

*Tanjona Ramiadantsoa*

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# A history of your new skills

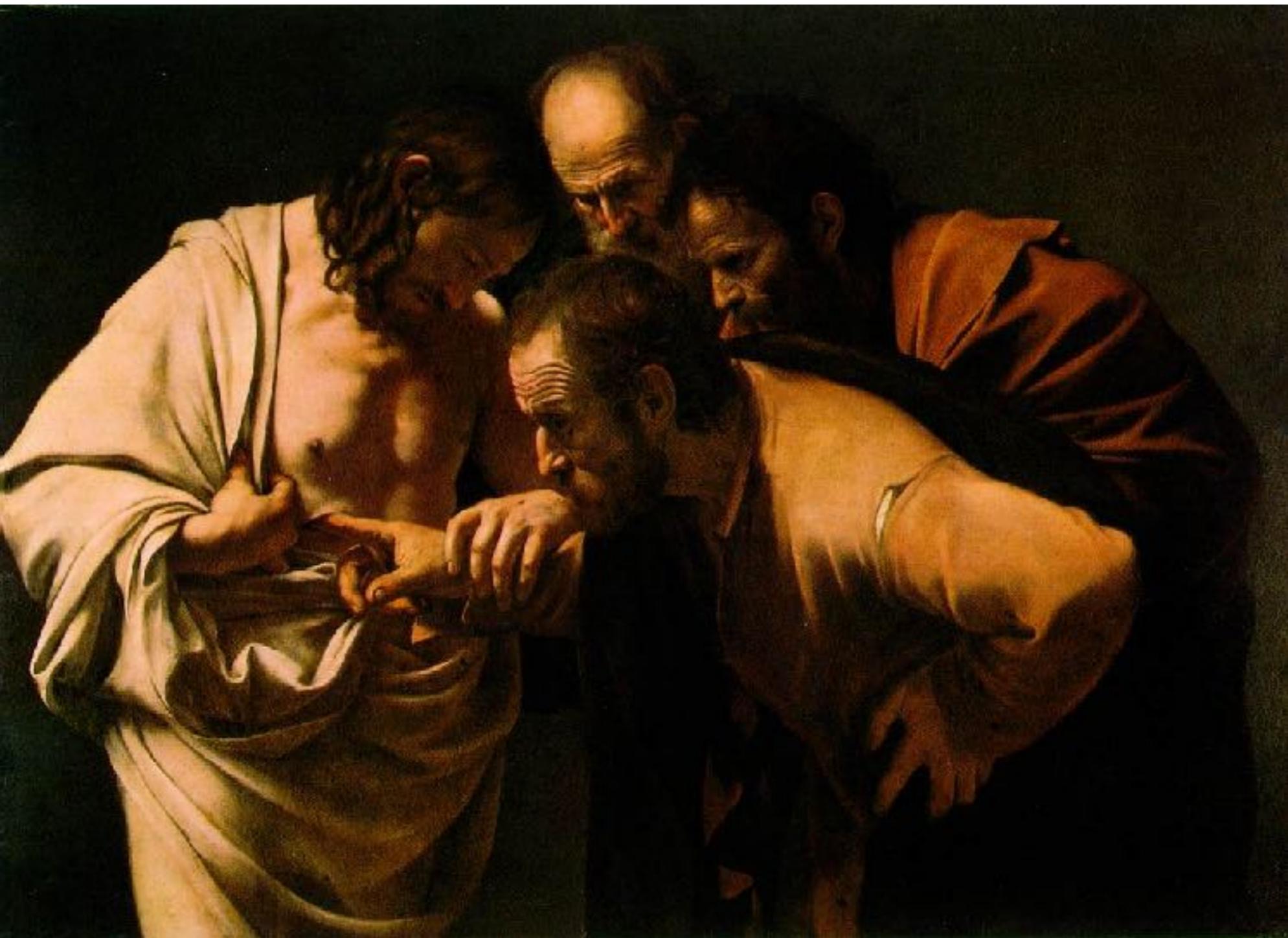
E2M2 2020  
Valbio Ranomafana  
January 10th, 2020

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# What is a scientist?

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## Steps in a modeling project

1. *Development of the study concept and question*
2. *Literature review*
3. *Data collection*
4. *Construction of model framework*
5. *Model analyses and selection*
6. *Model validation*
7. *Manuscript writing and submission*



### Literature review

- Who has tried to answer this before and how did they do it?
  - Empirical studies
  - Modeling studies
- What are these studies short-comings?
- Are there already parameter estimates or data sets to help you answer your question?



### Manuscript writing and submission

- What are the main results that provide the answer to my question?
  - 1 to 3 graphs
  - 1 to 3 tables
- What is the journal that best fits my study?
  - Scope, audience, impact factor, math focus
- How do I present my manuscript?
  - Introduction: set the stage to your question
  - Methodology: describe explicitly all steps for replicability
  - Results: clear and concise
  - Discussion: explain how your study improves previous knowledge



## Steps in a modeling project

### 1. Development of the study concept and question

### 2. Literature review

To help you prepare, try to answer the next few questions after reading:

- What are the context and key points of the paper?
- Identify the problem that inspired the authors to write the article.
- Do you agree with the methods, the results, and the discussion?
- Where do the authors use models to address their questions? What type of models can you identify?
- What do you like and what do you dislike about the paper?
- What did you not understand?
- What have you learned from the paper?
- And come up with three questions of your own.

ii Literature review

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  - Modeling studies
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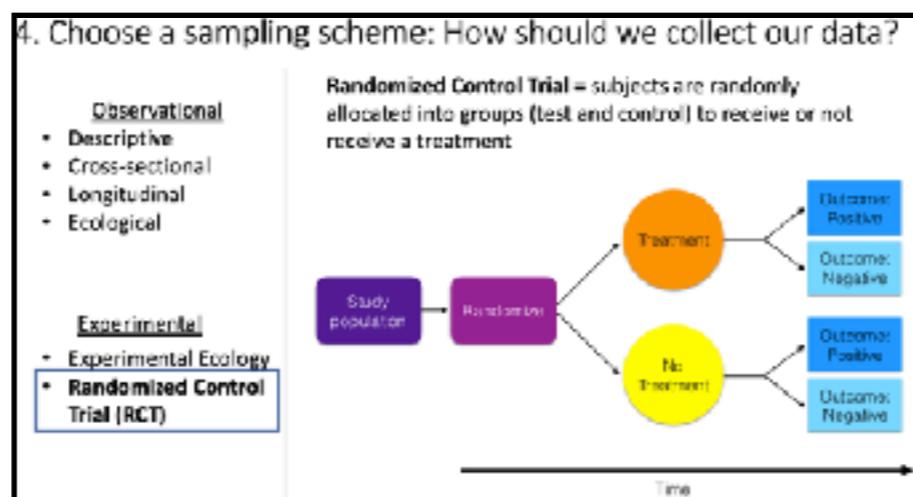


session  
the answer to my question?

- What is the journal that best fits my study?
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# Study Design

1. Define the **research question**: What are we trying to answer?
2. Define the **sample type**: What data do we need to answer our question?
3. Identify a **system**: Where can we collect our data?
4. Choose a **sampling scheme**: How should we collect our data?
5. Acknowledge **limitations**: What can we actually infer from our data?
6. Outline a **data organization plan**: How should we organize our data?
7. Be **flexible**: How can we prepare for potential/unanticipated challenges?



# Take homes

- ❖ Research question and hypothesis are more important than models
- ❖ Any data needs context: the X and Y should be clear
- ❖ Models are rigorous tools to assess how the data support the claim
  - ❖ There are figuratively an infinite number of models
  - ❖ Statistical model works with question starting with **what**
  - ❖ Mechanistic model **generates data** and works with question starting with **how**

# Two broad classes of models

Statistical



Correlative

Mechanistic



Causative

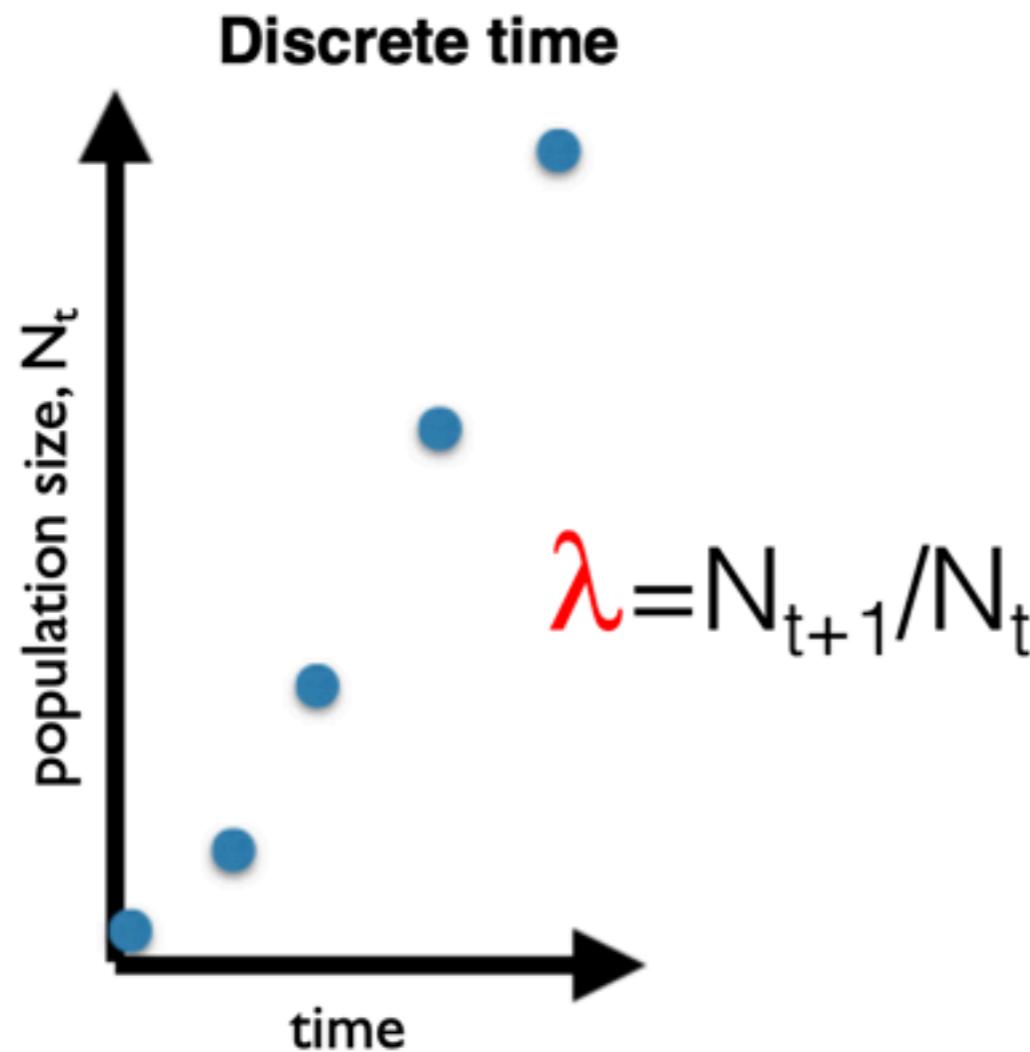
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# Mechanistic (mathematical) model

