

Introduction to Linear Regression



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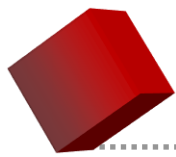
Association Ekipa Fanihy (Efa), Madagascar.

Zoologie
et
Biodiversité Animale



Adapted from Andrés Garchitorena, PhD
(E2M2, 2024)

E²M² Workshop
Andasibe, May 2025



Objectives of the lecture

- Remind some basic principles of linear regression
- Introduction to generalized linear models
- Provide an overview of the steps involved in developing a generalized linear mixed model (tutorial)
- « Understand alternatives to the use of mathematical models for the study of dynamical systems »

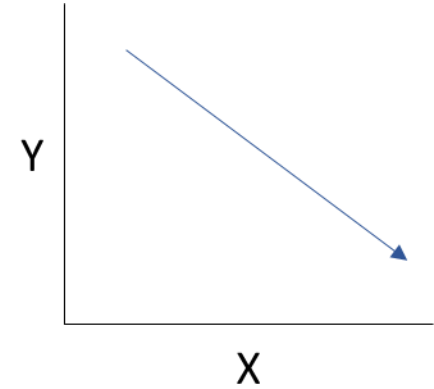
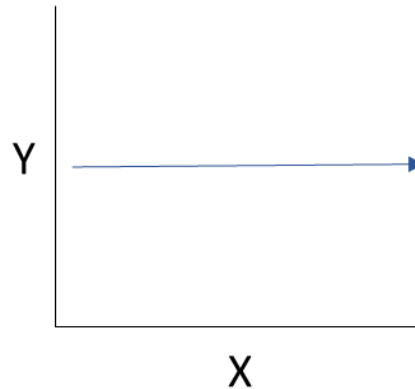
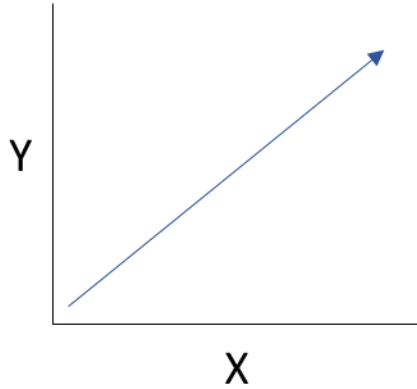
Being able to use a tool properly

‘ We are introducing to what is “available” (E2M2),
it is up to you to deepen what you think is “worth it” (C4C)’

SOME BASICS FIRST...

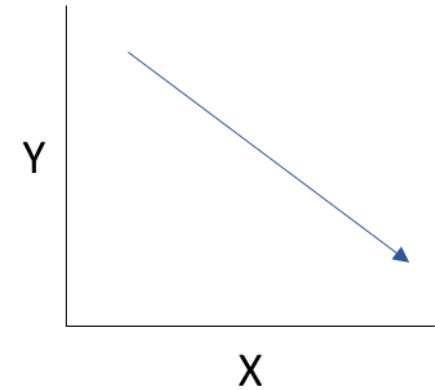
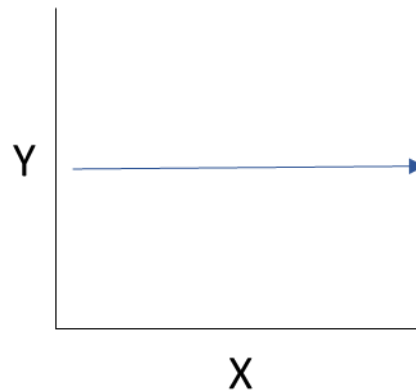
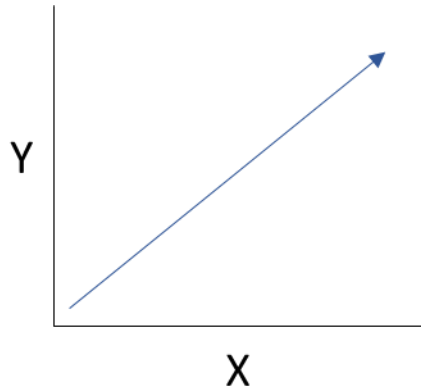
Linear Models

Think about a straight line to describe a trend



SOME BASICS FIRST...

Think about a straight line to describe a trend

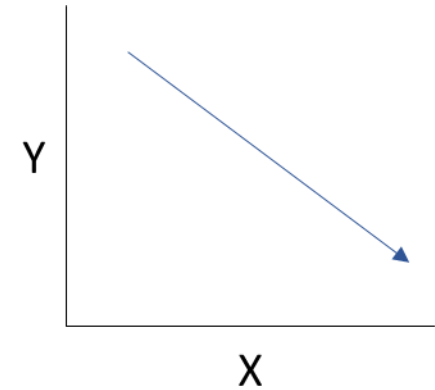
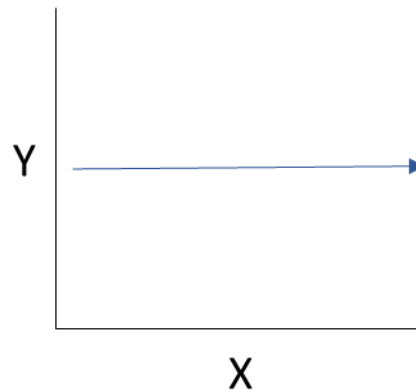
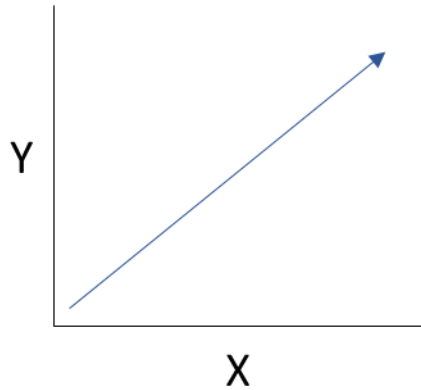


$$y = f(x)$$

SOME BASICS FIRST...

