

E²M²: Ecological and Epidemiological Modeling in Madagascar

Looking back: How far have we come?

Rétrospective : où en sommes-nous ?

Centre ValBio

Ranomafana National Park, Madagascar

December 2022

Friday: R Bootcamp

- Intro to R Studio
- Exploring and Visualizing Data in R
- For-loops, Functions, and If-Else Statements

Thursday: “Bells and Whistle”

- Introduction to Phylogenetic Modeling • Tutorial
- Introduction to Spatial Visualization and Plotting • Tutorial
- Modeling Insights from the Metapopulation Game • Tutorial
- Introduction to Network Modeling • Tutorial
- Research Snapshots

- Programming
- Data
- Models
- Research Development

Saturday: Travel

Sunday: “Getting Started with the Basics”

- Data and Models
- Student introductions & presentations
- Linear regression • Tutorial
- Formulating research questions

Monday: “Using Models with Data”

- Model-Guided Study Design
- Study design tutorial
- Intro to Compartmental Models & Differential Equations
- Building mechanistic models in R
- Refining research questions for modeling
- Defining a model world

Wednesday: “Refining Your Work”

- Model Fitting in Practice – the Basic Concept
- Epidemic Cards
- Model Fitting with Epidemic Cards
- Model Selection and Comparison
- Model Selection Tutorial
- Model Telephone

Tuesday: “Applying Simple Models”

- Dynamical Fever
- Introduction to Mixed Modeling
- Reading a Research Paper

Friday: “Putting it All in Perspective”

- The Life Cycle of a Modeling Project
- C4C Student Presentations

Saturday: Travel

January: “Sharing Your Work”

- Final student presentations

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```
dens <- density(data, n = npts)
dx <- dens$x
dy <- dens$y
if(add == TRUE)
  plot(0., 0., main, ylab)
if(orientati
  dx2 <- (dx - min(dx)) / (max(dx) - min(dx))
  x[1.]
  dy2 <- (dy - min(dy)) / (max(dy) - min(dy))
  y[1.]
seqbelow <- rep(y[1.], length(dx))
if(Fill == T)
  confshade(dx2, seqbelow, dy2)
```



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What is science?

Qu'est ce que la science?

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What is science?

Qu'est ce que la science?

the **systematic observation** of natural events and conditions in order to **discover facts** about them and to **formulate laws and principles**

l'observation systématique des événements et des conditions naturelles afin de découvrir des faits à leur sujet et de formuler des lois et des principes fondés

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What are data?
Quelles sont les données?

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What are data?

Quelles sont les données?

evidence to support a claim

preuves à l'appui d'une réclamation

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What is a model?
Qu'est-ce qu'un modèle?

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What is a model?

Qu'est-ce qu'un modèle?

an abstract representation of a phenomenon

une représentation abstraite d'un phénomène

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Statistical
Statistique

Mechanistic
Mécaniste

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Statistical Statistique	Mechanistic Mécaniste
Correlation Corrélation	Causation Causalité
What? Que?	How? Comment?

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Formulating research questions
Formuler des questions de recherche

Statistical:

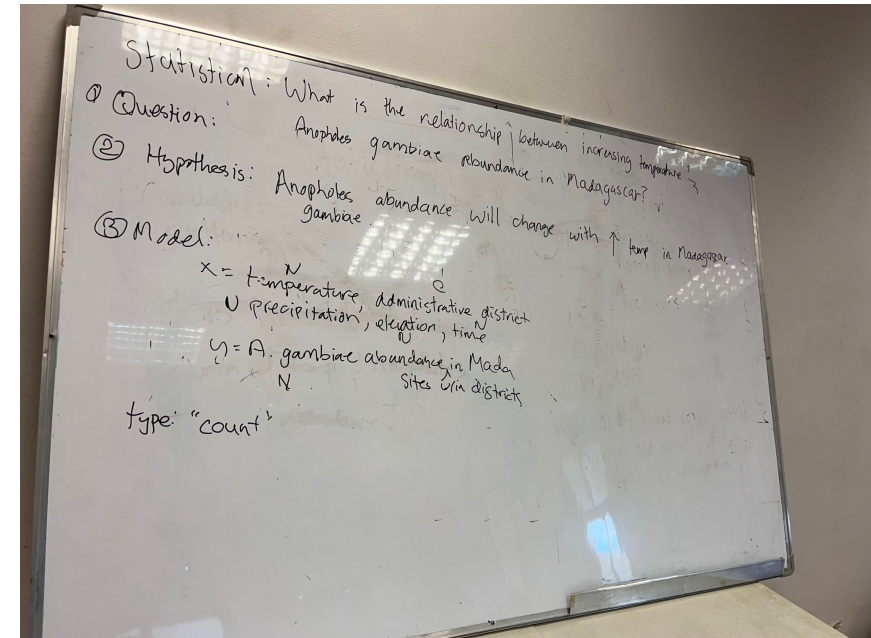
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Formulating research questions Formuler des questions de recherche

Statistical: What is the relationship between increasing temperature and *A. gambiae* occurrence in Madagascar?

Statistique : Quelle est la relation entre l'augmentation de la température et la présence d'*A. gambiae* à Madagascar ?

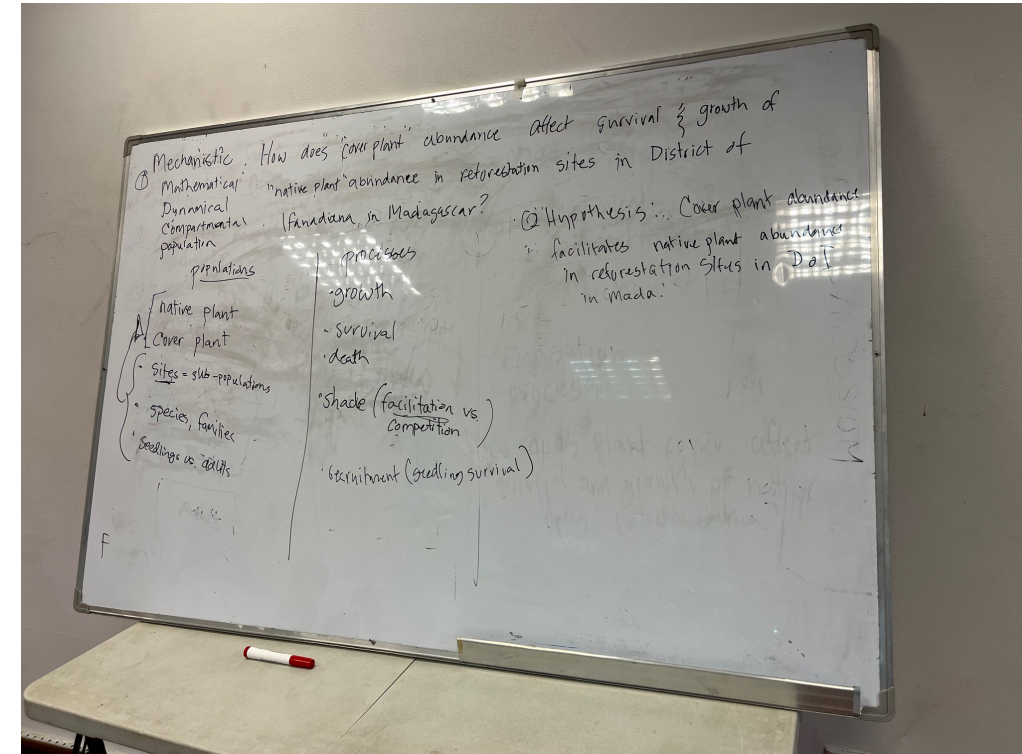


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Mechanistic:



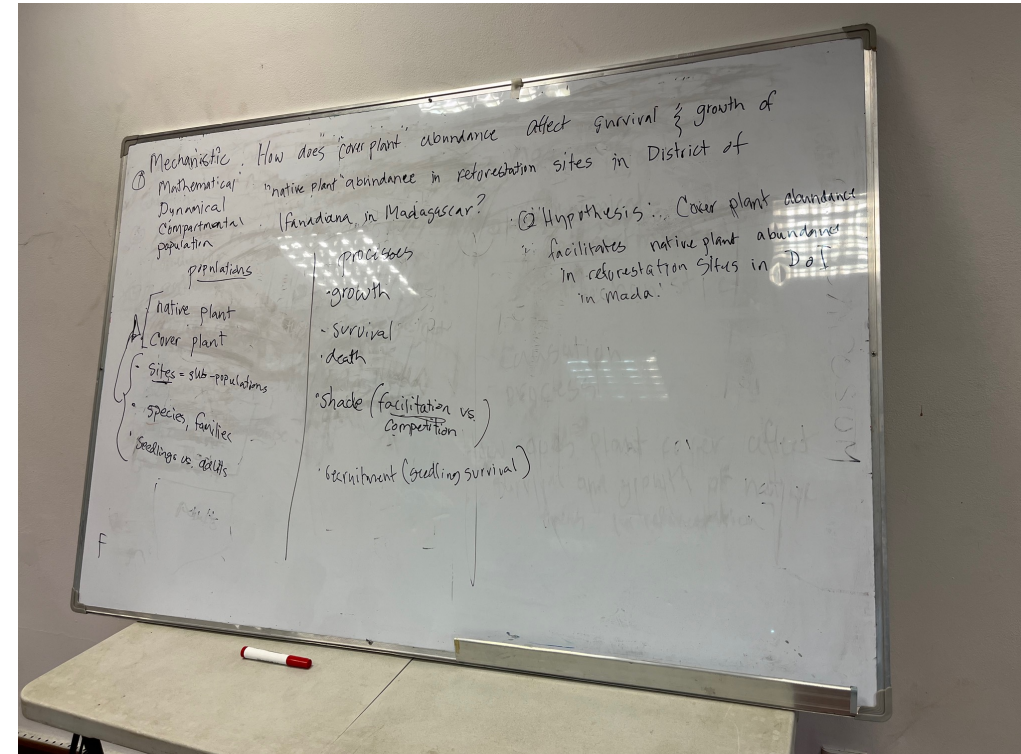
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Formulating research questions Formuler des questions de recherche

Mechanistic: How does cover plant abundance affect survival and growth of native plants in reforestation sites in the District of Ifanadiana?

Mécaniste : comment l'abondance des plantes de couverture affecte-t-elle la survie et la croissance des plantes indigènes dans les sites de reboisement du district d'Ifanadiana ?



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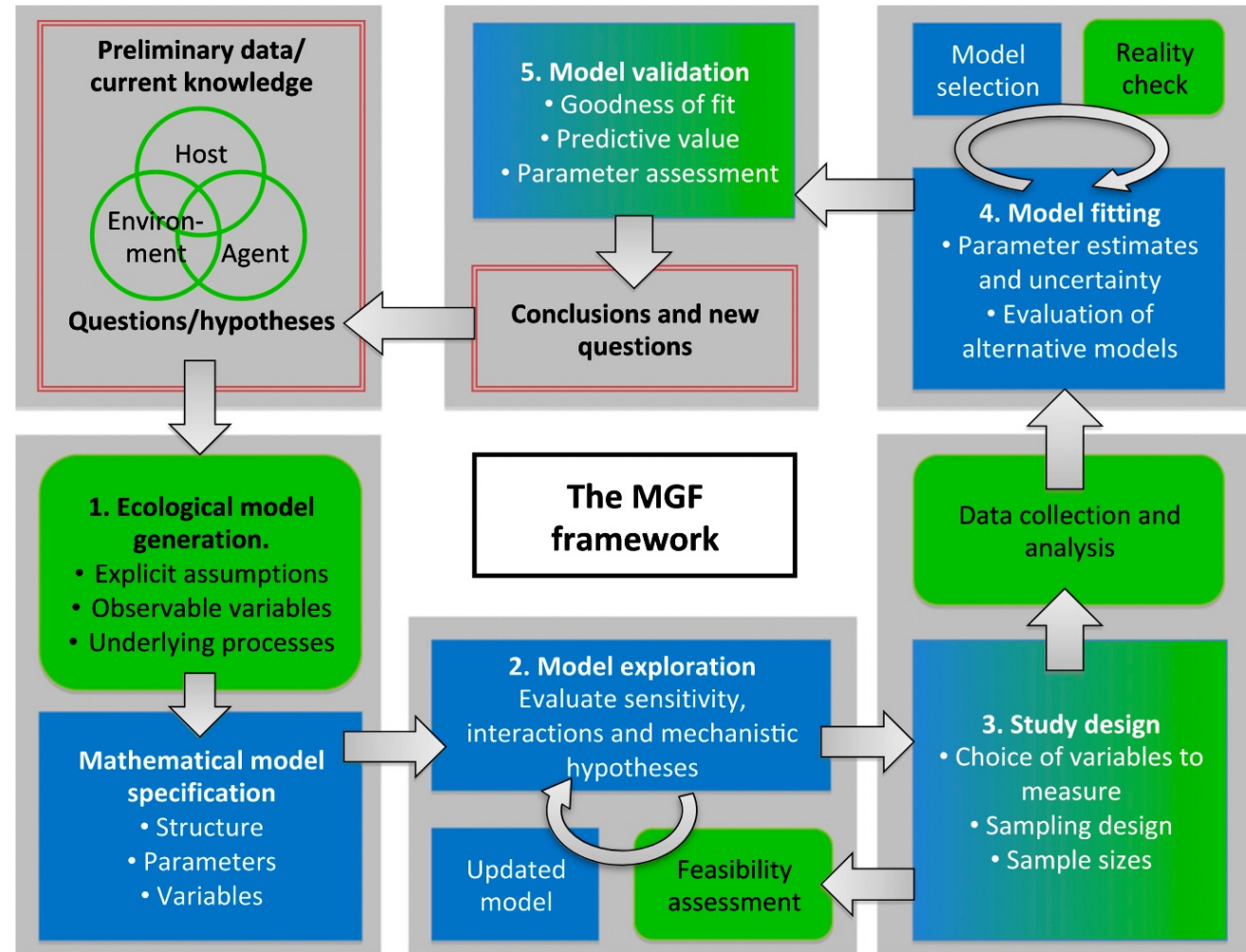
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Model-Guided Study Design

Conception d'étude guidée par modèle

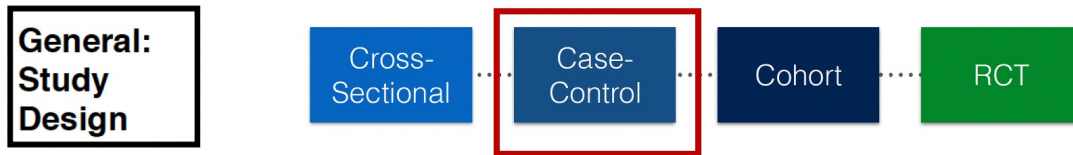


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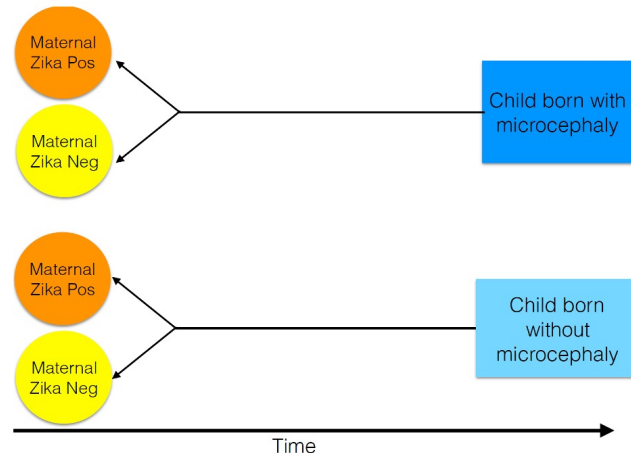
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Model-Guided Study Design

Conception d'étude guidée par modèle



Case Control Study