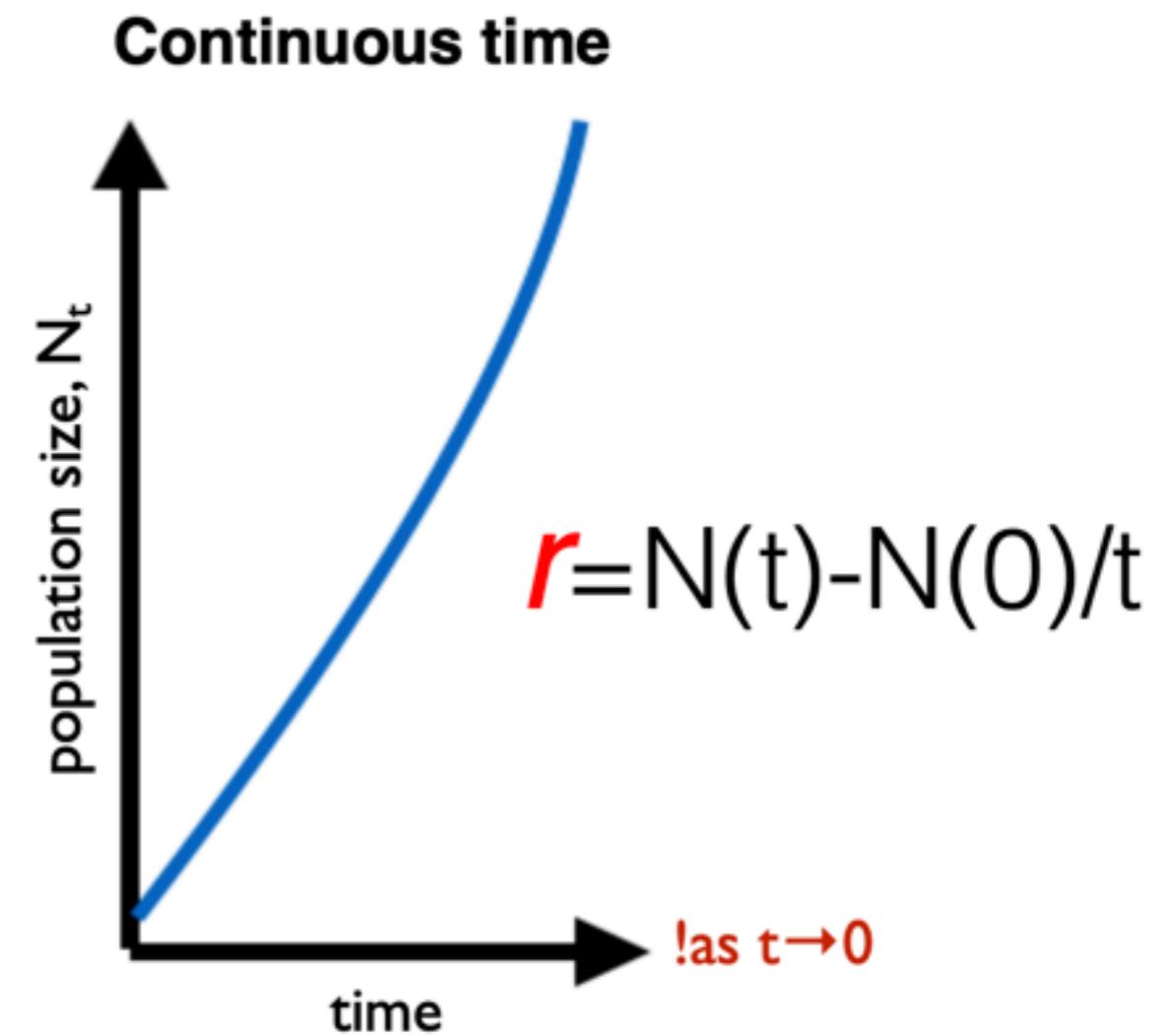
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- ❖ **Process-driven** with question starting with **How**
- ❖ You generate data with simulation

# Discrete vs. continuous



$$N_1 = \lambda N_0$$
$$N_2 = \lambda[\lambda N_0] = \lambda^2 N_0$$
$$N_3 = \lambda^3 N_0$$
$$N_t = \lambda^t N_0$$



$$dN(t)/dt = rN(t)$$

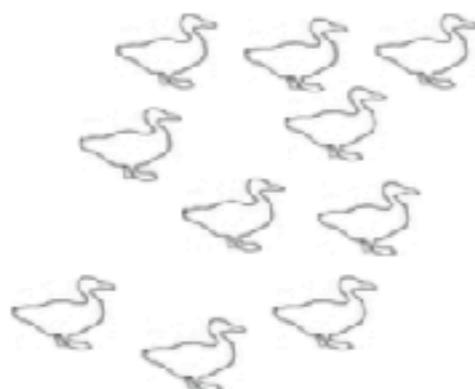
$$N(t) = N(0)e^{rt}$$

# Deterministic vs. stochastic

## The basic population model



starting population

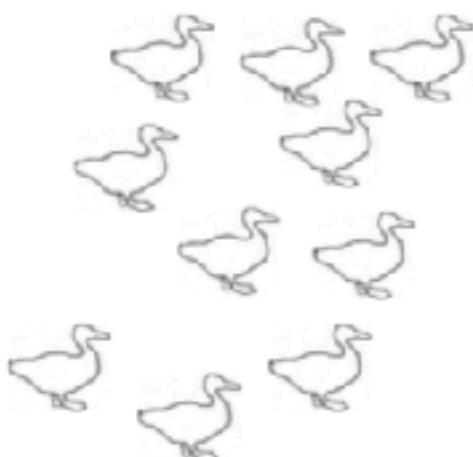


probability of  
death = 0.5

if deterministic    "always the same"



