s$  is the vector of unknown parameters,  $\mathbf{x}^D = (\mathbf{x}_{ti}^D)_{t,i}$  the known incidence

- Let  $N_{st}$  be the number of recruits:

$$\begin{aligned} N_{st} &\sim \mathcal{P}(r_{st}) \\ \log(r_{st}) &= \mathbf{x}_t^N \gamma_s \end{aligned}$$

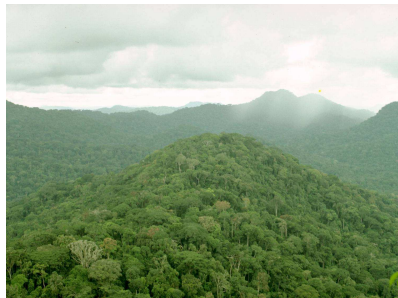
- Let  $M_{sti}$  be the mortality

$$\begin{aligned} M_{sti} &\sim \mathcal{Ber}(m_{sti}) \\ \text{logit}(m_{sti}) &= \mathbf{x}_{ti}^M \alpha_s \end{aligned}$$

# Main concern with tropical forests: high specific richness

## + 300 species/ha

- many species with very few individuals
- high intra specific variability of dynamic variables
- ➔ one species one model: poor fit of models



👉 **building groups of species according to their dynamics**

# Mixture of growth, mortality and recruitment processes

## Mixture models

- For growth and mortality processes, let assumed:

$$\ell_n(\psi|\mathbf{Y}) = \sum_{s=1}^S \sum_{t=1}^T \sum_{i=1}^{n_{st}} \log \left[ \sum_{k=1}^K \pi_k f(Y_{sti}|\mathbf{X}, \psi_k) \right]$$

with  $f$  Gaussian density and  $Y_{sti} = \Delta D_{sti}$  or the masse function associated to the Bernoulli distribution and  $Y_{sti} = M_{sti}$ .

- For the recruitment process, let assumed

$$\ell_n(\psi|\mathbf{Y}) = \sum_{s=1}^S \sum_{t=1}^T \log \left[ \sum_{k=1}^K \pi_k f(N_{st}|\mathbf{X}, \psi_k) \right]$$

where  $f$  is the masse function associated to the Poisson distribution.

# Mixture Models and variable selection

Co-variates can differ from a group to another  
stepwise selection can be computational intensive

□ LASSO (**L**east **A**bsolute **S**hrinkage & **S**election **O**perator)

Selection (Khalili and Chen, 2007)

The estimator  $\hat{\psi}$  of the model's parameters  $\psi$  corresponds to the maximum of a penalized version of the log-likelihood:

$$\hat{\psi} = \arg \max_{\psi} \left\{ \ell_n(\psi | \mathbf{Y}) - p_n(\psi) \right\}$$

where  $p_n$  is a penalty. We used the lasso penalization to perform variable selection in each component:

$$p_n(\psi) = \sum_{k=1}^K \pi_k \left( \sum_{j=1}^{p+1} \gamma_{nk} \sqrt{n} |\beta_{kj}| \right)$$

where  $\beta_{kj}$  is the  $j$ th element of  $\beta_k$  and  $\gamma_{nk}$  is a tuning parameter.



# EM algorithm with LASSO

- **Estimate**  $\hat{c}_{sik} = P \{ \text{tree of species } s \in k \mid \text{data } (\mathcal{X}), \text{ parameters } (\psi) \}$   
to start we choose randomly  $\hat{c}_{sik}$

- **Maximise** the penalized log likelihood (Khalili and Chen, 2007)

$$\log L^*(\psi \mid \mathcal{X}) = \log L(\psi \mid \mathcal{X}) - p(\psi \mid \mathcal{X})$$

with  $p(\psi \mid \mathcal{X})$  penalty term

covariates selection by shrinkage : coefficient  $\beta_{kp} < \text{threshold} = 0$

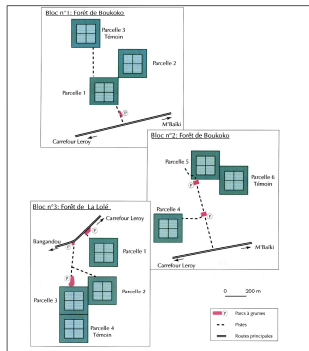
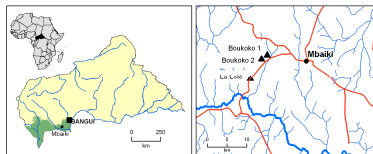
$$\rightarrow \hat{\psi}_k \rightarrow \hat{\pi}_k$$

- **Classification of species** (not individuals) into groups  
 $\rightarrow$  tree  $i$  of species  $s$  belongs to groups  $k$  for  $\max(\hat{c}_{sik})$

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# The M'Baïki experimental site (Central African Republic)



N. Faure CIRAD - UR 105  
Septembre 2011

- Permanent sample plots (annual)
- 1982
- 40 ha
- semi-deciduous forest
- 239 species / morphospecies
- silvicultural treatments  
(undisturbed,  
logging,  
logging + thinning)  
→ disturbance gradient