```
> # independent t-test
> pwr2t <- pwr.t2n.test(d=0.2,
+                       n1=175,
+                       power = .8,
+                       sig.level=0.05,
+                       alternative="greater")
> # inspect
> pwr2t
```

t test power calculation

```
n1 = 175
n2 = 1333.83
d = 0.2
sig.level = 0.05
power = 0.8
alternative = greater
```


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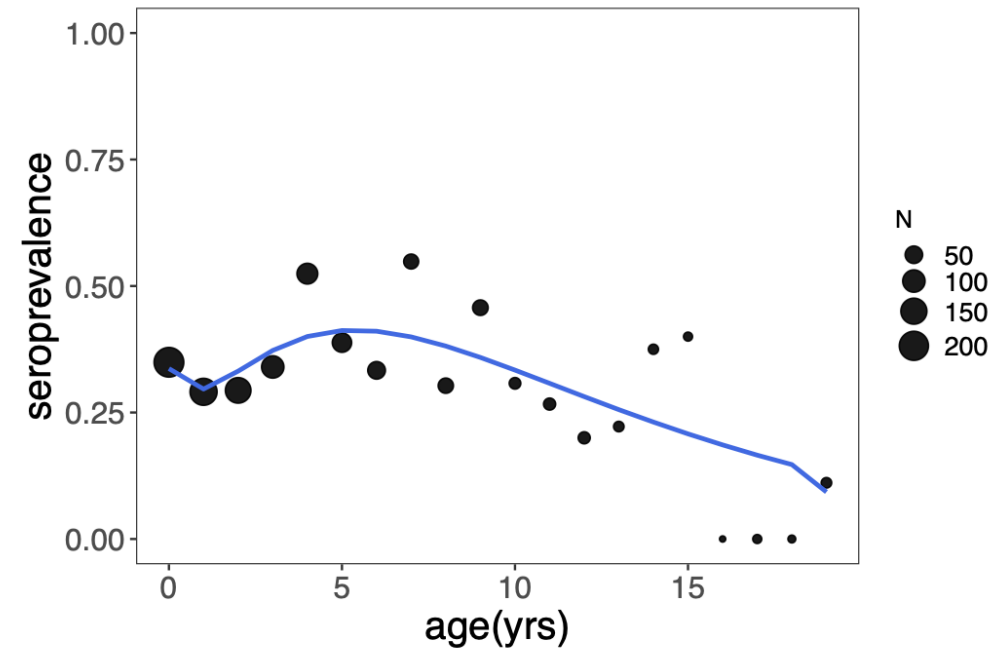
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Model-Guided Study Design

Conception d'étude guidée par modèle

Simulation and evaluation for mechanistic models

fitted lifelong immunity model (hyp1) to data subsample



AIC = 1277

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Intro to Compartmental Models

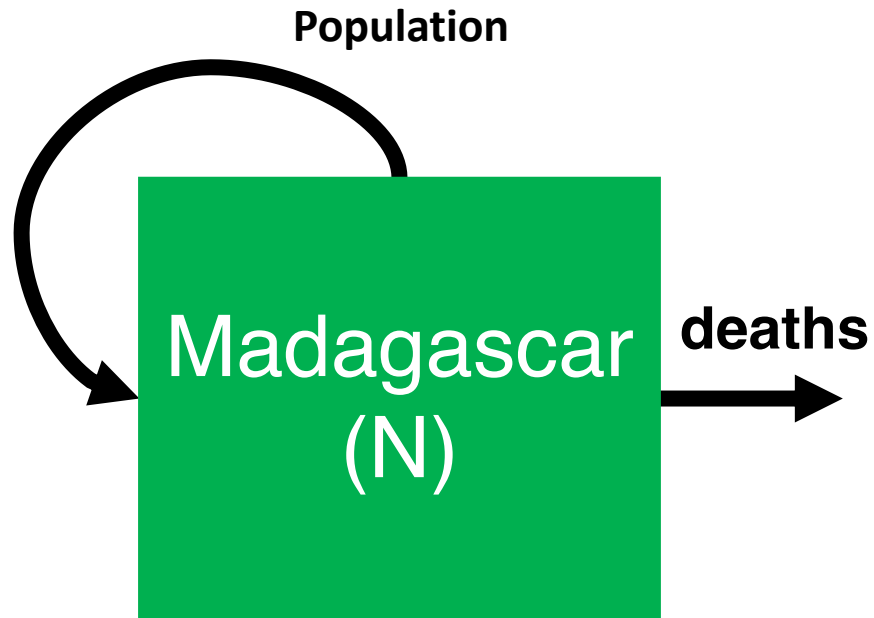
Introduction aux modèles compartimentés

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Intro to Compartmental Models

Introduction aux modèles compartimentés



$$N_{t+1} = \text{births} * N_t - \text{deaths} * N_t$$

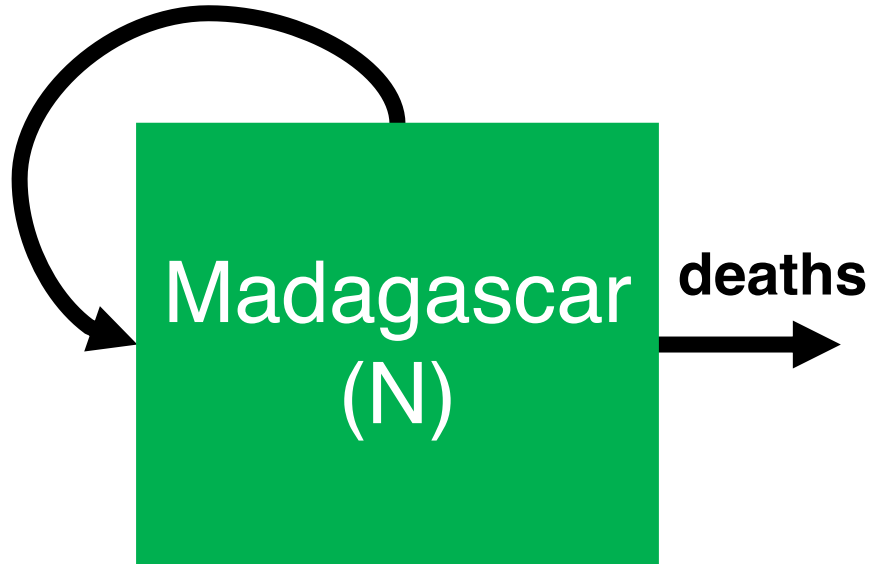
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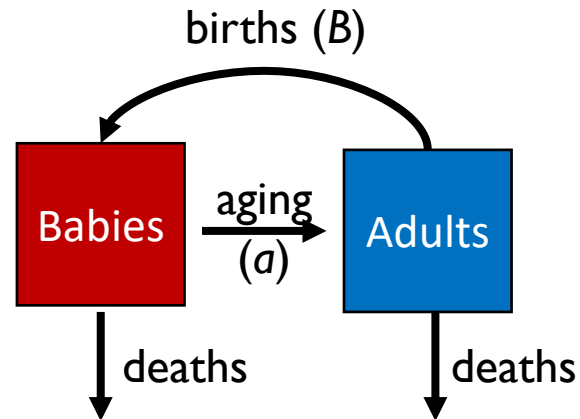
Introduction aux modèles compartimentés