starting population

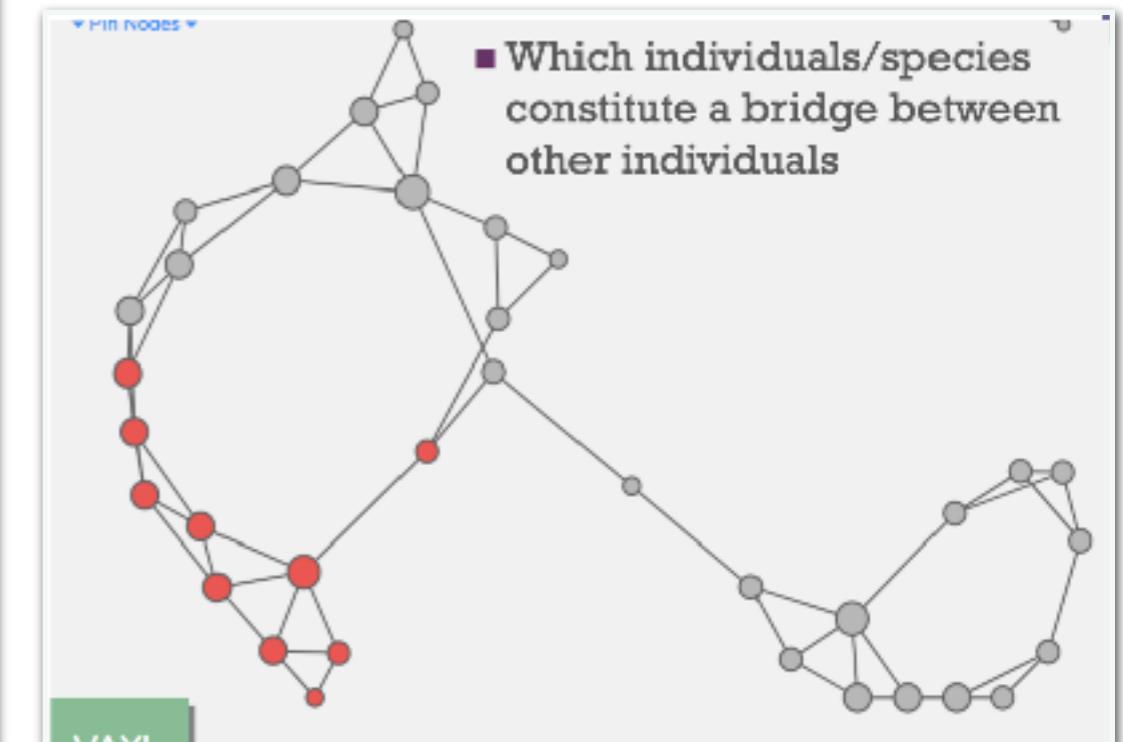
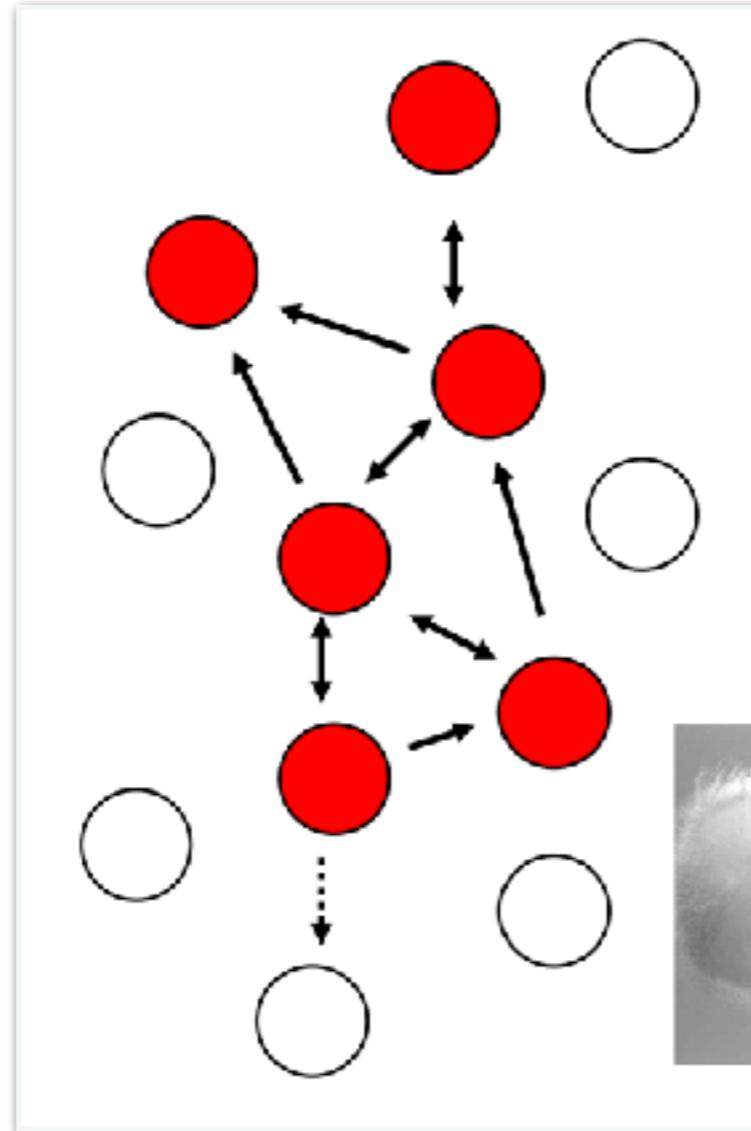
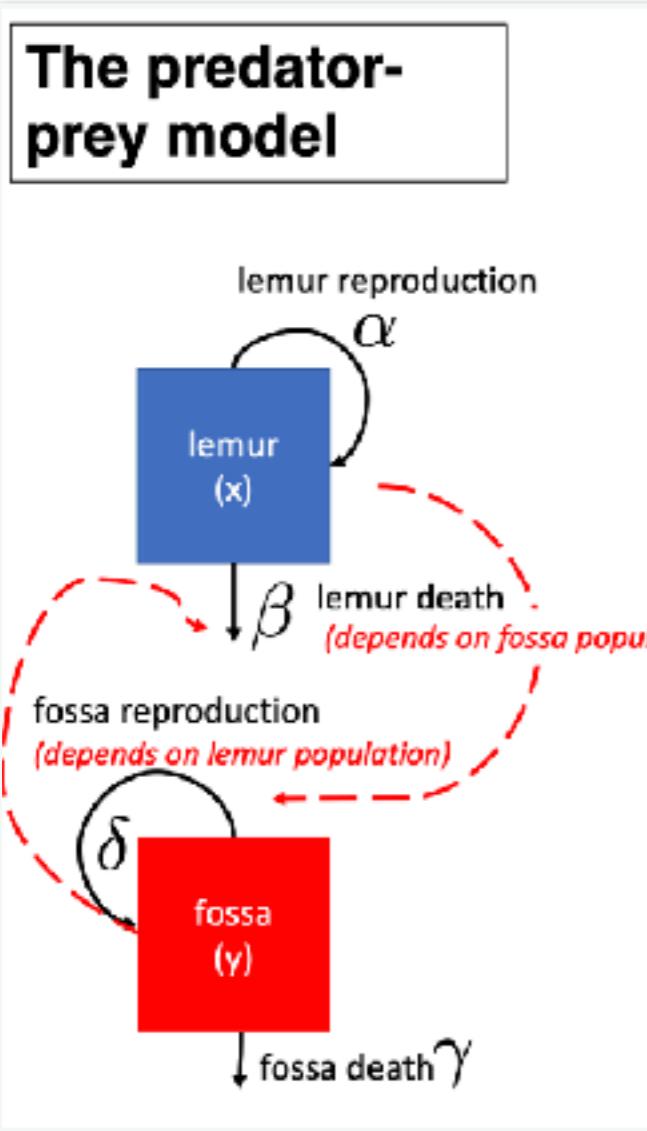


probability of  
death = 0.5

if stochastic?    "up to chance"

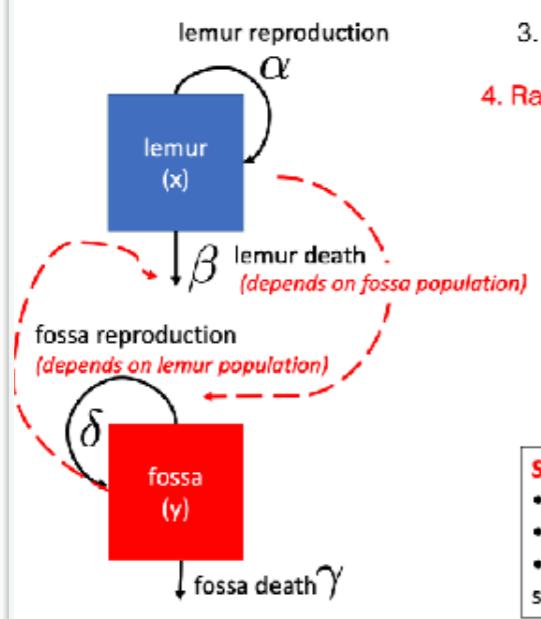


# Non-spatial vs. spatial vs. network



# Compartment models

## The predator-prey model



### Compartmental models (Mechanistic Models)

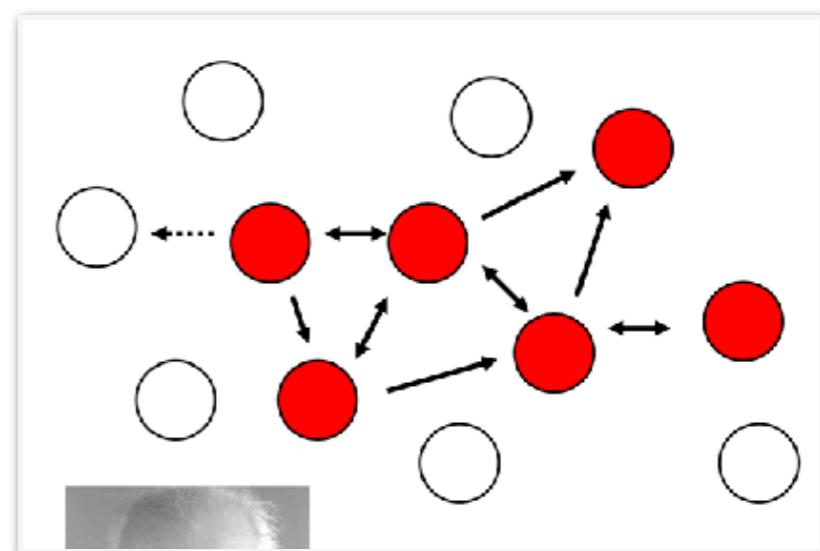
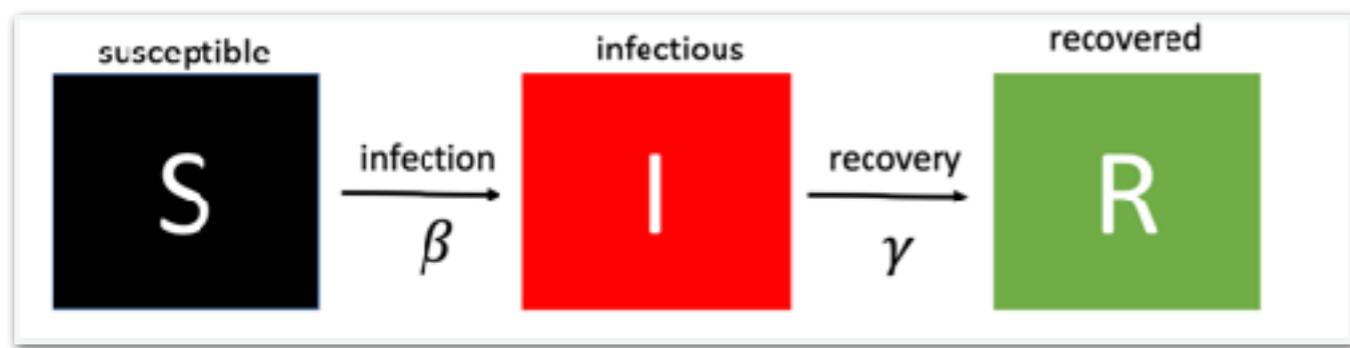
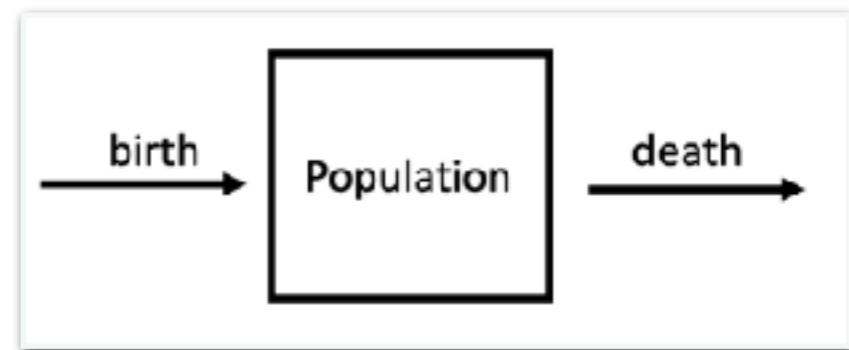
1. Populations are divided into compartments
2. Individuals within a compartment are homogeneously mixed
3. Compartments and transition rates are determined by biological systems
4. Rates of transferring between compartments are expressed mathematically