# The M'Baïki experimental site (2)

Every year since 1982 (except in 1997, 1999, 2001), all trees  $\geq 10$  cm diameter at breast height (dbh)

- individually marked
- measured for dbh
- mapped
- identified

+ inventory of dead trees and newly recruited trees with dbh  $\geq 10$  cm  
annual diameter growth, mortality, and recruitment

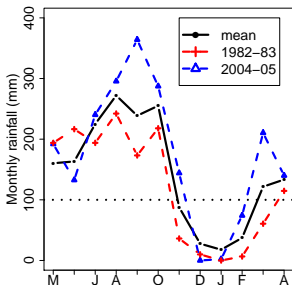
## Data

- growth, mortality more than 200,000 observations
- recruitment more than 100,000 observations
- more than 200 species



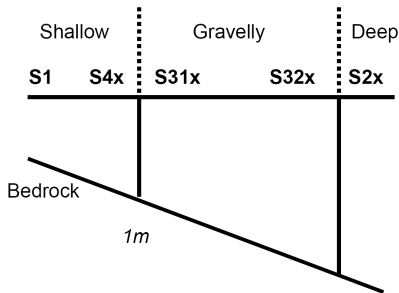
# Covariates (1) : Drought indices

- Length of *the dry season* (**LDS**, nb months)
- Average rainfall during *the dry season* (**RDS**, mm)



- Average annual soil water content (**MSW**, mm)  
simple water balanced model  
5 soil depth categories

$$SW_{t+1} = SW_t + P_t - E_t$$



Positive correlation

## Covariates (2): Light availability indices

Indirectly measured

- stand basal area (**BAst**,  $\text{m}^2 \text{ha}^{-1}$ )
- stand density (**Dst**, number of stems  $\text{ha}^{-1}$ )

Positive correlation

## Covariates (3): Tree development stage

- tree diameter ( $D_i$ , cm)



# Mixture models outputs I

## Separate species groups for growth, mortality, recruitment

- Species groups **characteristics**
- Species groups **response to drought**  
Coefficients

$$\begin{aligned}\Delta D_{kti} &= \mathbf{X}_{ti}^D \beta_k + \varepsilon_k \\ \text{logit}(m_{kti}) &= \mathbf{X}_{ti}^M \alpha_k \\ \log(r_{kt}) &= \mathbf{X}_t^N \gamma_k\end{aligned}$$

### Covariates

Length of the dry season (**LDS**)

Average rainfall during the dry season (**RDS**)

Average annual soil water content (**MSW**)



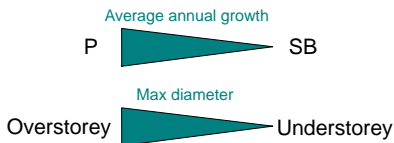
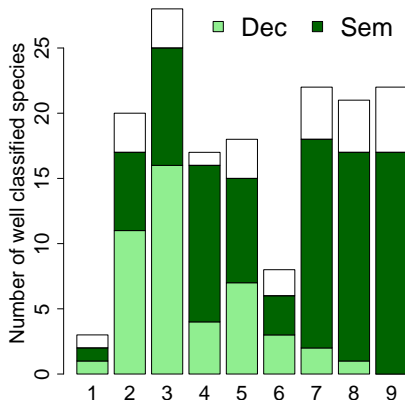
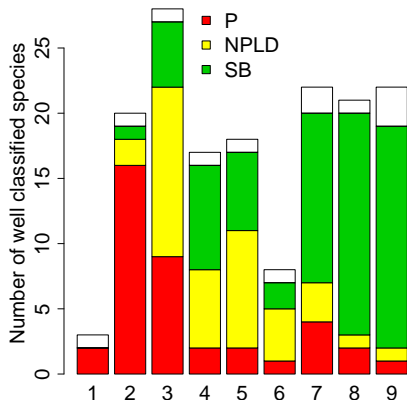
# Mixture models outputs II

## Classification results

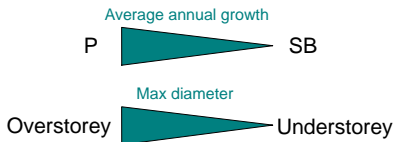
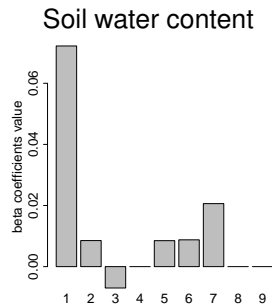
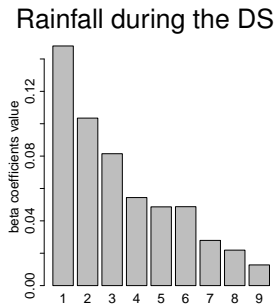
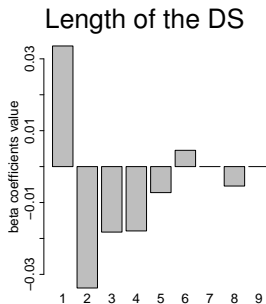
- 9 **growth species groups**
- 3 **mortality species groups**
- 5 **recruitment species groups**