Linear Models

Think about a straight line to describe a trend



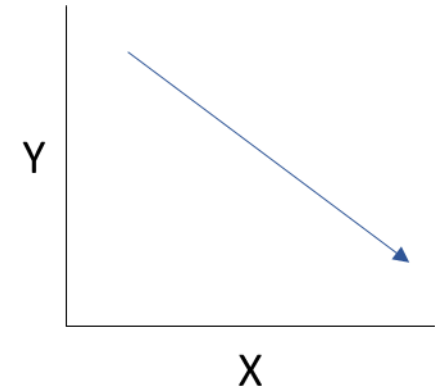
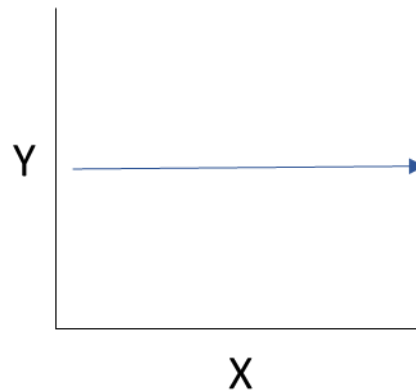
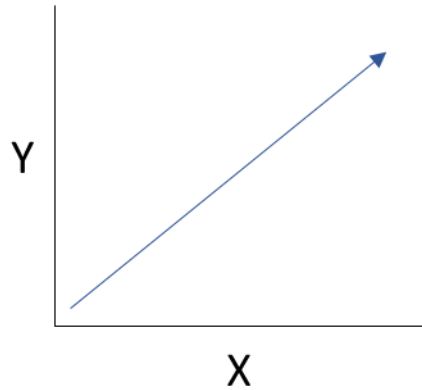
$$y = f(x)$$

Explanatory variable
Independent variable
(Variable explicative, axe des abscisses)

SOME BASICS FIRST...

Linear Models

Think about a straight line to describe a trend



$$y = f(x)$$

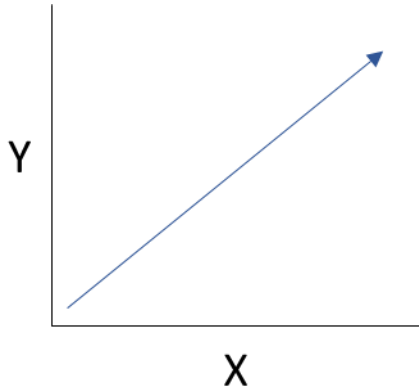
Response variable
Dependent variable
(Variable à expliquer, axe des ordonnées)

Explanatory variable
Independent variable
(Variable explicative, axe des abscisses)

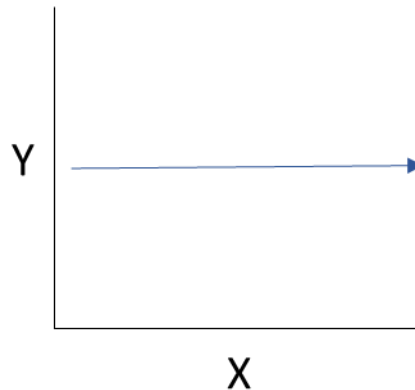
SOME BASICS FIRST...

Linear Models

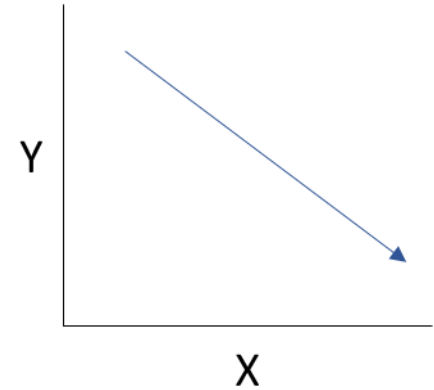
Think about a straight line to describe a trend



Going up (Miakatra)



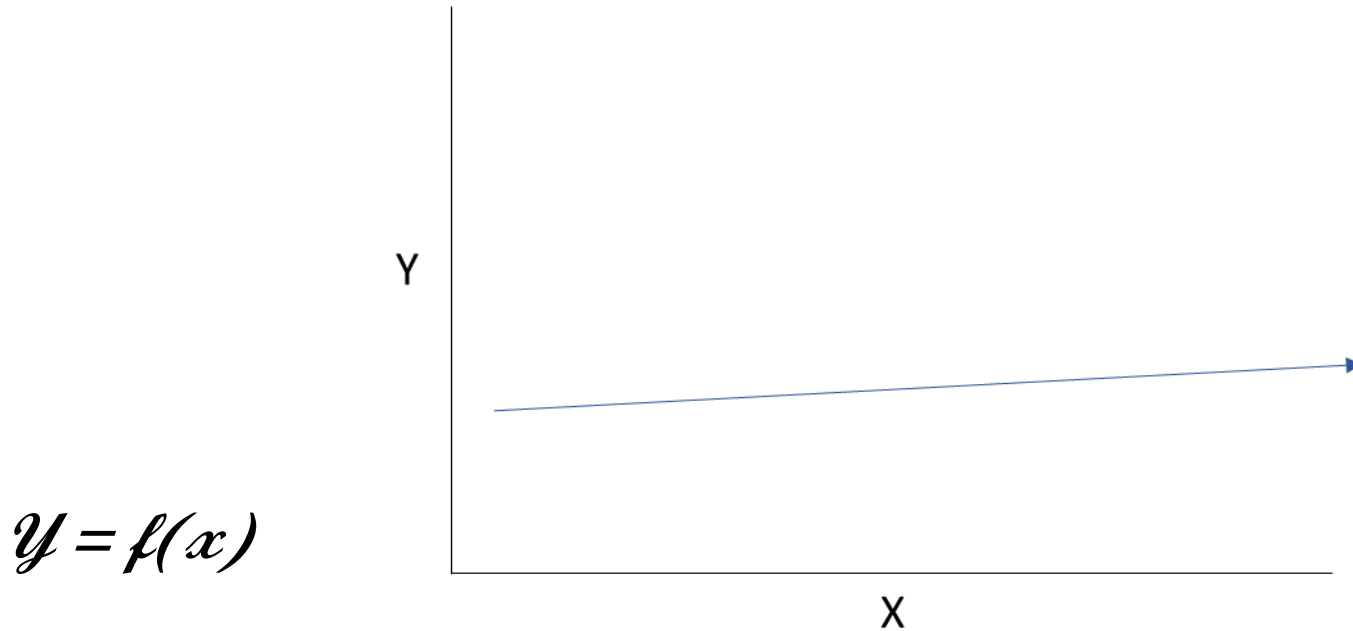
Steady (Tsy miova)