$$\frac{dx}{dt} = x(\alpha - \beta y)$$

$$\frac{dy}{dt} = y(\delta x - \gamma)$$

#### SOME ASSUMPTIONS

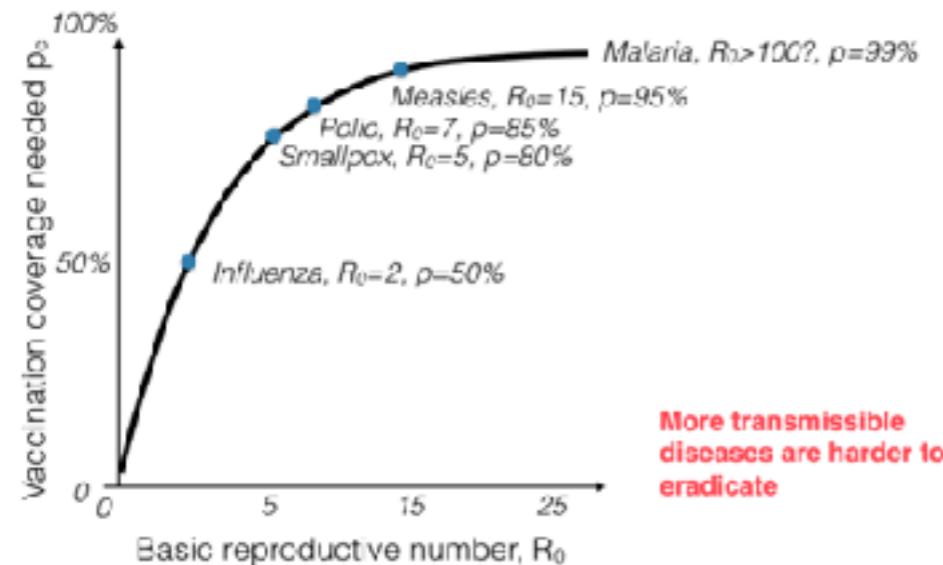
- the **lemur** has an unlimited food supply
- the **lemur** only dies from being eaten by fossa
- the **fossa** is totally dependent on a single prey species (the lemur) as its only food supply