54 non-empty matrix population groups

# 9 growth species groups

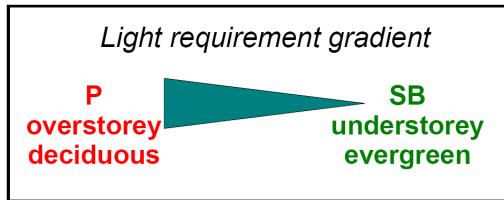


# Growth species groups response to drought



# Comments I

Drought indices: capture several effects



Average annual **growth**

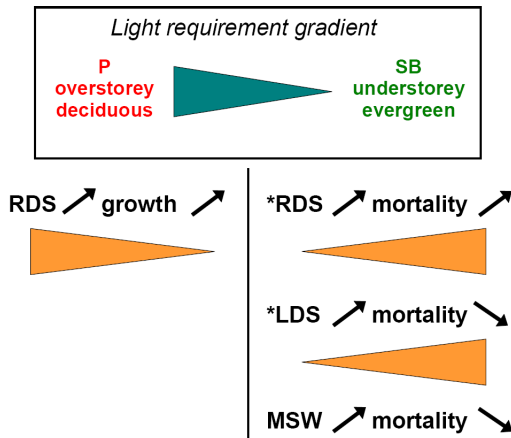


Average annual **mortality**



# Comments II

## Drought indices: capture several effects



- LDS/RDS**

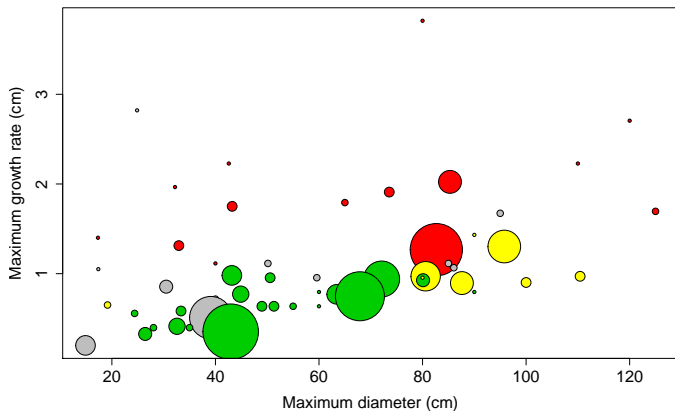
indirect measure of light availability for the understorey

(Lingenfelder and Newbery, 2009;  
Newbery et al., 2011)

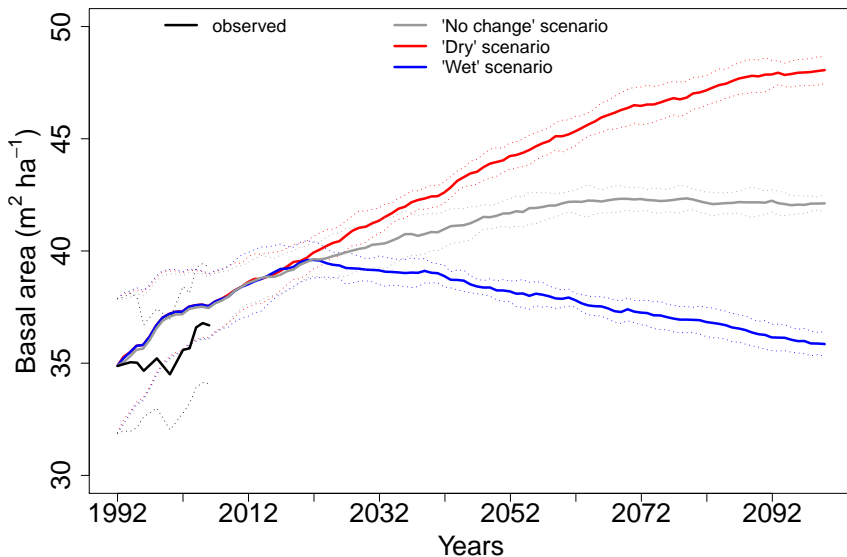
- MSW**

measure of water stress

# Dynamics species groups



# Dynamics response to drought: basal area



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# Conclusions I

- **A species classification without *a priori***

based on (growth + mortality + recruitment) response to  
light, drought (+ size)

⇒ leads to ecological groups of tree species

- M'Baïki **undisturbed** forest seems to be resilient to drought  
+ pioneer trees more sensitive to drought  
Disturbance increased the proportion of pioneer trees

Ouédraogo et al. (2011)

↪ **combination of logging + drought disturbance?**

# Conclusions II

- **A powerful method for species classification**

- species classification
- estimation of the response to light, drought, and size
- covariates selection

**simultaneously!**

- but Estimation and selection realized process by process

➡ **How to combine estimation, classification and selection for the three processes simultaneously?**

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