Population



$$N_{t+1} = \text{births} * N_t - \text{deaths} * N_t$$

Structured Population



$$N_{t+1} = \lambda * N_t$$

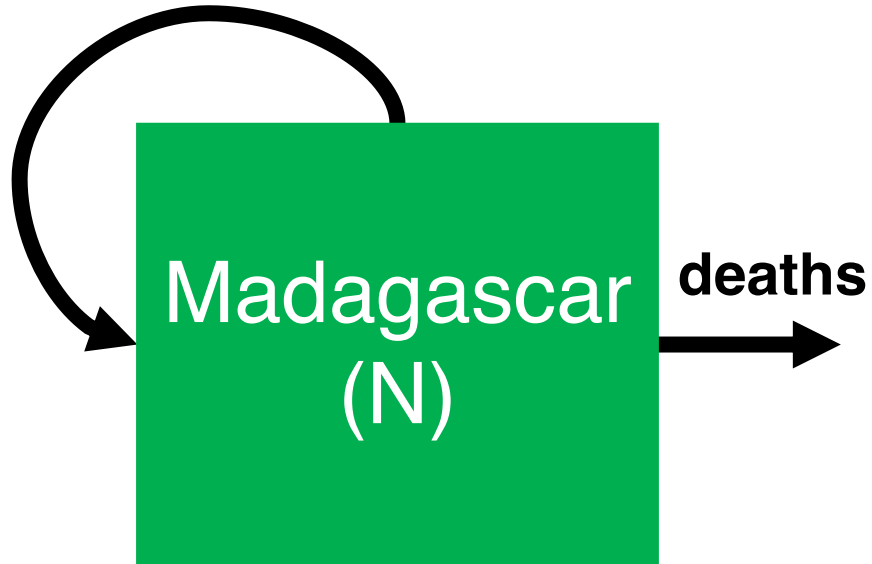
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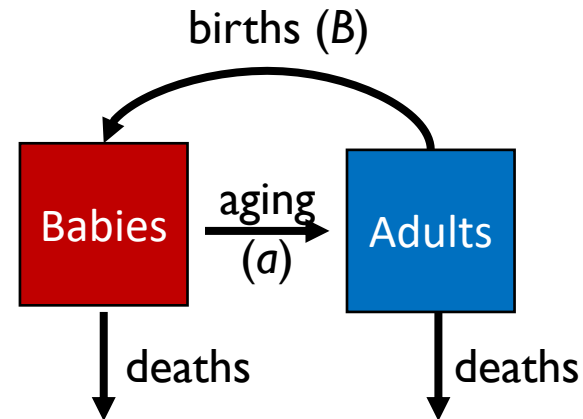
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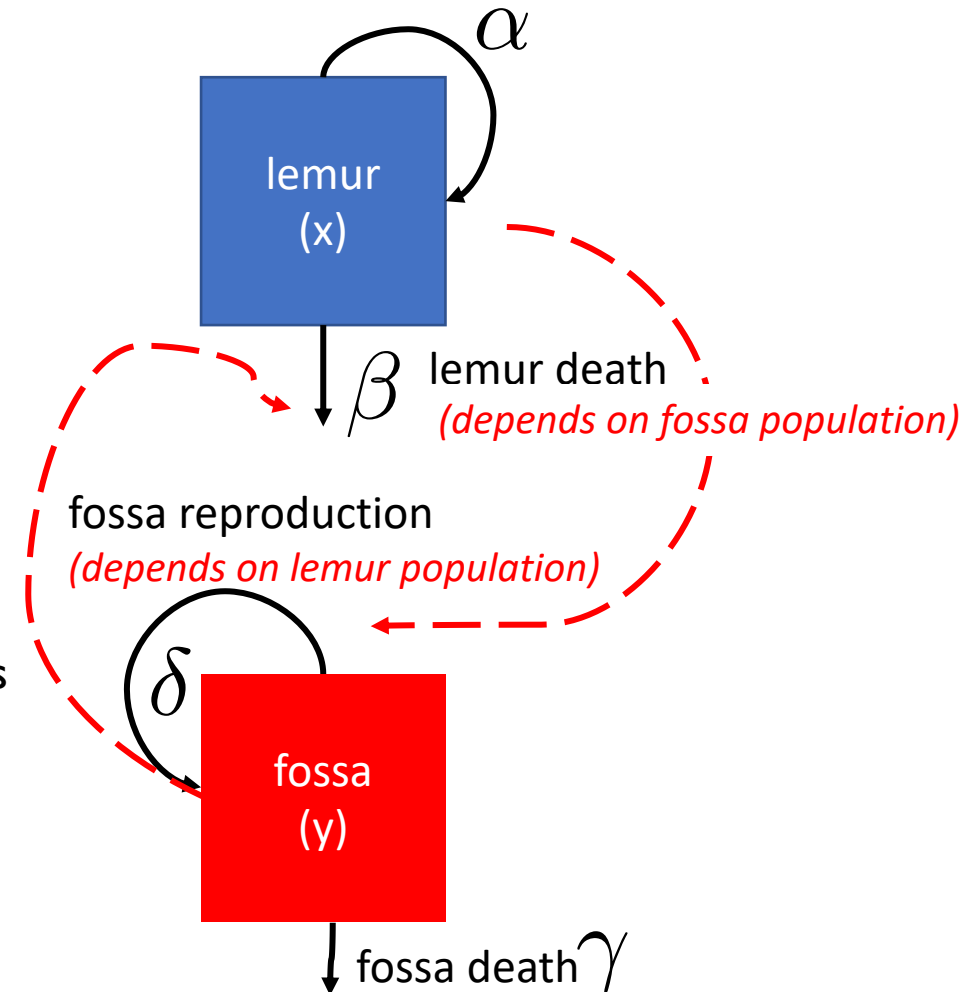
Structured Population



$$N_{t+1} = \lambda * N_t$$

Predator-Prey

lemur reproduction

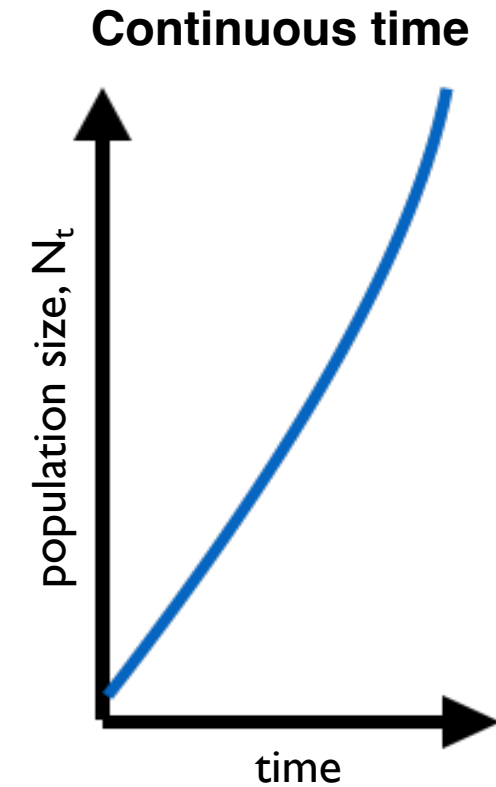
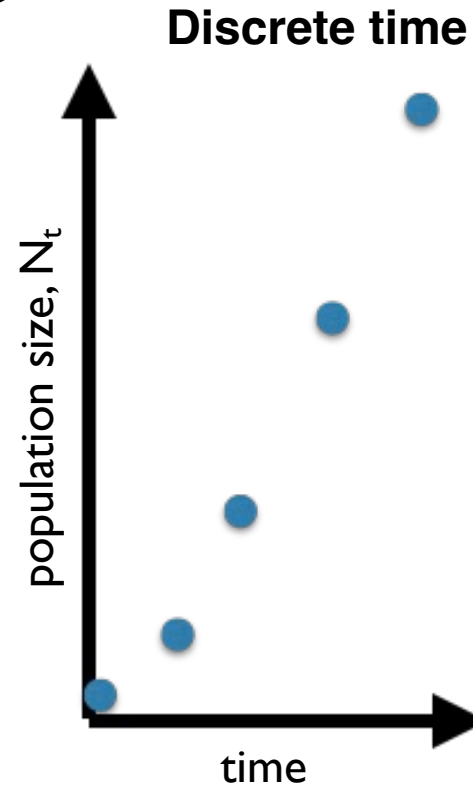


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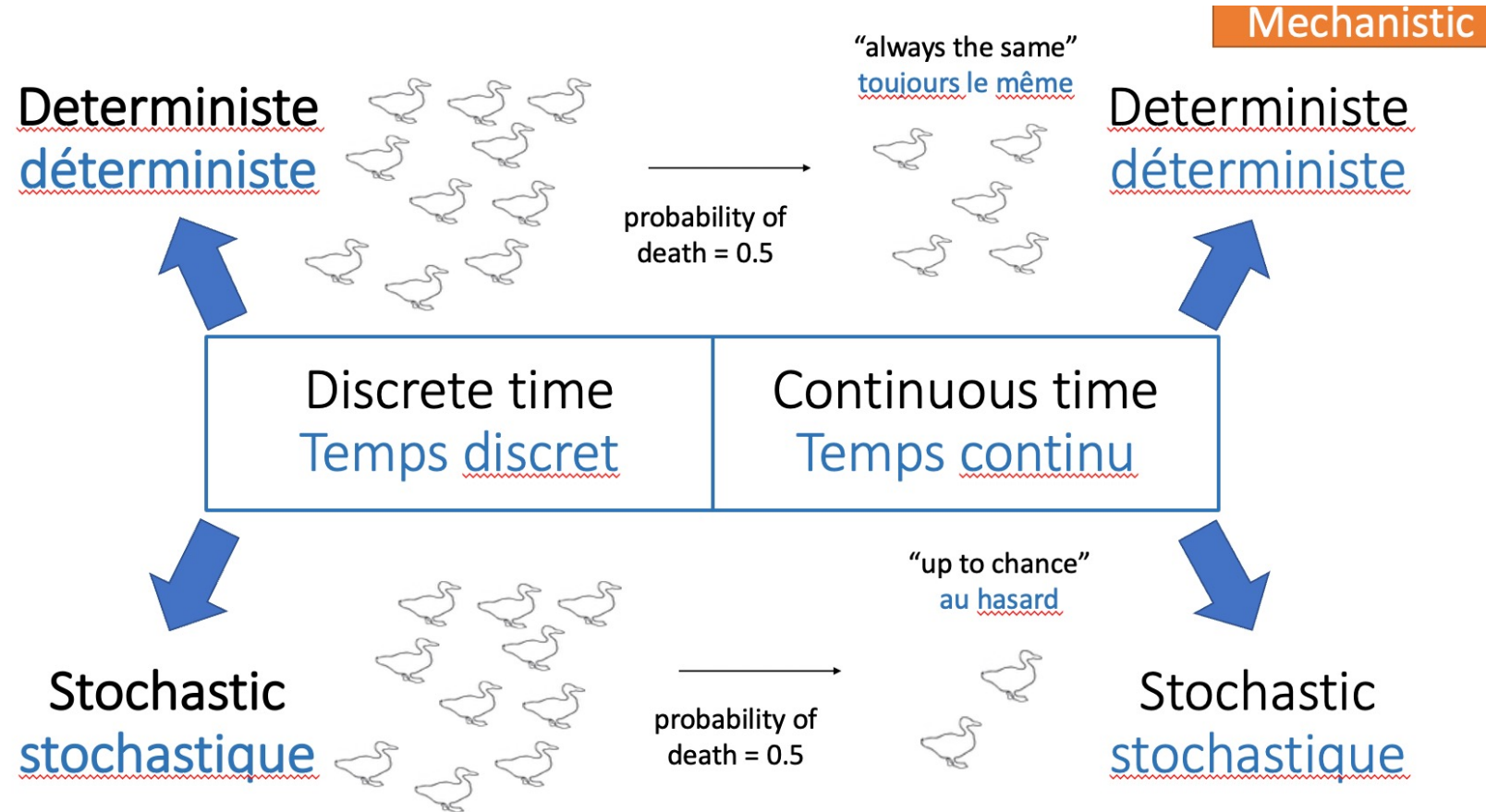


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Refining research questions for modeling
Affiner les questions de recherche pour la
modélisation

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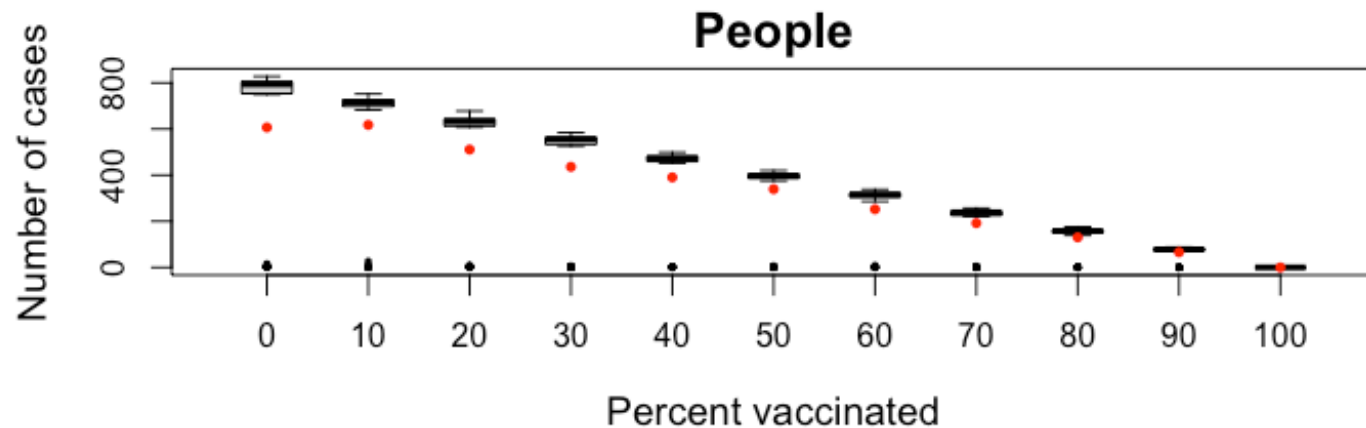
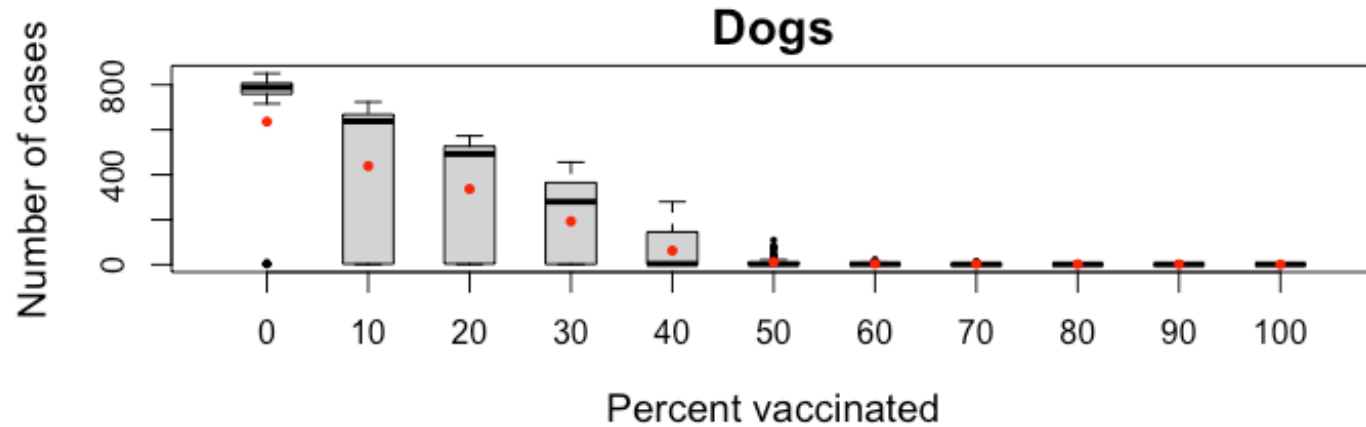
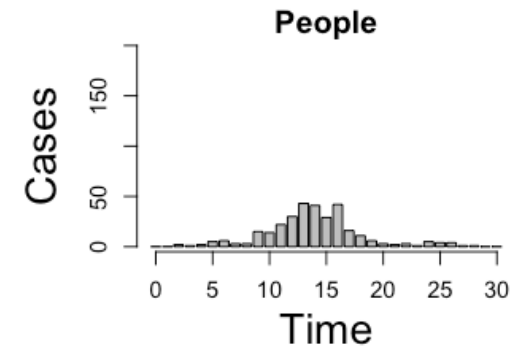
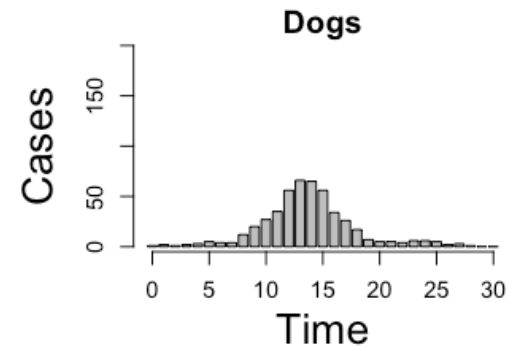
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Dynamical Fever Fièvre dynamique



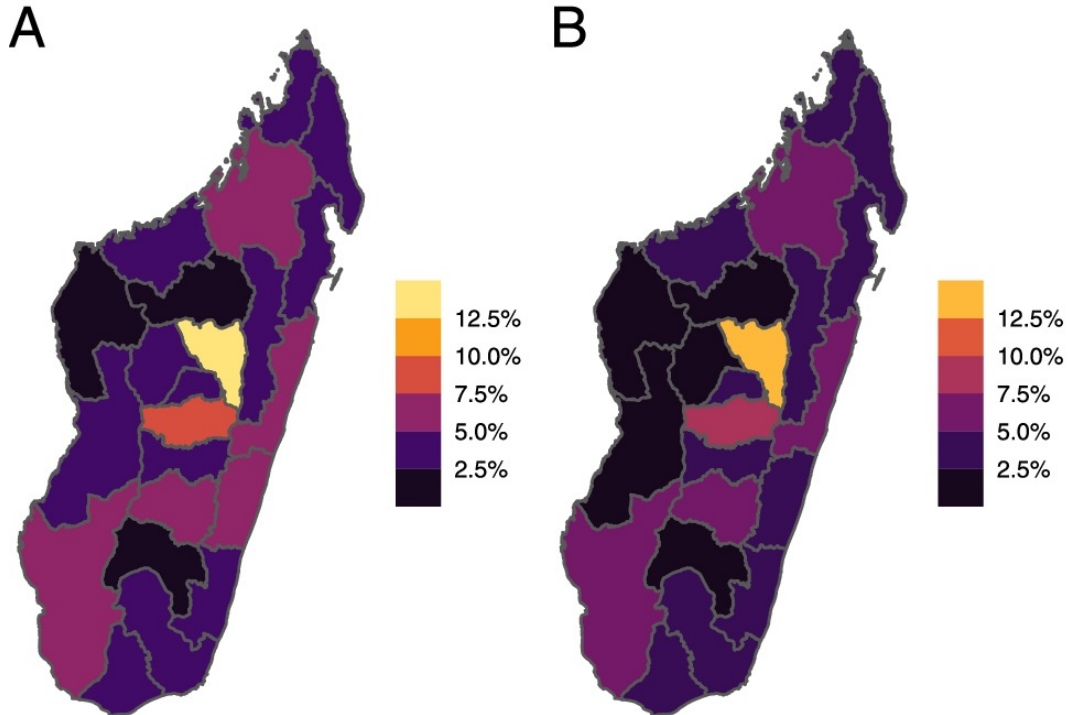
$$P_V = 1 - \frac{1}{R_0}$$