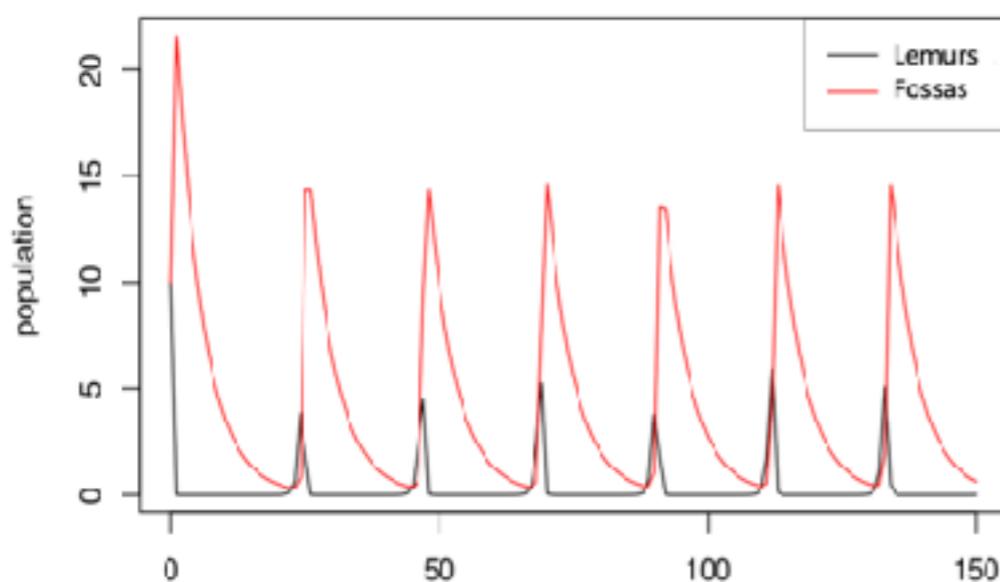
# Insights

## The SIR model : eradication

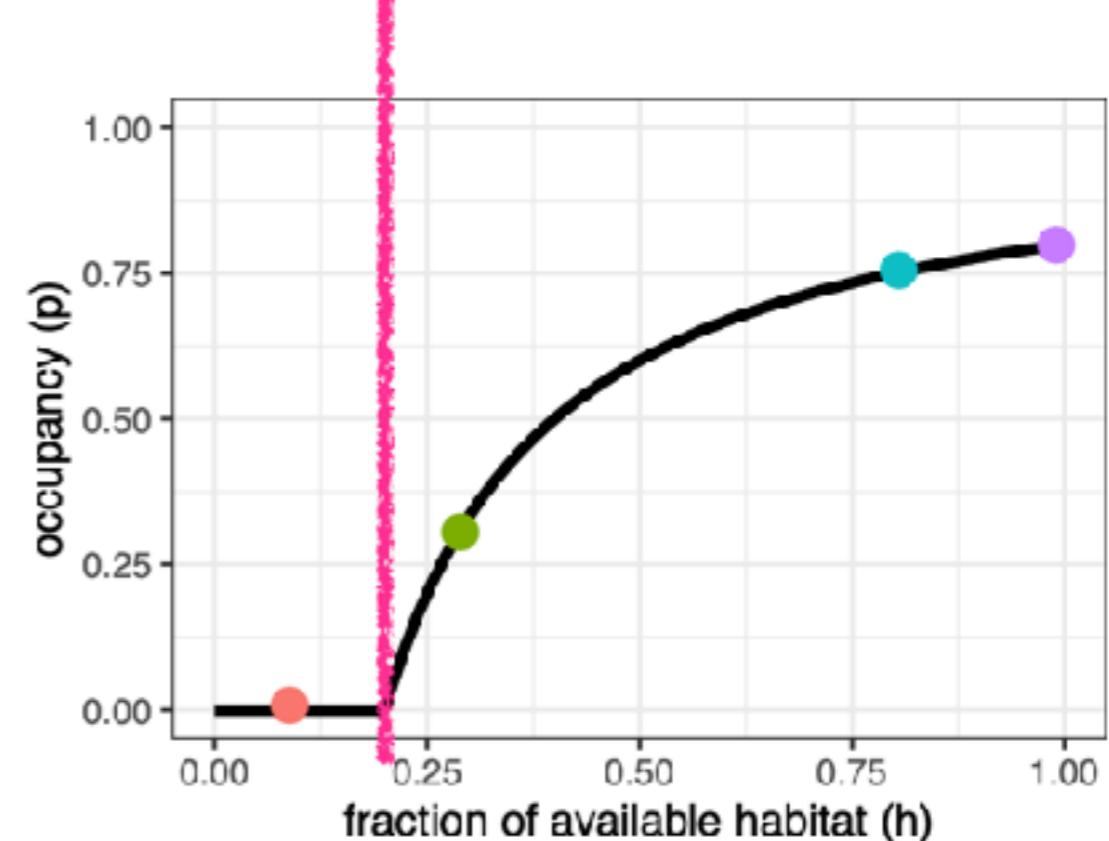
$$p_c = 1 - \frac{1}{R_0}$$



## The predator-prey model



Extinction threshold = the root



Extinction threshold:  $h = \frac{e}{c}$

# Two broad classes of models

Statistical



Correlative

Mechanistic



Causative

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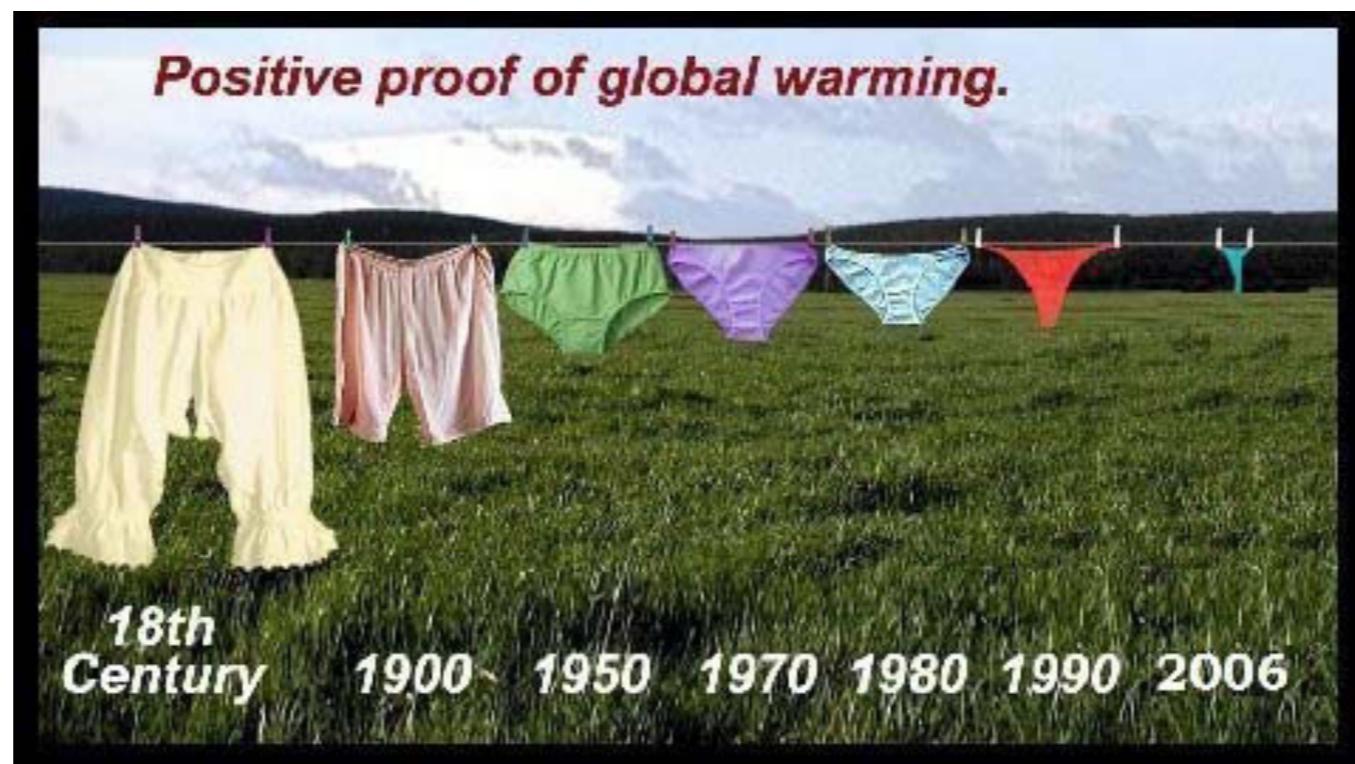
# Statistical model

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- ❖ **Data-driven** with question starting with **What**
- ❖ Test patterns in data using predefined functions