Going down (Mihidina)

SOME BASICS FIRST...

What about this one? (Dia ahoana izany ity?)



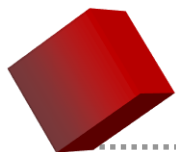
SOME BASICS FIRST...

Considering the effect of **only one explanatory variable (x)**
on a response variable **(y) with a normal distribution**

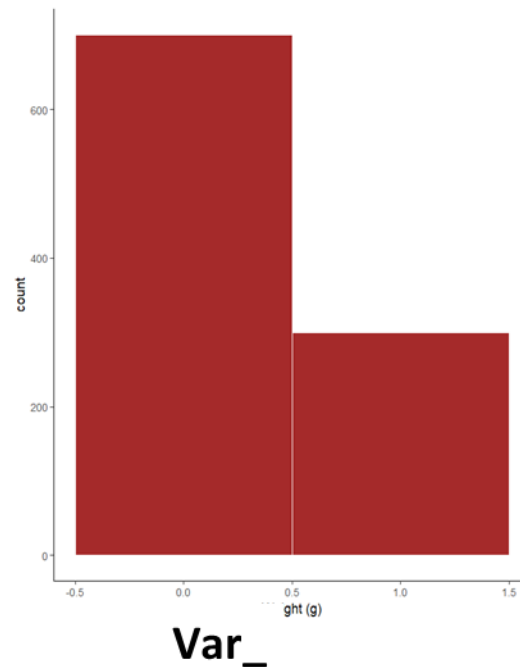
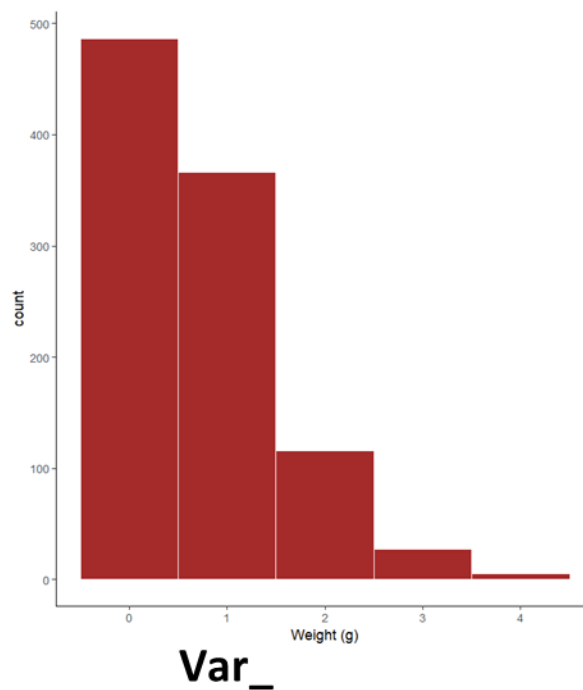
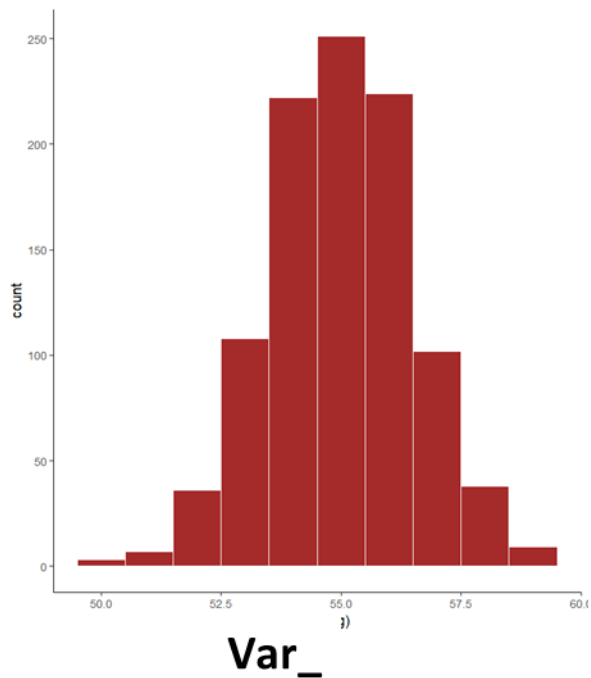
(Etudier « l'effet » une seule variable explicative sur une variable
réponse ayant une distribution normale)

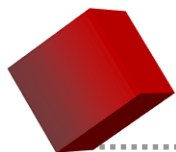
(Azo tsotsorina hoe fiakinan'ny “y” amin'ny “x” iray ihany)

SOME BASICS FIRST...