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Reading a Research Paper

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Research | [Open Access](#) | [Published: 12 April 2022](#)

Prioritizing COVID-19 vaccination efforts and dose allocation within Madagascar

[Fidisoa Rasambainarivo](#) , [Tanjona Ramiadantsoa](#), [Antso Raherinandrasana](#), [Santatra Randrianarisoa](#), [Benjamin L. Rice](#), [Michelle V. Evans](#), [Benjamin Roche](#), [Fidiniaina Mamy](#), [Randriatsarafara](#), [Amy Wesolowski](#) & [Jessica C. Metcalf](#)

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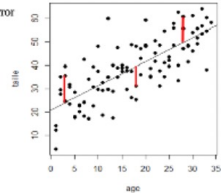
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Generalized linear mixed effect models modèles linéaires généralisés à effets mixtes

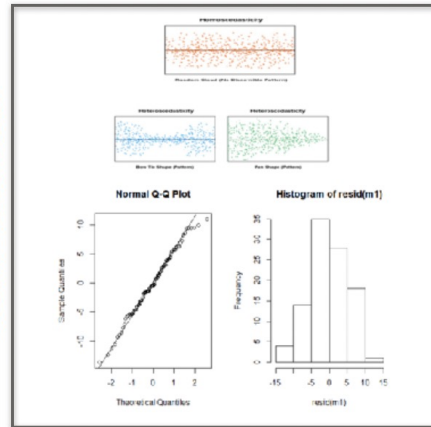
Univariate linear model: simple linear regression

- Quantify the relationship between the response variable and each explanatory variable
- Linear relationship: $y = a + bx + \epsilon$
 - ▶ y : response variable, x : explanatory variable
 - ▶ a : intercept, b : slope, ϵ : Error or residual
- Minimize the error

Taille = 20 + 1.15 * Age(Months) + Error



Model validation



Multilinear model