# Statistical model

- ❖ Data-driven with question starting with What
- ❖ Test patterns in data

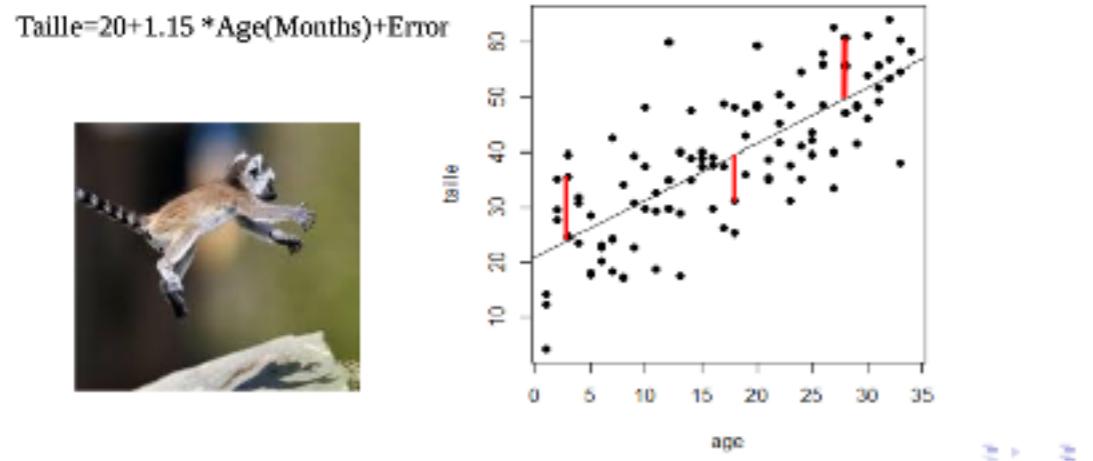


POSITIVE PROOF OF GLOBAL WARMING

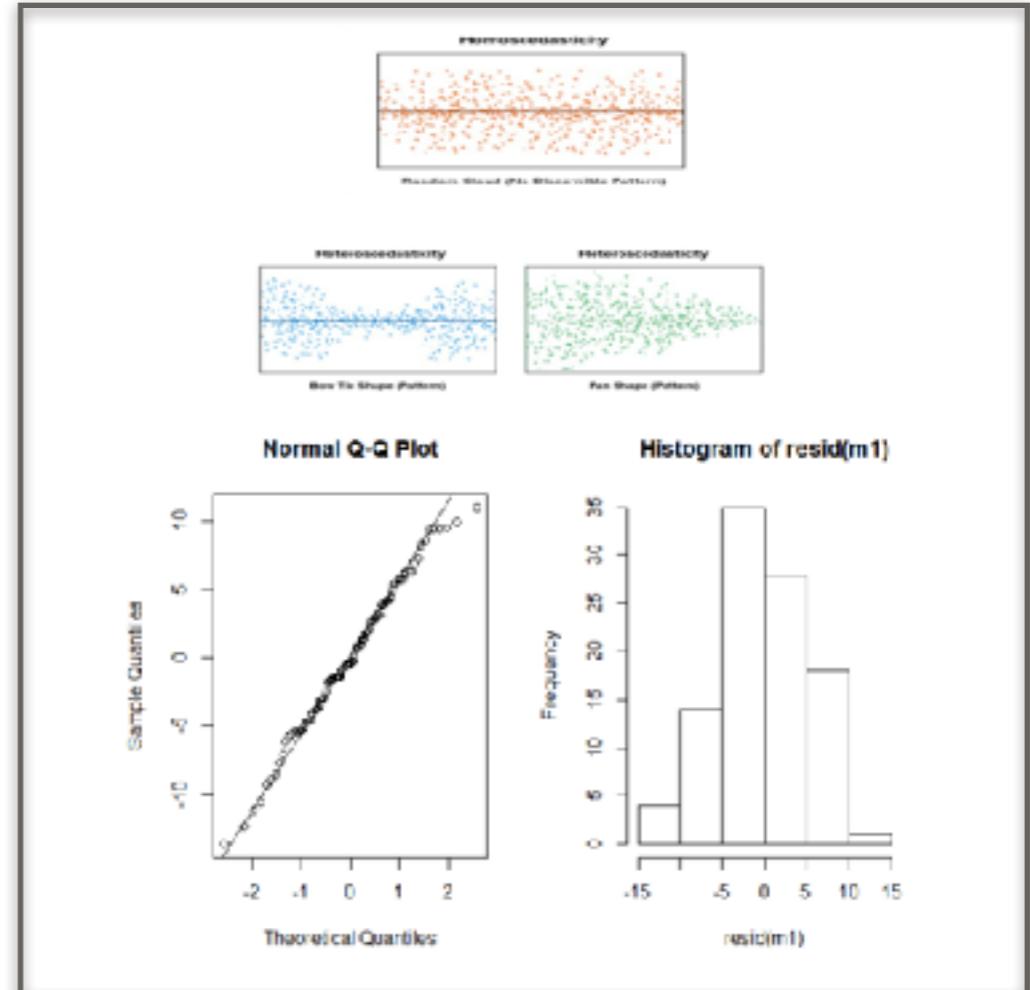
Correlation does not imply causation

## 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,  $\epsilon$ : Error or residual
- Minimize the error



## Model validation



## Multilinear model

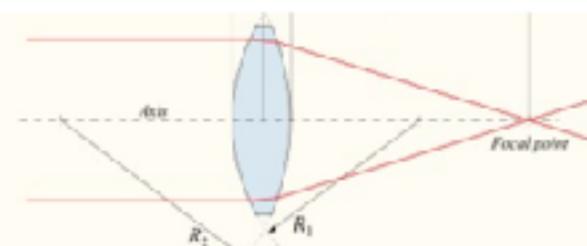
```
Call:  
lm(formula = taille ~ age + sexe + GIparasites + malaria, data = lemur.)  
  
Residuals:  
    Min      1Q  Median      3Q     Max  
-12.3695 -4.2168  0.0111  3.8716  9.9466  
  
Coefficients:  
            Estimate Std. Error t value Pr(>|t|)  
(Intercept) 24.37448   1.40044 17.405 < 2e-16 ***  
age          0.87527   0.05423 15.141 < 2e-16 ***  
sexeMale    10.28143   1.04410  9.771 5.11e-16 ***  
GIparasites -0.38178   0.02681 -11.598 < 2e-16 ***  
malariaOui  -0.18413   1.04683 -0.180    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:

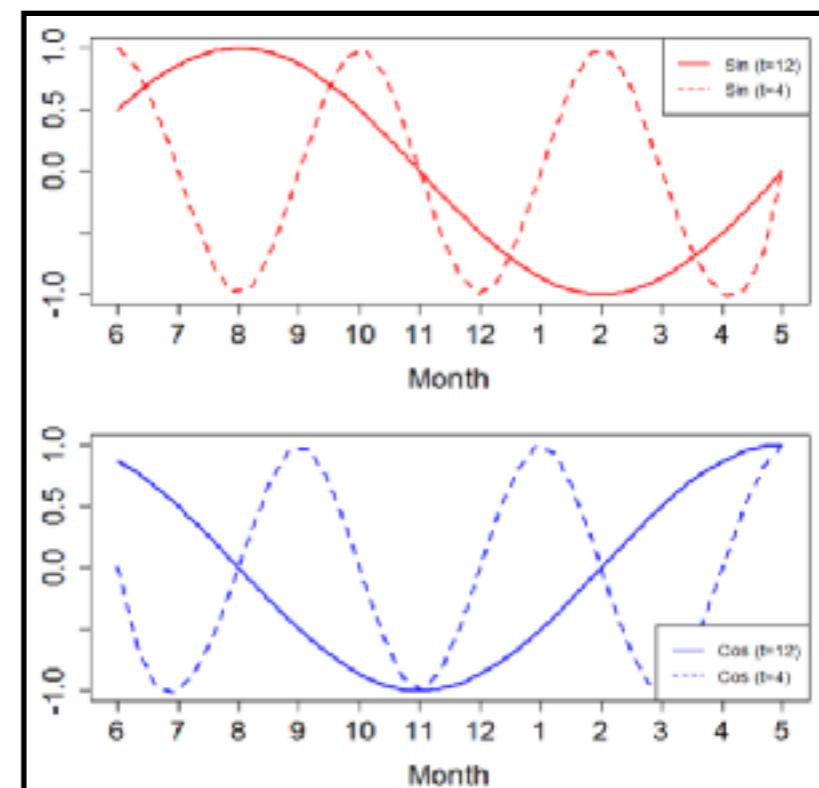
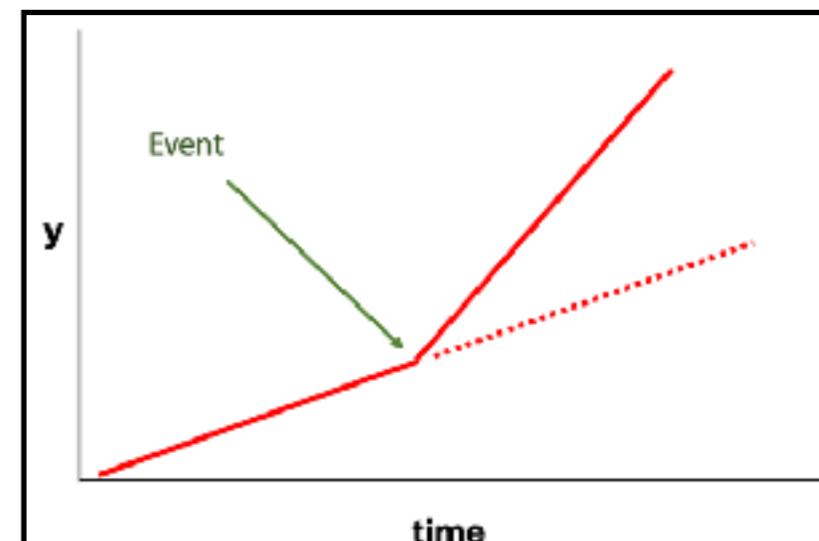
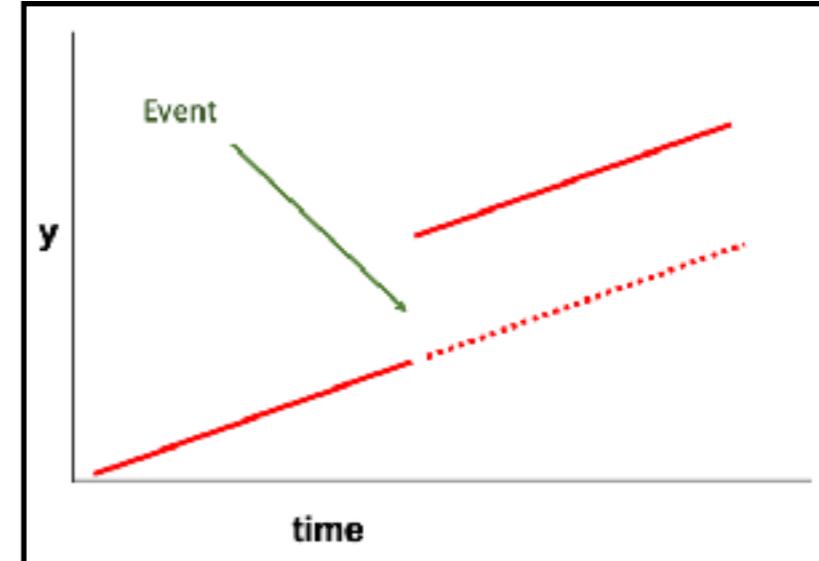
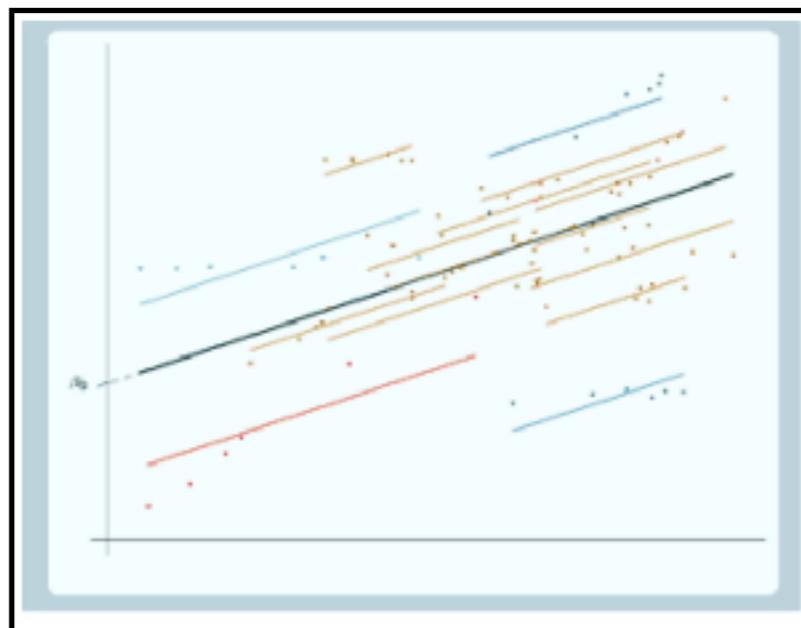
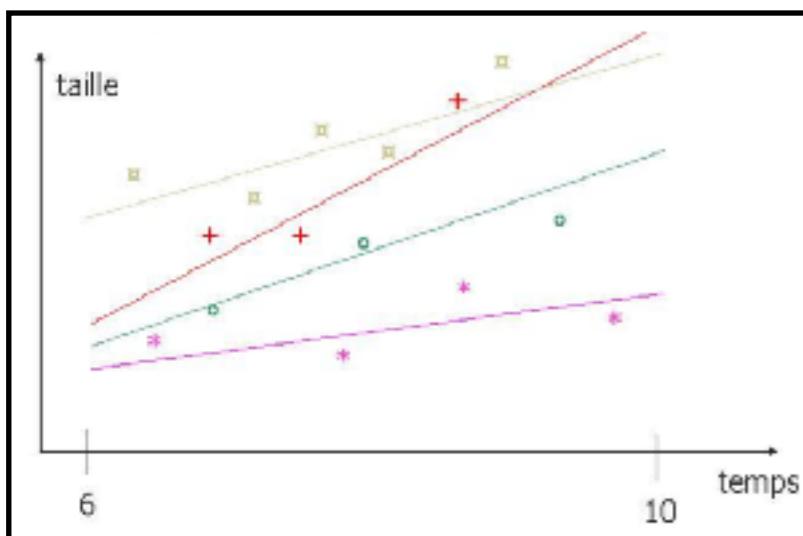
- Repeated measures
- Temporal correlation
- Spatial correlation
- Heterogeneity
- Nested data

$$y_i = X_i\beta + Z_i b_i + \varepsilon_i$$

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)`



# Network analyses



## MRQAP (Quadratic Assignment Procedure)

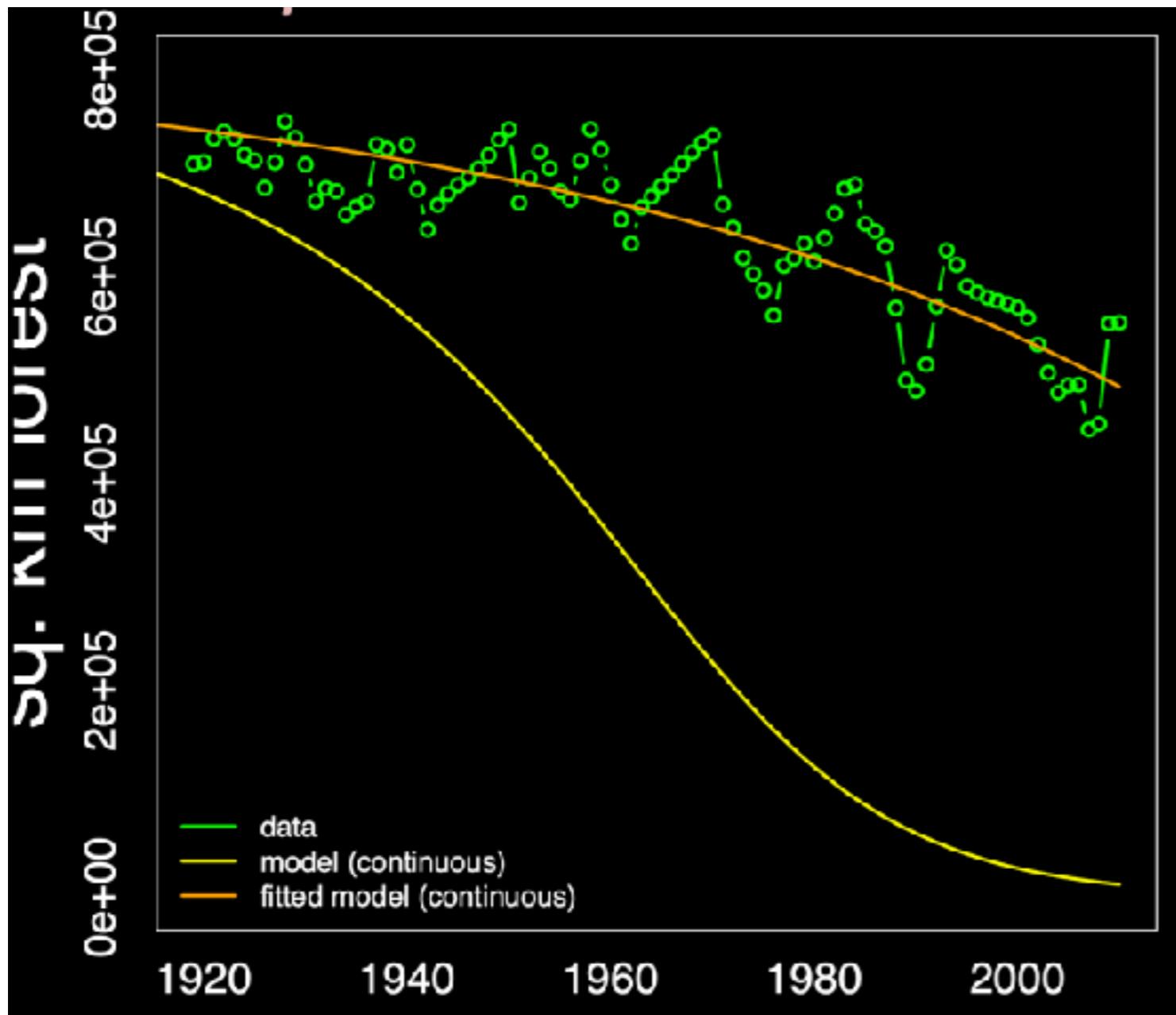


- Multiple Regression Quadratic Assignment Procedure
  - Basically logistic regression analysis applied to matrix data.
  - Is your response variable linked to explanatory variable 1 while controlling for all other variables?

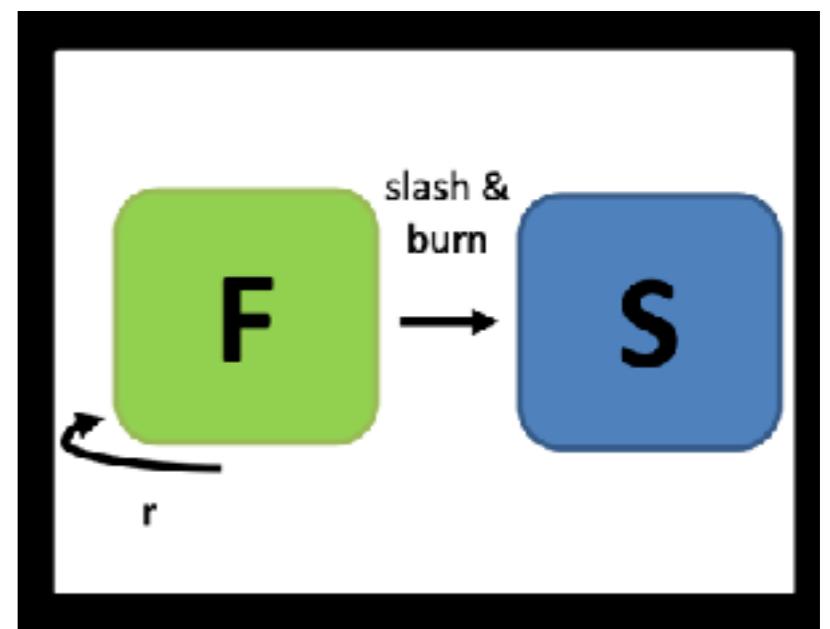