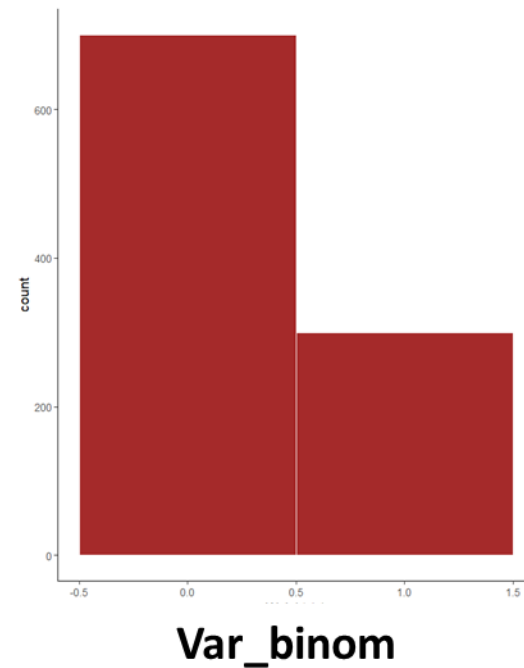
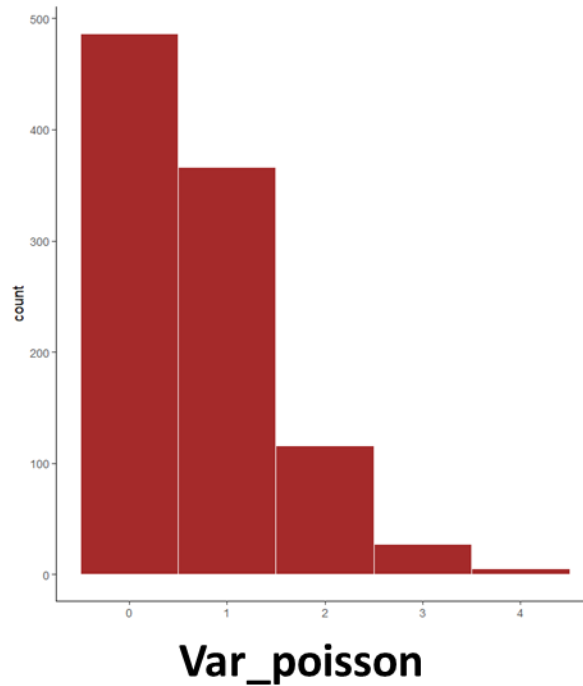
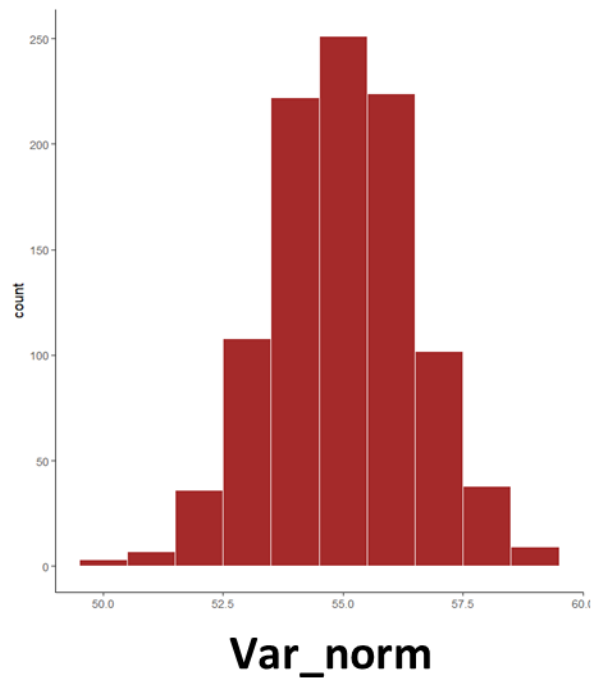