```
Call:
lm(formula = taille ~ age + sexe + Giparastes + malaria, data = lenur...)

Residuals:
    Min       1Q   Median       3Q      Max
-12.3695  -4.2168   0.0111   3.8715   9.9466

Coefficients:
(Intercept) 24.37448  1.40044 17.405 < 2e-16 ***
age          0.87527  0.05423 15.141 < 2e-16 ***
sexeMale    10.26143  1.04410  9.771 5.11e-16 ***
Giparastes  -0.36176  0.07601 -11.598 < 2e-16 ***
malariaOut  -0.16413  1.04605  -0.108  0.921

Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

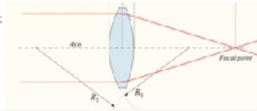
Residual standard error: 5.203 on 95 degrees of freedom
Multiple R-squared:  0.8463, Adjusted R-squared:  0.8399
F-statistic: 130.8 on 4 and 95 DF, p-value: < 2.2e-16
```

Generalized linear model

- Extend the linear model framework by using a linear predictor and a link function
- link function: describe the relationship between the linear combination of the explanatory variables and the mean of the response variable
- R command: `glm(response_variable ~ explanatory_variable, family = family_distribution)`

Most common family function:

Gaussian: identity
Binomial: logit
Poisson: log
Neg binomial: log



Why use GLMMs?

Generalized linear mixed models include both **fixed effects** and **random effects** in order to allow for:

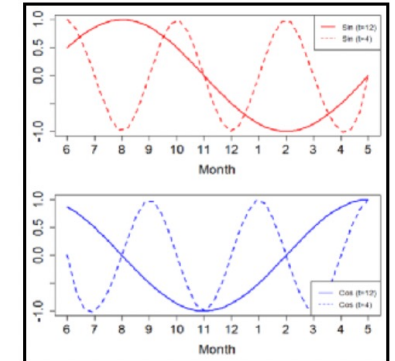
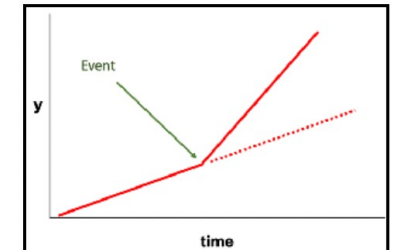
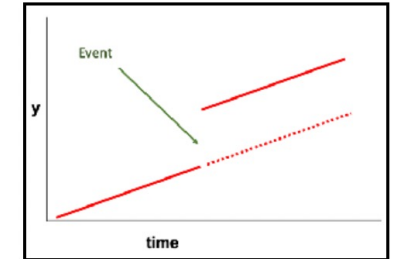
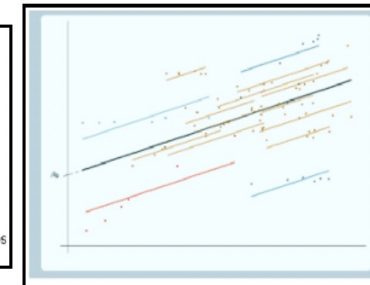
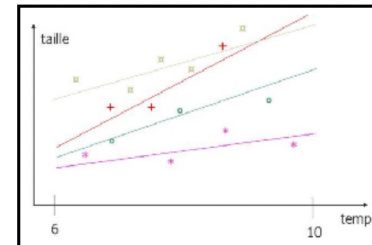
- Repeated measures
- Temporal correlation
- Spatial correlation
- Heterogeneity
- Nested data

$$y = X\beta + Zb + \epsilon_i$$

Fixed Effects

Random Effects

The R function to fit a generalized linear mixed model is `glmer()` which uses the form `fitted.model <- glmer(formula, family = "model family", data = data.frame)`



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Generalized linear mixed effect models modèles linéaires généralisés à effets mixtes

	linear	generalized
fixed	<p>linear regression régression linéaire</p> <p>normal distribution <code>lm()</code></p>	<p>generalized linear regression régression linéaire généralisée</p> <p>non-normal distribution <code>glm()</code></p>
mixed	<p>linear mixed model modèle mixte linéaire</p> <p>normal distribution random effects effets aléatoires <code>lmer()</code></p>	<p>generalized linear mixed model modèle mixte linéaire généralisé</p> <p>non-normal distribution random effects effets aléatoires <code>glmer()</code></p>



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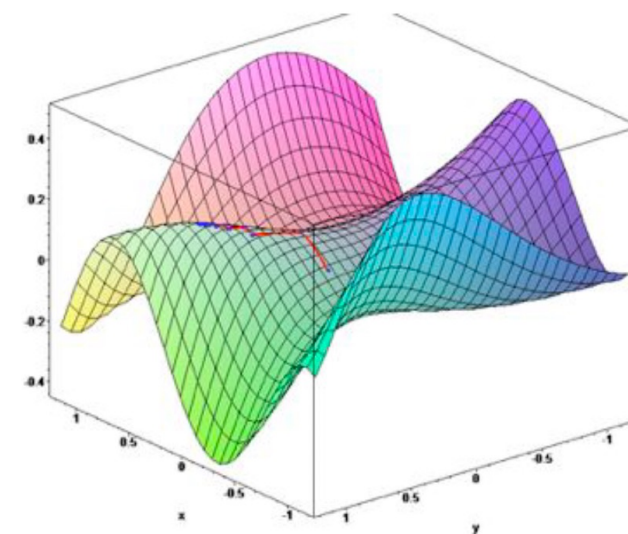
- Model Fitting with Epidemic Cards

Fitting Models to Data

Adapter les modèles aux données

Model Fitting in Science

1. Define your research question (*Définir votre question de recherche*)
2. Formulate a hypothesis (*Formuler une hypothèse*)
3. Collect Data (*Collection des données*)
4. Construct a model that demonstrates your hypothesis (*Construction d'un modèle qui démontre ton hypothèse*)
5. Assess model fit: assuming our model is true, how likely are we to recover the observed data? (*Evaluation du modèle: si le modèle est vrai, quelle est la probabilité qu'on récupère les données observées?*)