Fit mechanistic models to data

# Fit mechanistic model to data

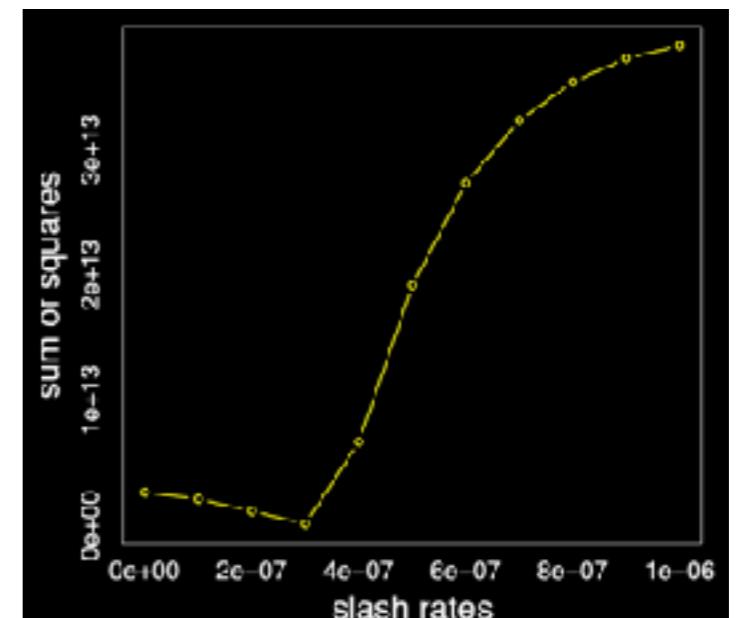
Combines the best of mechanistic and statistical method



Mechanistic model



Minimize sum of squares



# Model selection

- ❖ Some methods
  - ❖  $R^2$ , adj- $R^2$
  - ❖ Least square
  - ❖ Log likelihood
  - ❖ AIC
  - ❖ BIC
  - ❖ RMSE
  - ❖ ...
- ❖ There is no consensus among statisticians...it is an art

You can always fit a model,  
and select the best model  
But it is just the best based on what  
you tried



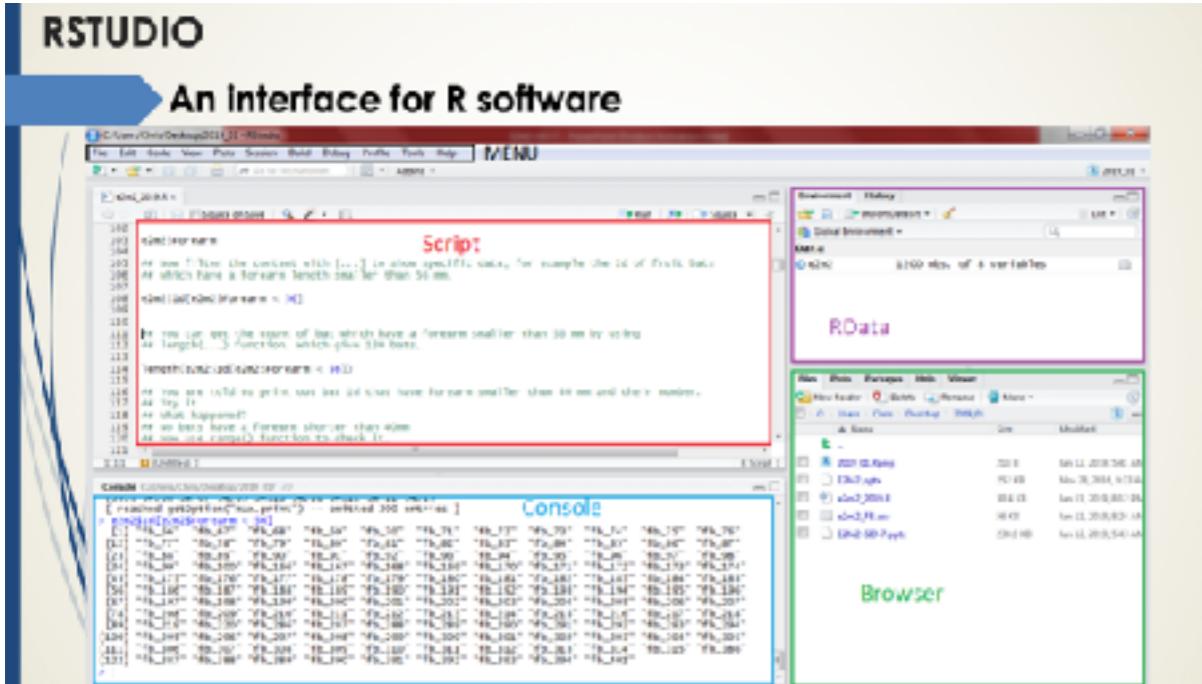
# More on quantitative skills



# Basic R

## RSTUDIO

### An Interface for R software



## Data cleaning with R

### Consistent data

#### What is wrong with this data frame?

Id	Site	Date	Sex	Weight	Balanced	gmean	stdev
bat_12	Site_3	41586	F	650	140mm	NA	1.90
bat_33	Site_3	41586	F	70	140mm	NA	3.89
bat_34	Site_2	41593	M	780	136mm	NA	30.05
bat_35	Site_2	41593	M	690	136mm	NA	30.02
bat_36	Site_2	41593	F	390	136mm	3.88	NA
bat_37	Site_2	41593	M	120	136mm	NA	9.39
bat_38	Site_2	41593	M	160	136mm	NA	NA
bat_39	Site_2	41593	F	530	136mm	4.03	NA
bat_40	Site_2	41593	M	120	136mm	NA	90.63
bat_28	Site_2	41594	F	640	136mm	4.26	NA
bat_22	Site_2	41594	F	390	136mm	4.18	NA
bat_23	Site_2	41594	M	140	136mm	NA	30.02
bat_24	Site_2	41594	F	520	136mm	4.07	NA
bat_25	Site_2	41594	M	150	136mm	NA	50.00
bat_26	Site_2	41596	F	650	136mm	4.34	NA
bat_27	Site_2	41596	F	105	136mm	4.34	NA
bat_28	Site_2	41596	M	130	136mm	NA	30.34

## Writing For Loops, If-Else Statements, and Functions in R

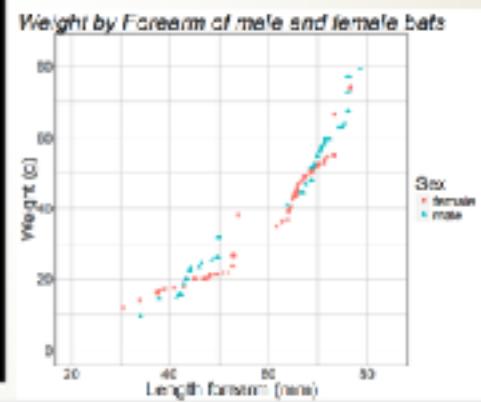
- Institut Pasteur de Madagascar
- January 2020

- EPM: Ecological and Epidemiological Modeling in Madagascar

## Visualizing Data (Present de data)

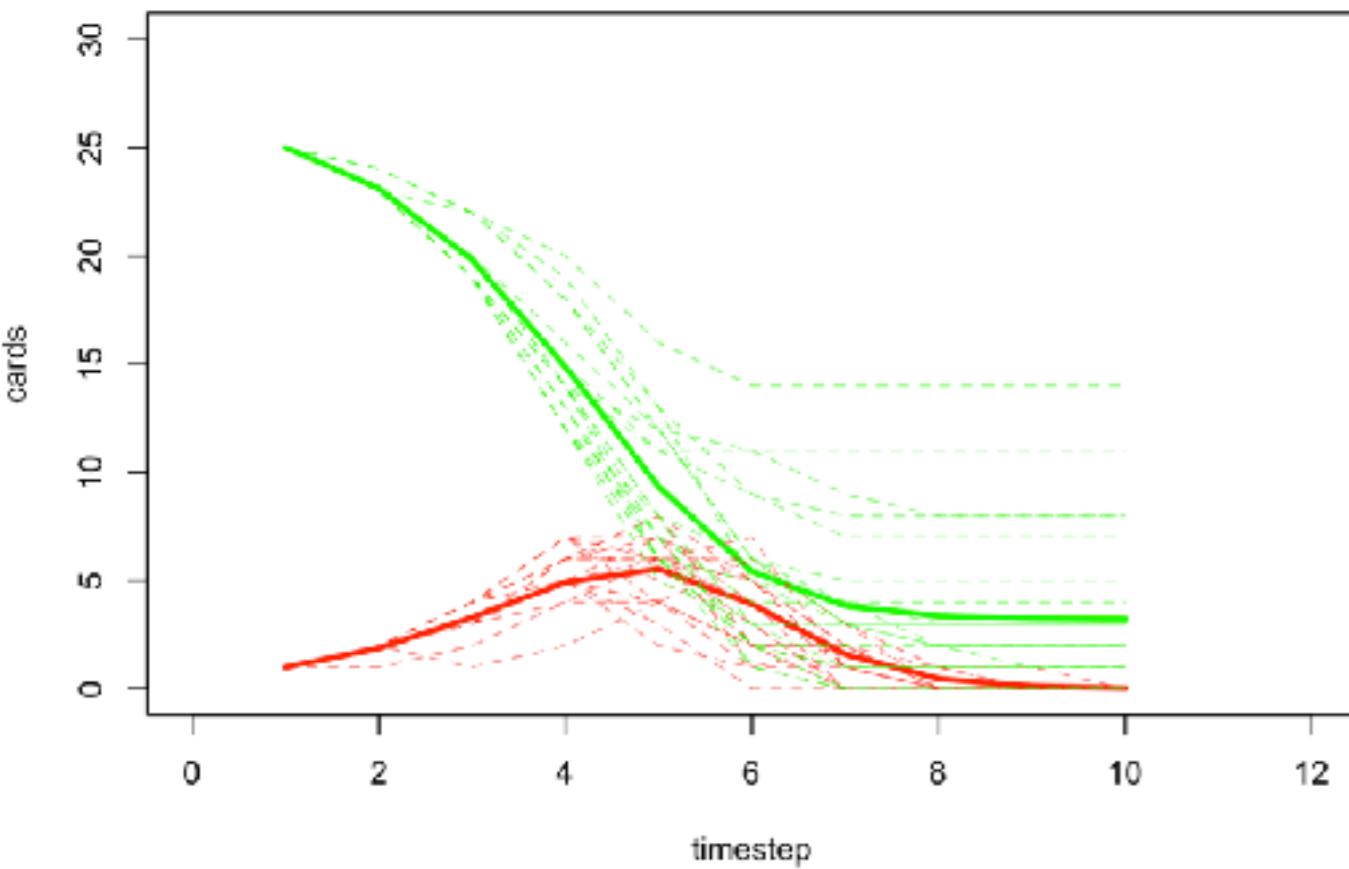
### Polish the plot