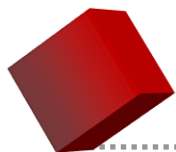
Variables and distributions



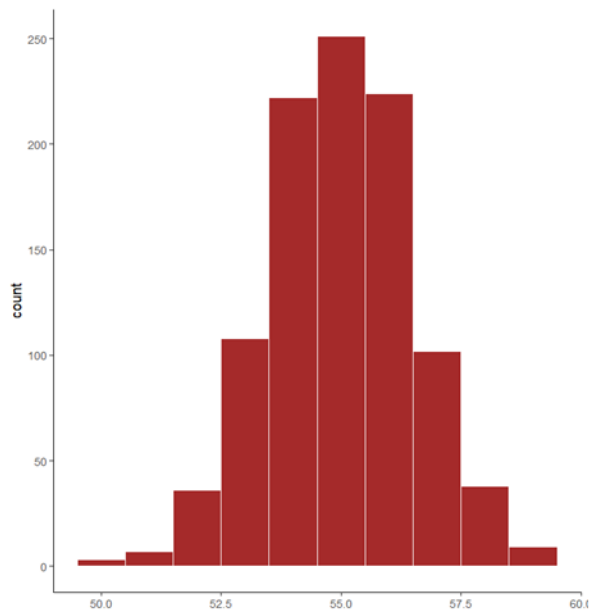


Variables and distributions



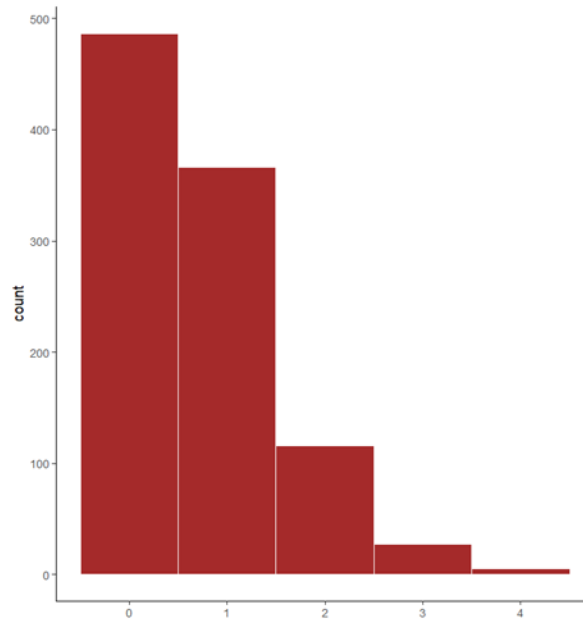


Variables and distributions



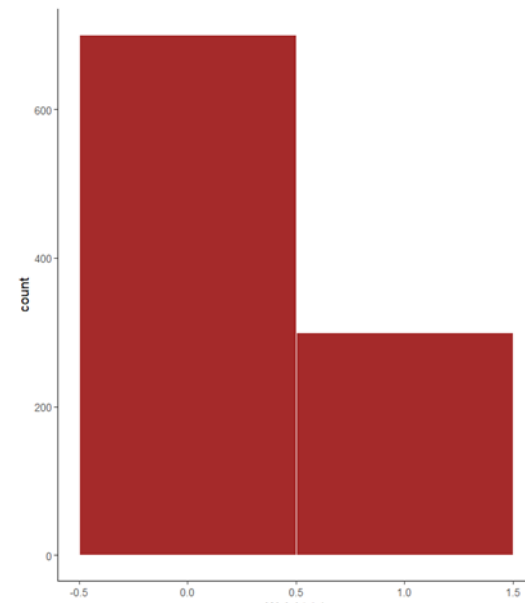
Var_norm

Continuous



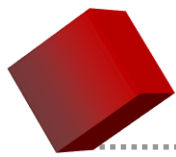
Var_poisson

Integer (Count)