6. Optimize parameters behind the model to result in best model fit (*Optimization des paramètres du modèle pour avoir un modèle bien ajusté*)



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Fitting Models to Data

Adapter les modèles aux données

Statistical models are **data-driven**

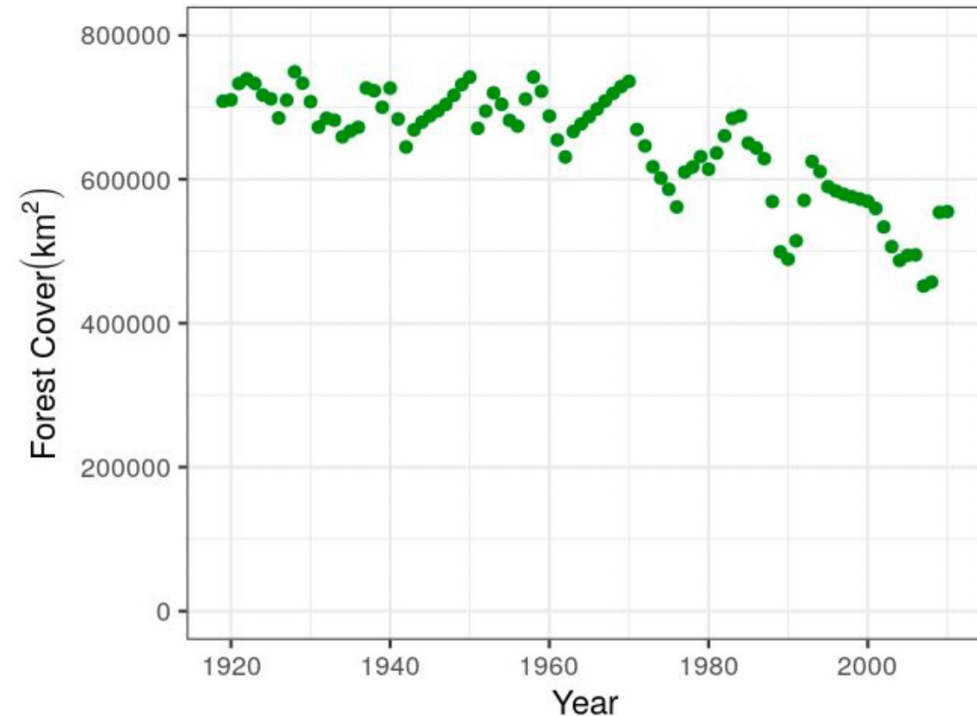
Les modèles statistiques sont basées sur les données

Goal: find patterns and correlations in data

Objectif: révéler des tendances et des corrélations dans les données

What is the trend in Madagascar's forest cover through time?

Quelle est la tendance de la couverture forestière de Madagascar dans le temps?



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Fitting Models to Data

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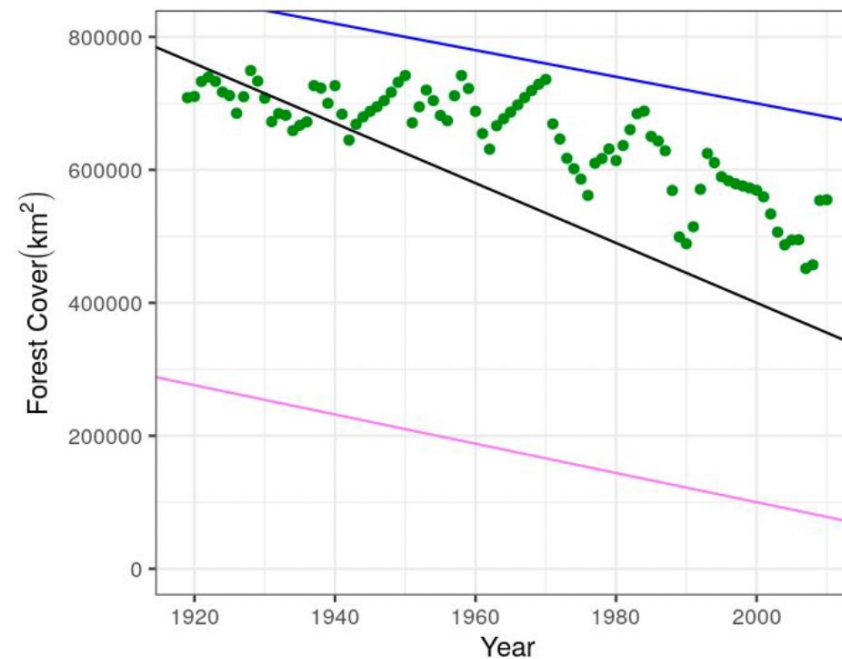
3. Optimize the parameters

Optimisation des paramètres

Optimize slope (m) and intercept (b)

Optimisez la pente (m) and intercept/ordonnée d'origine (b)

Slope (m)	Intercept (b)
-2200	4.5e6
-4500	9.4e6
-2000	4.7e6



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- Model Fitting with Epidemic Cards

Fitting Models to Data

Adapter les modèles aux données

Mechanistic modeling is **process-driven**

La modélisation mécanistique est basée sur les processus

We want to understand what happened,
when it happened, and why it happened

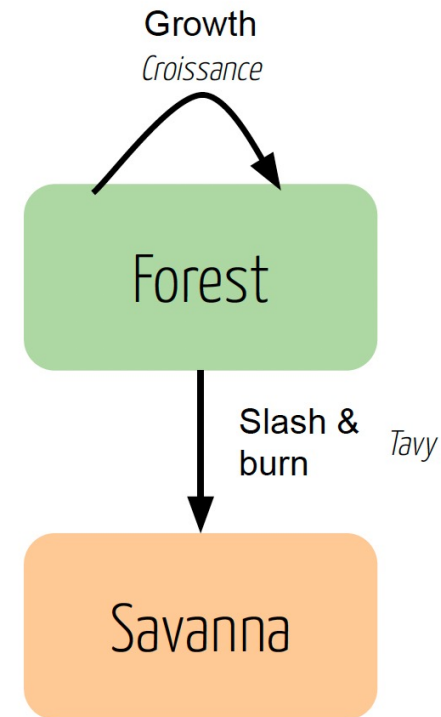
Build a model that uses explicit **processes**
to recover the same outcomes
 (“populations”) as our data

What are the populations in our data?

Quelles sont les populations dans notre système?

What processes are in our data?

Quels sont les processus?



Wednesday: “Fitting Models to Data”

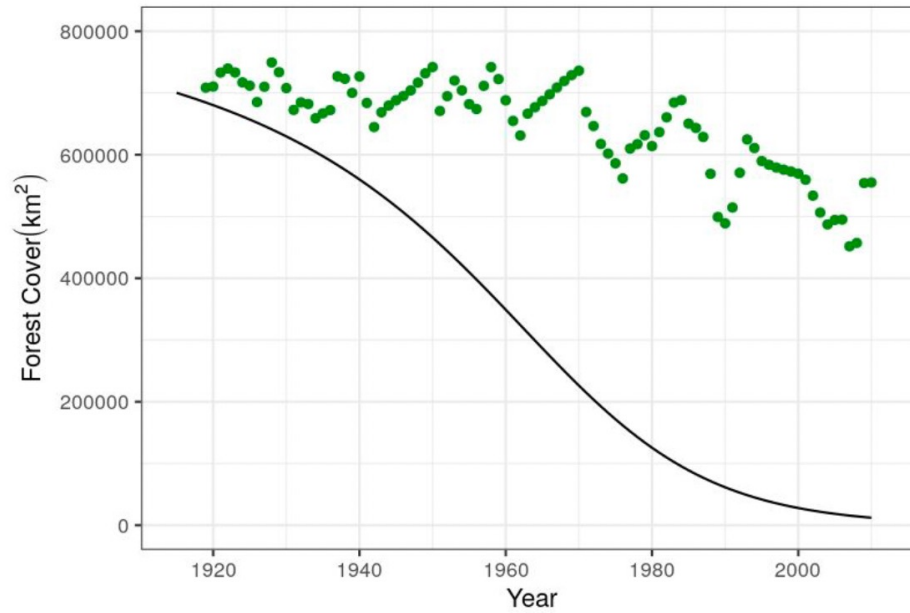
- Model Fitting in Practice – the Basic Concept
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Fitting Models to Data

Adapter les modèles aux données

2. Assess model fit

Évaluation du modèle



Our model predicts forest would decline faster than the data do

Notre modèle prévoit une réduction de la couverture forestière plus rapide que les données

What does this suggest about our guess for the slash and burn rate?

Qu'est-ce que cela suggère à propos de notre estimation du taux du tavy ?

Wednesday: “Fitting Models to Data”

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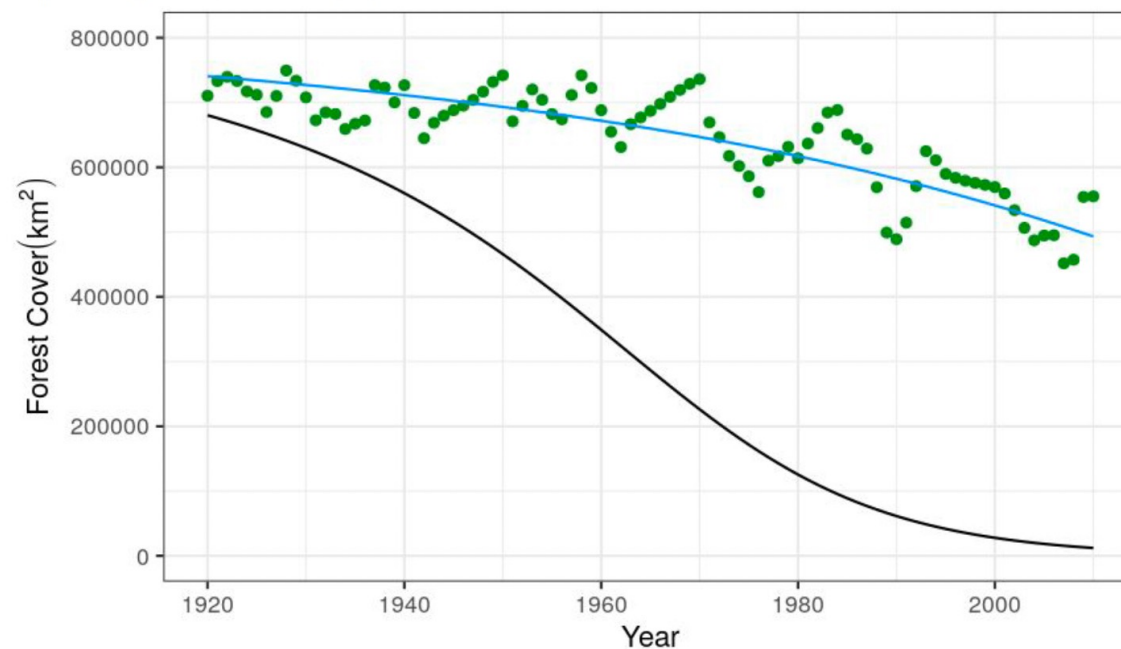
- Model Fitting with Epidemic Cards

Fitting Models to Data

Adapter les modèles aux données

Does this optimal value result in a model that better matches the data?

Est-ce que cette valeur optimale (en bleu) mieux explique les données?

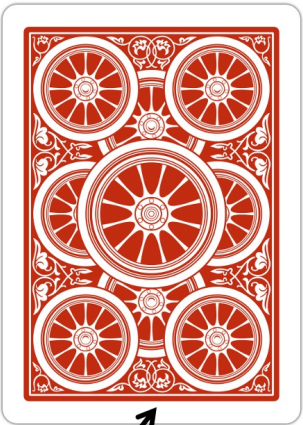


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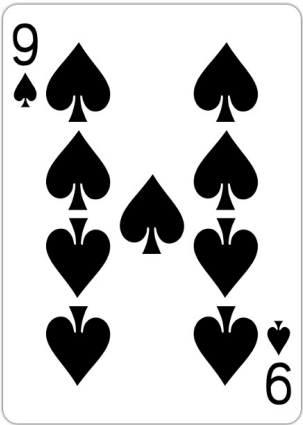
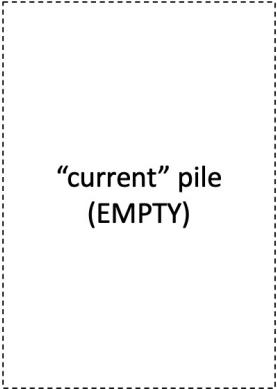
Epidemic Cards
Cartes Épidémie

The Set-Up



The population pile must ALWAYS maintain exactly 26 cards.

What is the equivalent assumption in defining a model world to represent an epidemic?



Round One

Timestep	Infecteds	Susceptibles	Trial	R0
1	1	25	1	2
2			1	2
3			1	2
4			1	2
5			1	2
6			1	2
7			1	2
8			1	2
9			1	2
10			1	2

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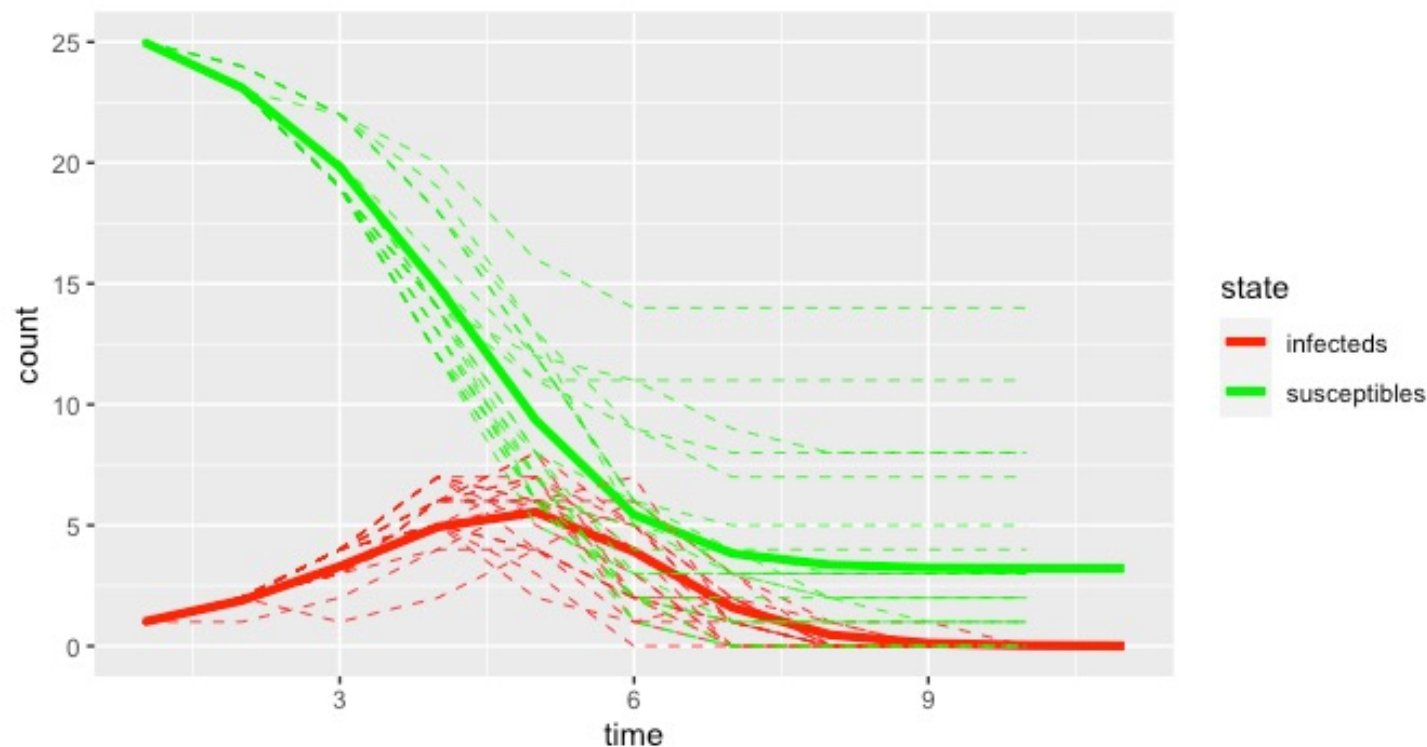
Epidemic Cards Cartes Épidémie