```
ggtitle("Weight by Forearm of male and female") +  
  scale_x_continuous(name="Length forearm (mm)",  
    limits=c(20,85)) +  
  scale_y_continuous(name="Weight (g)",  
    limits=c(0,85)) +  
  scale_color_discrete(name="Sex",  
    breaks=c("f","m"),  
    label=c("female","male")) +  
  scale_shape_discrete(name="Sex",  
    breaks=c("f","m"),  
    label=c("female","male"))
```



# More advanced R

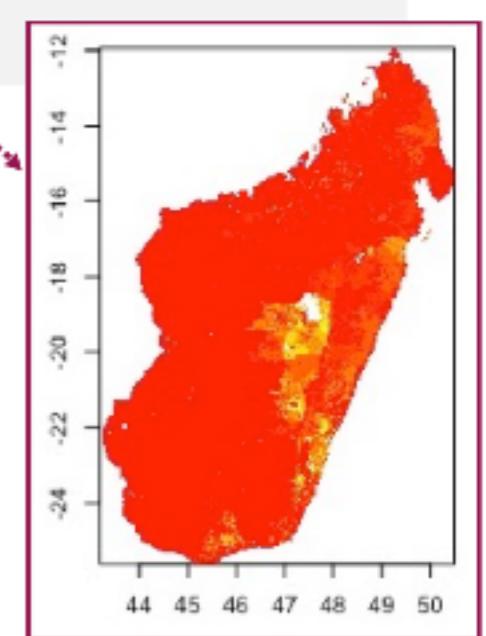
Model fitting



Spatial visualization

```
par(mfrow=c(1,3))
image(mdg_preg, col = blue2red(10))
image(log(mdg_preg+1), col = blue2red(10))
image(mdg_preg, zlim = c(0,10))
```

Plotting



Model construction

Forward simulation in time

Optimization

Precise visualization

Solving differential equations?

# Useful math

## Greek letters

A α	B β	Γ γ	Δ δ	E ε	Z ζ	H η	Θ θ
alpha	beta	gamma	delta	epsilon	zeta	eta	theta
α	β	γ	δ	ε	ζ	η	θ
[aɪə]	[b]	[g]	[d]	[e]	[zɪ/dz]	[eɪ]	[θ]
I i	K κ	Λ λ	M μ	N ν	Ξ ξ	O o	Π π
iota	kappa	lambda	mu	nu	xi	omicron	pi
ι	κ	λ	μ	ν	ξ	ο	π
[aɪ]	[k]	[l]	[m]	[n]	[zɪ]	[o]	[p]
P ρ	Σ σ/ς	T τ	Υ υ	Φ φ	X χ	Ψ ψ	Ω ω
rho	sigma	tau	upsilon	phi	chi	psi	omega
ρ	σ	τ	υ	φ	χ	ψ	ω
r-1	r-1	r-1	r-1	r-1	r-1	r-1	r-1

## Eigenvectors and eigenvalues

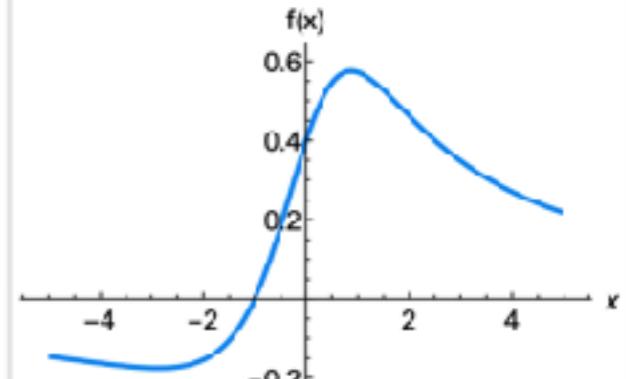
$$M = \begin{bmatrix} 1 & 2 & 0 \\ 0 & 4 & 1 \\ 0 & -2 & 1 \end{bmatrix} \quad v_1 = \begin{bmatrix} -1 \\ -1 \\ 1 \end{bmatrix} \quad v_2 = \begin{bmatrix} -2 \\ -1 \\ 2 \end{bmatrix} \quad v_3 = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix}$$

- $M \cdot v_1 = 3v_1$  if  $V = [v_1 \ v_2 \ v_3] = \begin{bmatrix} -1 & -2 & 1 \\ -1 & -1 & 0 \\ 1 & 2 & 0 \end{bmatrix}$
- $M \cdot v_2 = 2v_2$
- $M \cdot v_3 = 1v_3$  and  $\Lambda = \begin{bmatrix} 3 & 0 & 0 \\ 0 & 2 & 0 \\ 0 & 0 & 1 \end{bmatrix}$  then  $M = V^{-1}\Lambda V$

## Function properties

- Intercept/root(s)
- Positive/negative value
- Maximum/minimum value
- Increasing/Decreasing/ Constant
- Concave/Convex
- Asymptotic

$$f(x) = \frac{x+1}{x^2 + 2.5}$$



GLMM Network

LM, GLM

The new you

Advanced R

Basic R

Compartment model

Basic math

Research questions





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7. *Manuscript writing and submission*

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# You are now well equipped

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