Var_binom

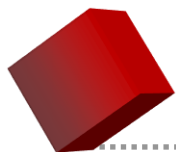
Presence/Absence



Let's work through a cute example

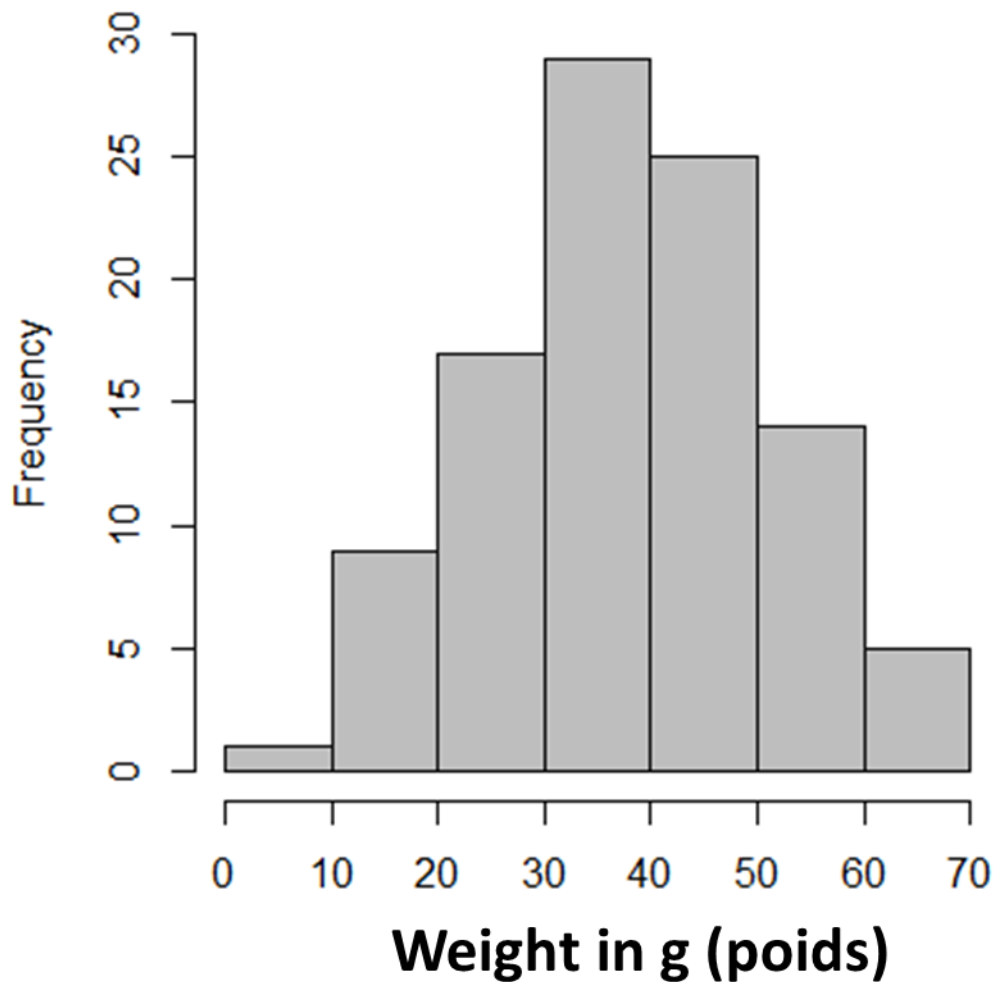


Lemurs weight (poids) determinants



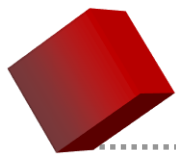
Lemur weight and determinants

Histogram of weight (poids) of lemurs

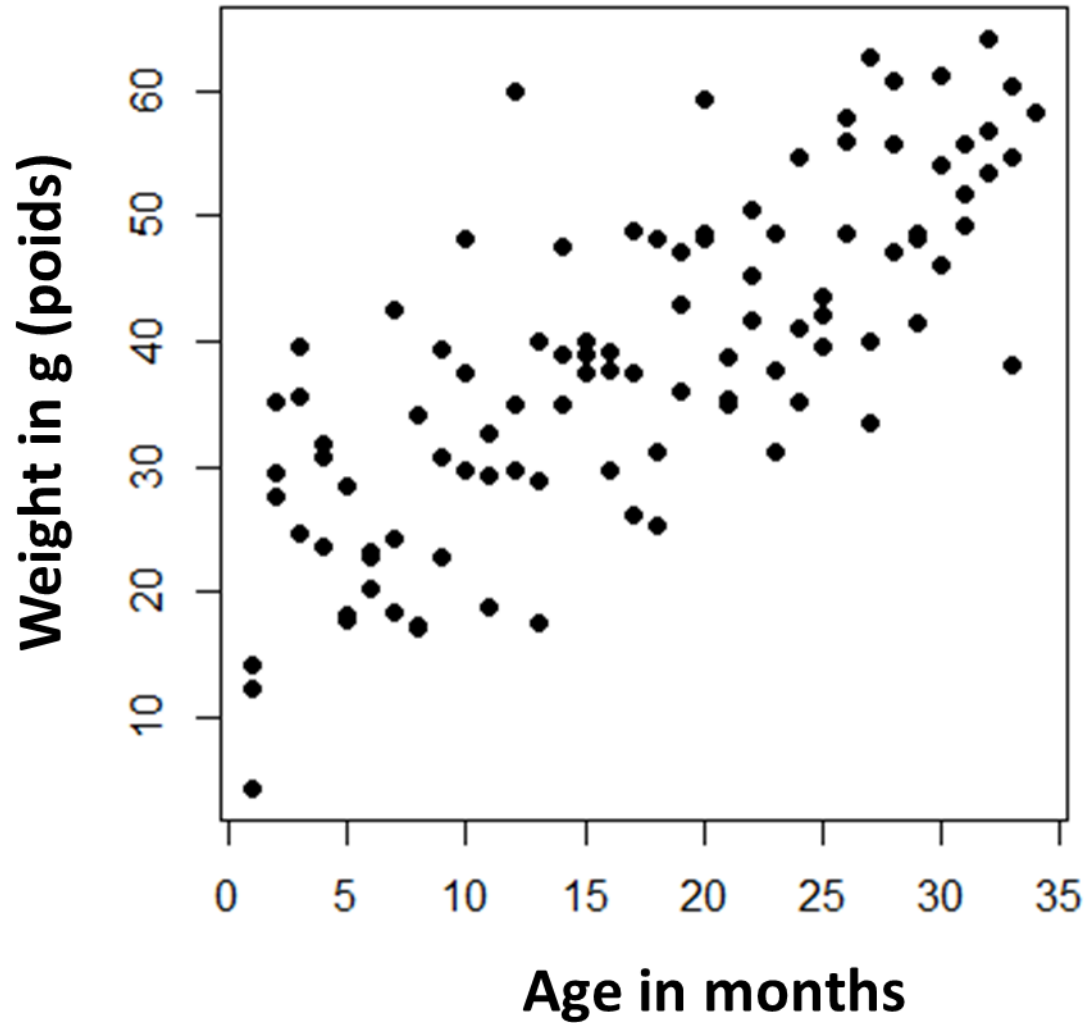


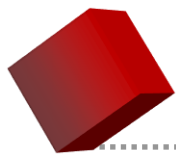
$$Z_i = \frac{X_i - \bar{X}}{\sigma}$$

$$Z \sim N(0, 1)$$

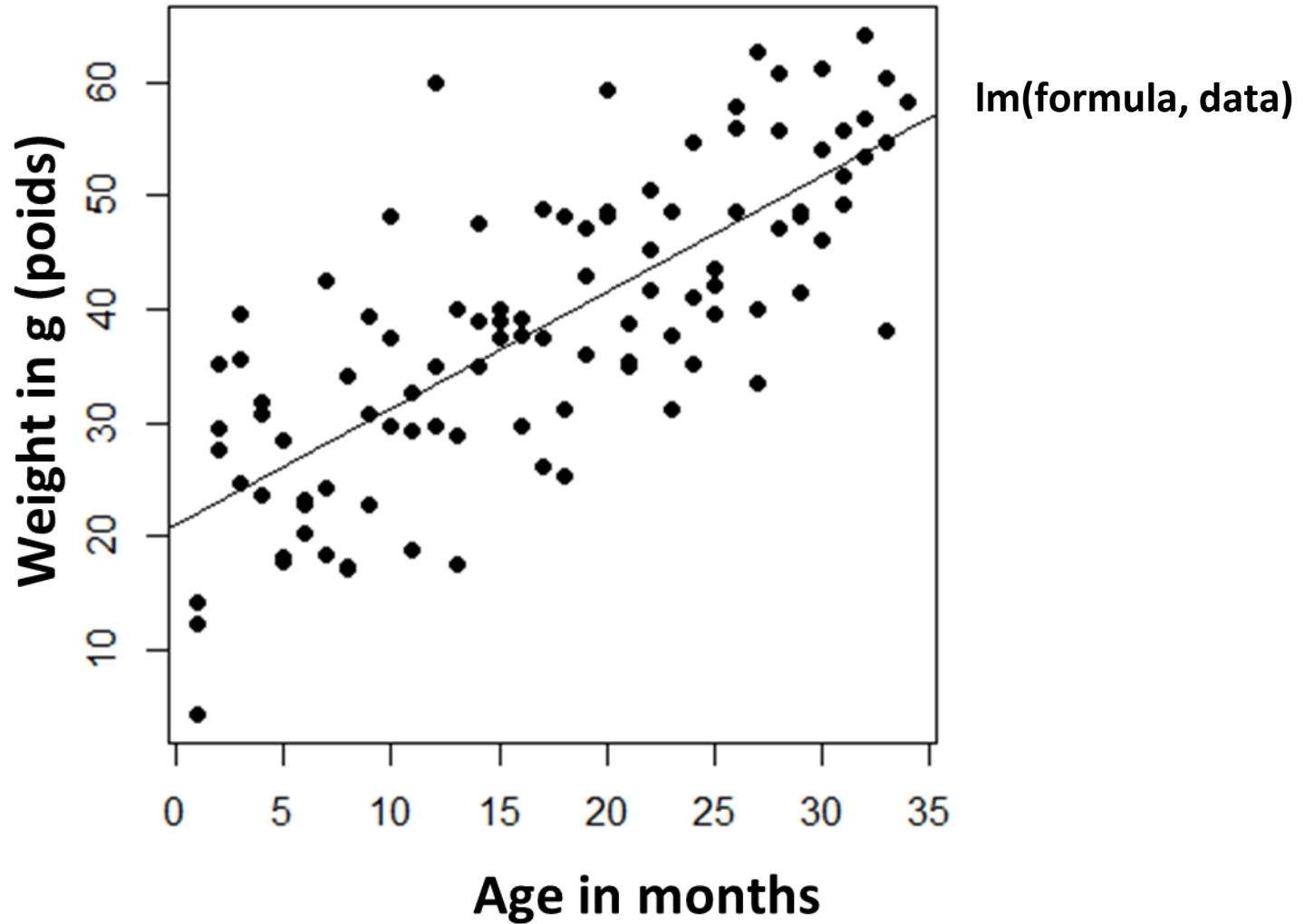


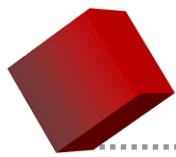
Lemur weight and determinants



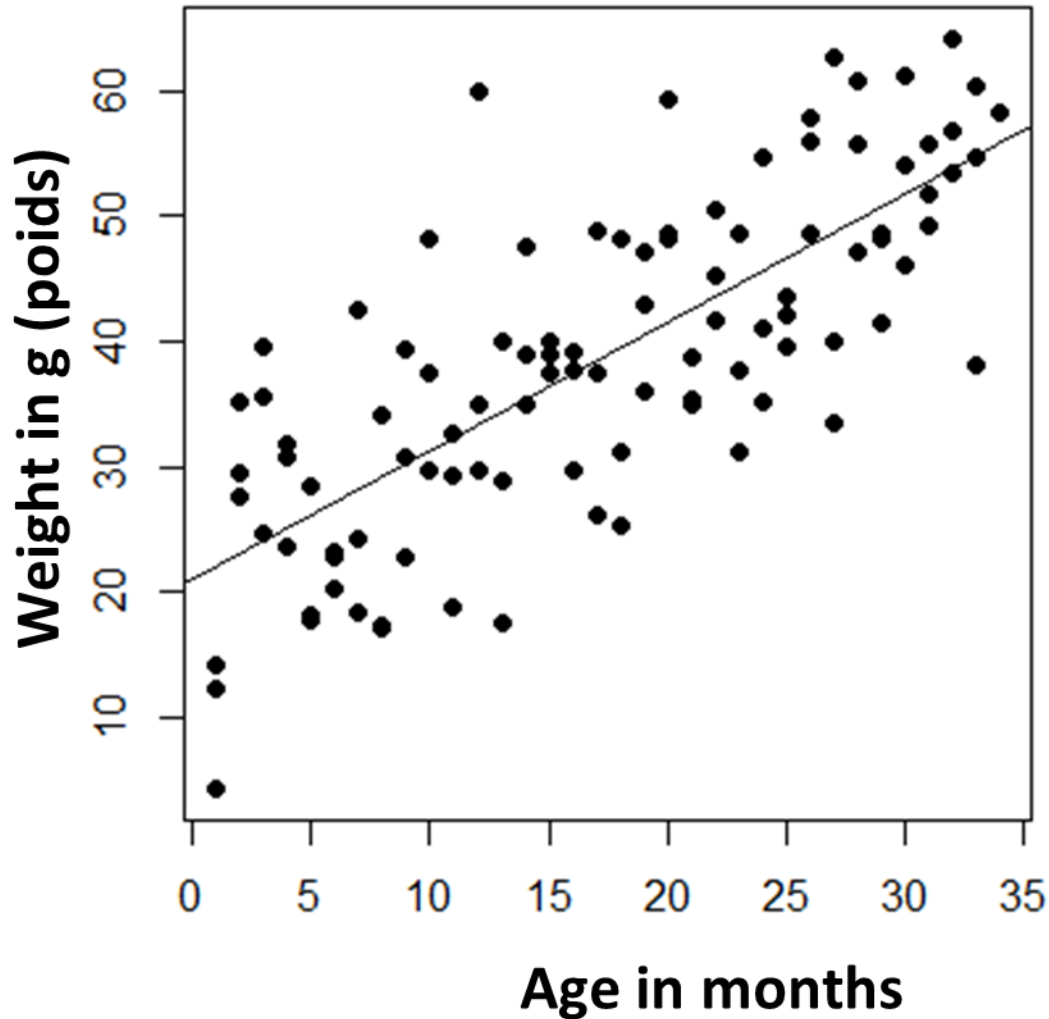


Lemur weight and determinants



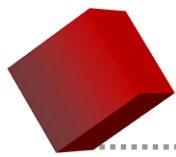