```
## (12) Now, write a for-loop that iterates your discrete time model across the full length of the time series  
## Essentially, write the R language that says the following:
```

```
##      for (all variables, t, in the length of our time vector){  
##          run my discrete time model (hint, use the equations from above, lines 88 and 89)  
##      }
```

```
## S[t+1] = S[t] - (R0 * S[t]/N)*I[t]  
## I[t+1] = (R0 * S[t]/N)*I[t]
```

```
for(t in 1:length(time)){  
  model.S[t+1] = model.S[t] - (R0 * model.S[t]/N)*model.I[t]  
  model.I[t+1] = (R0 * model.S[t]/N)*model.I[t]  
}
```



Thursday: “Refining Your Work + Bells and Whistle”

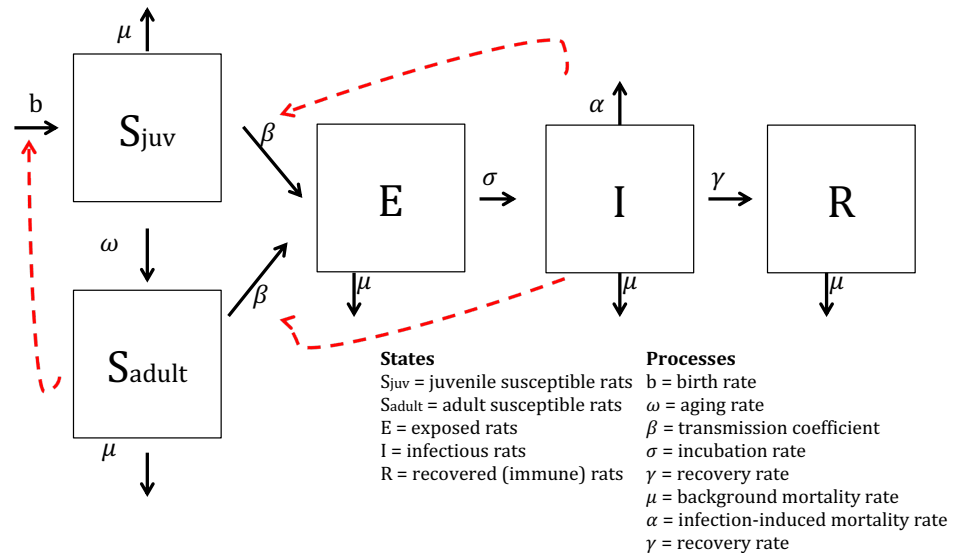
- Model Telephone
- Model Selection and Comparison • Tutorial
- Introduction to Phylogenetic Modeling • Tutorial
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Model Telephone Modèle de téléphone



Example:

Can the Malagasy black rat (*Rattus rattus*) population independently maintain transmission of the plague bacterium, *Yersinia pestis*?



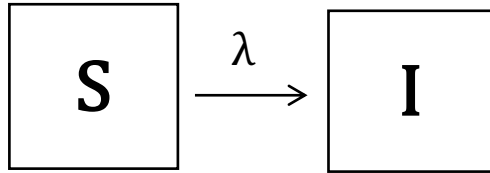
Example Model Description

Can the Malagasy black rat (*Rattus rattus*) population independently maintain transmission of the bacterium, *Yersinia pestis*, responsible for human plague?

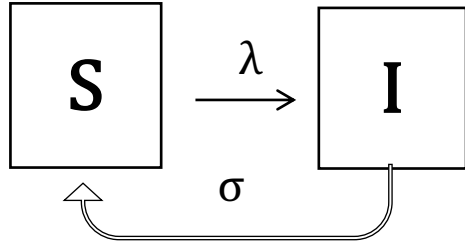
Susceptible juvenile rats enter the population through birth, at rate b , which is influenced by the proportion of uninfected (susceptible) adult rats in the population at a given time. Juvenile rats age into the adult class, on average $1/\omega$ time units after they are born. Both juvenile and adult susceptible rats can be infected by contact with infectious rats of any age, based on a force of infection proportional to the prevalence of infectious rats in the population. Once infected, rats enter the exposed class. The incubation period is $1/\sigma$ time units (on average), after which the animals develop clinical plague, which is equivalent to transitioning from the exposed class to the infectious class. A subset of rats recover from plague to become immune, based on rate γ . All rats in the population experience background mortality with hazard μ , and infectious rats experience an additional disease-induced hazard of mortality, α .

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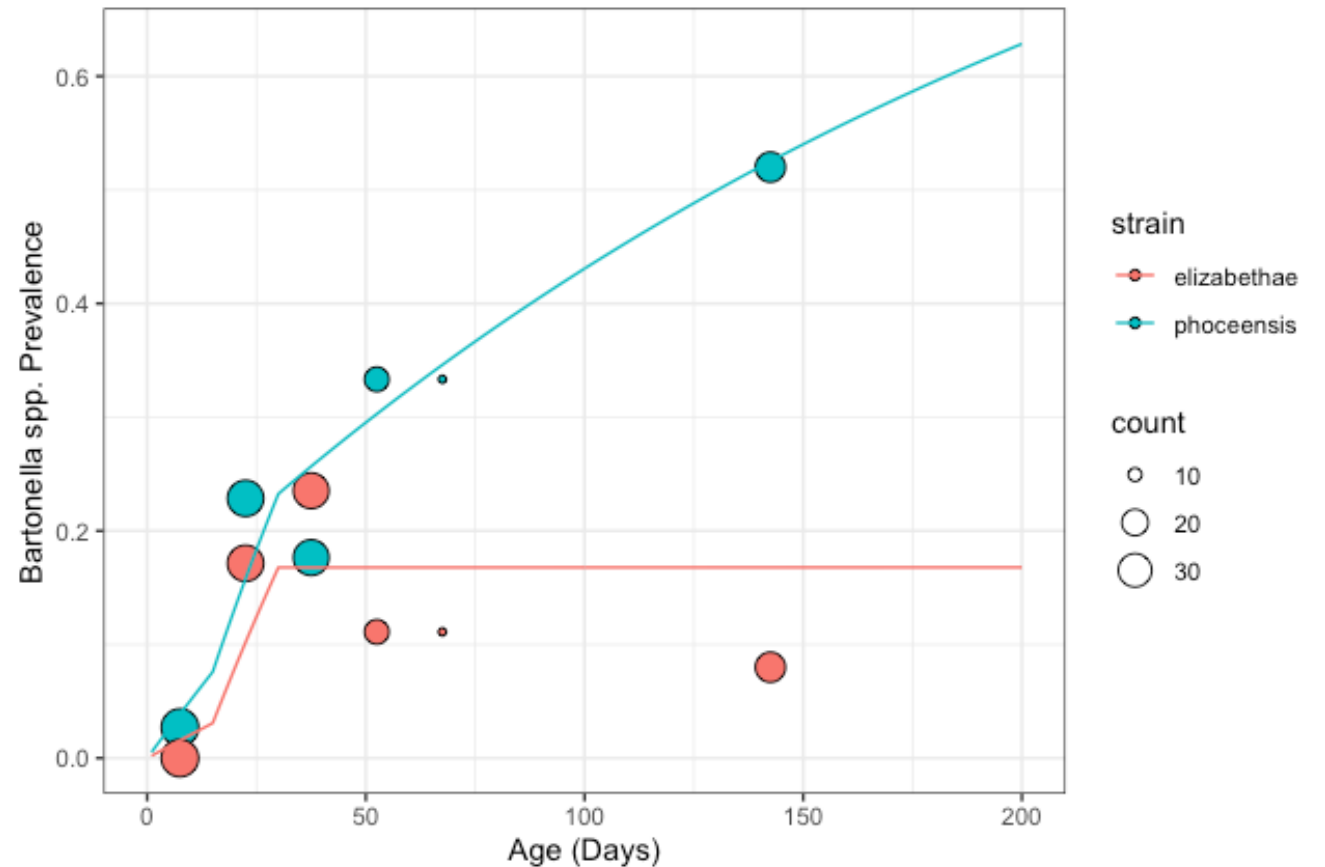


$$\frac{dI(a)}{da} = \lambda(a)(1 - I(a))$$



$$\frac{dI(a)}{da} = \lambda(a)(1 - I(a)) - \sigma I(a)$$

Model Selection and Comparison Sélection et comparaison de modèles



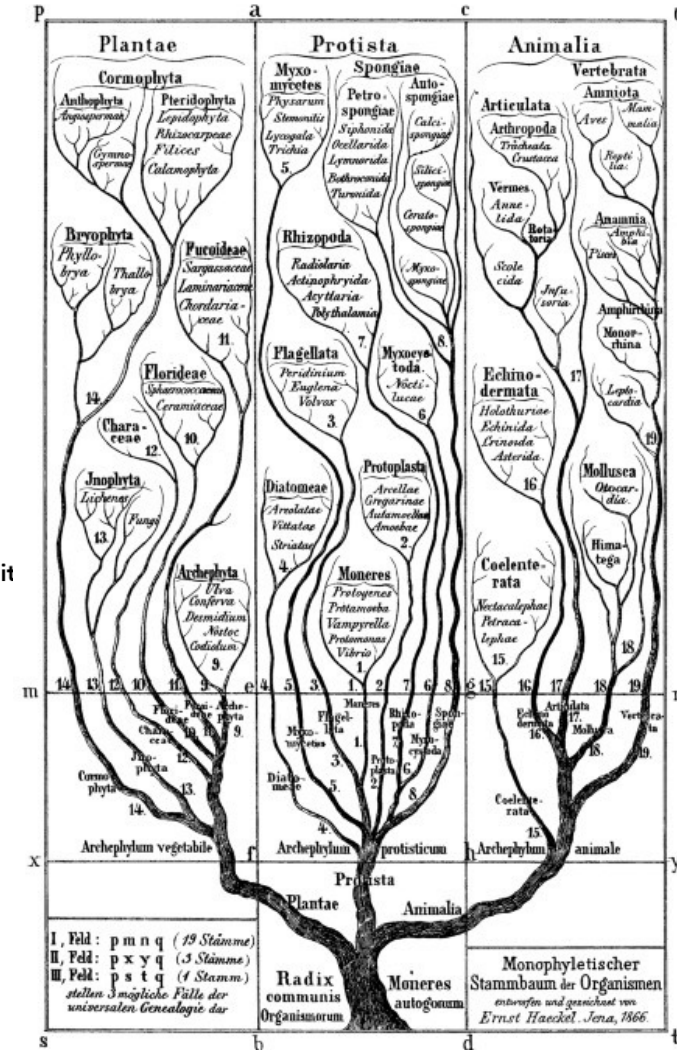
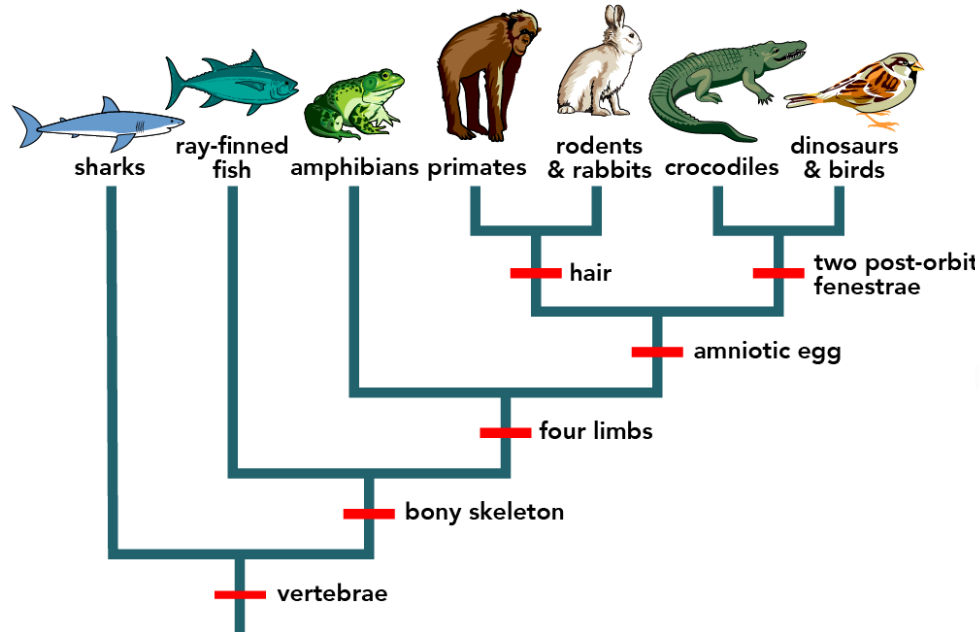
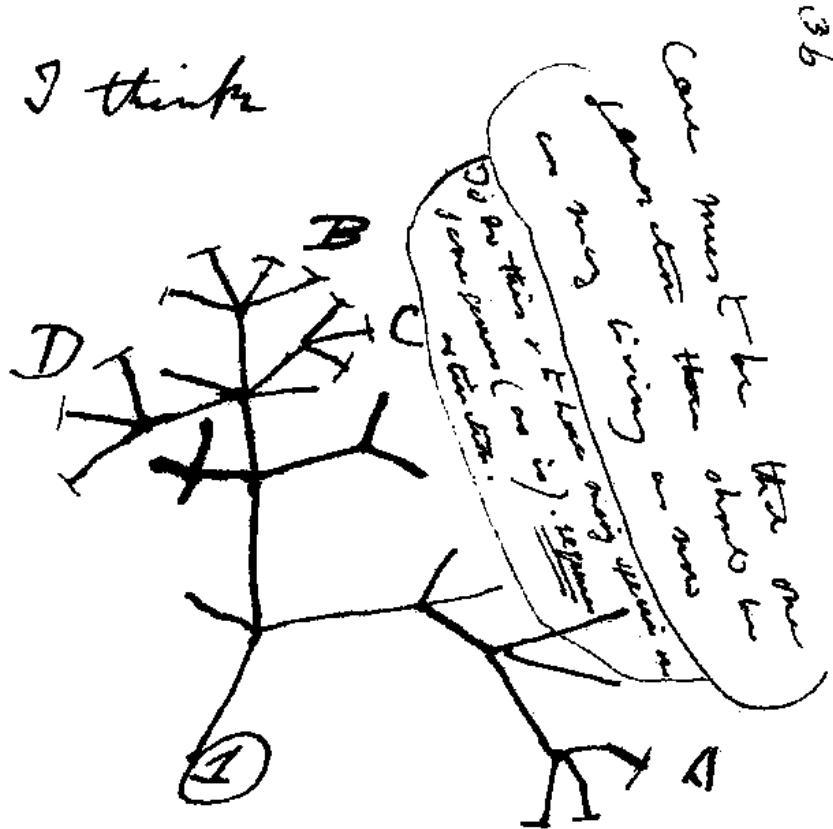
Thursday: “Refining Your Work + Bells and Whistle”

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Intro to Phylogenetic Modeling

Introduction à la modélisation phylogénétique



Hossfeld and Levit, Nature, 2016

Thursday: "Refining Your Work + Bells and Whistle"

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Intro to Network Modeling

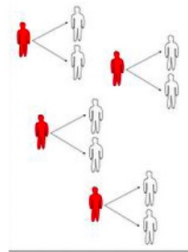
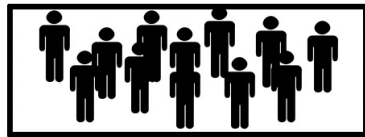
Introduction à la modélisation de réseau

Why do we use networks in epidemiology?

Pourquoi utiliser les réseaux dans épidémiologie?

Classic epidemiological models assume full mixing (everyone can contact everyone)

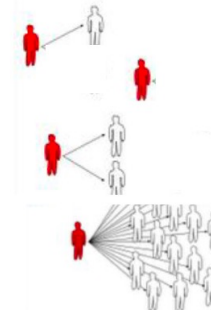
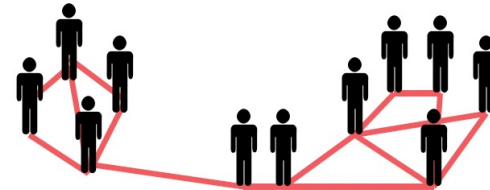
Les modèles épidémiologiques classiques supposent un mélange complet (tout le monde peut contacter tout le monde)



$$R_0 = 2$$

Reality is better represented by a network

La réalité est mieux représentée par un réseau



Average $R_0 = 2$,

But more realistic heterogeneity

Friday: R Bootcamp

- Intro to R Studio
- Exploring and Visualizing Data in R
- For-loops, Functions, and If-Else Statements

Thursday: "Bells and Whistle"

- Introduction to Phylogenetic Modeling • Tutorial
- Introduction to Spatial Visualization and Plotting • Tutorial
- Modeling Insights from the Metapopulation Game • Tutorial
- Introduction to Network Modeling • Tutorial
- Research Snapshots

- Programming
- Data
- Models
- Research Development

Saturday: Travel

Sunday: "Getting Started with the Basics"

- Data and Models
- Student introductions & presentations
- Linear regression • Tutorial
- Formulating research questions

Monday: "Using Models with Data"

- Model-Guided Study Design
- Study design tutorial
- Intro to Compartmental Models & Differential Equations
- Building mechanistic models in R
- Refining research questions for modeling
- Defining a model world

Wednesday: "Refining Your Work"

- Model Fitting in Practice – the Basic Concept
- Epidemic Cards
- Model Fitting with Epidemic Cards
- Model Selection and Comparison
- Model Selection Tutorial
- Model Telephone

Tuesday: "Applying Simple Models"

- Dynamical Fever
- Introduction to Mixed Modeling
- Reading a Research Paper

Friday: "Putting it All in Perspective"

- The Life Cycle of a Modeling Project
- C4C Student Presentations

Saturday: Travel

January: "Sharing Your Work"

- Final student presentations

Any aminareo ny baolina



E²M²: Ecological & Epidemiological Modeling in Madagascar