Lemur weight and determinants

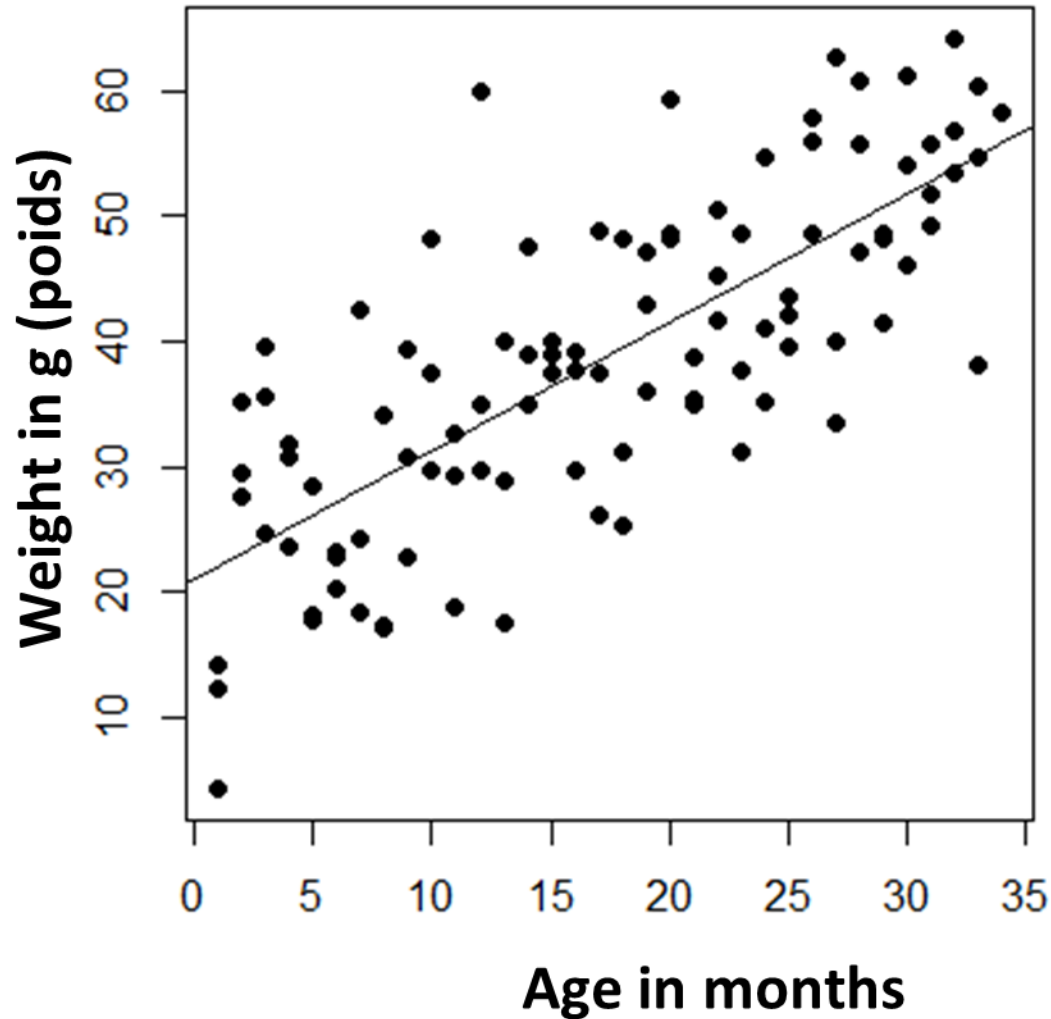


`lm(formula, data)`

Formula : $y = f(x)$

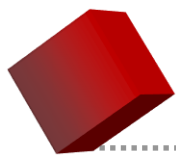


Lemur weight and determinants



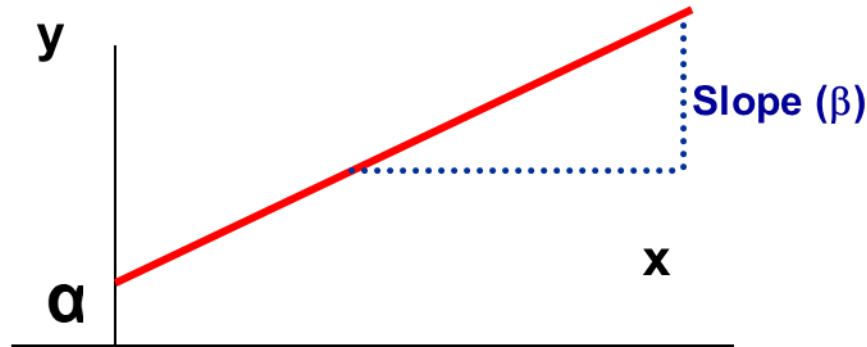
`lm(weight ~ Age, data)`

`"lm(formula, data)"`

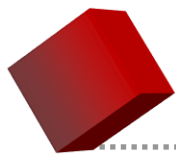


Simple linear regression

- Relation between 2 continuous variables



- *Intercept (α)*
 - Value of y when x is 0
- *Regression coefficient β_1*
 - Measures association between y and x
 - Amount by which y changes on average when x changes by one unit
- *Error (ε)*
 - Difference between the predicted values and observed values of y



Simple linear regression

$$f(x) = \alpha + \beta x + \varepsilon$$

Response variable =

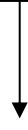
Systematic
component

+

Residual
component



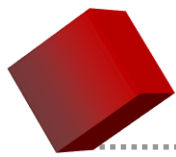
Intercept and
explanatory variables



- Null mean
- Independence
- Fixed variance
- Normality

$$y = \alpha + \beta x + \varepsilon$$

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



Simple linear regression

$$f(x) = \alpha + \beta x + \varepsilon$$

Response variable =

Systematic
component

+

Residual
component

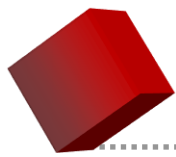
Intercept and
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- Null mean
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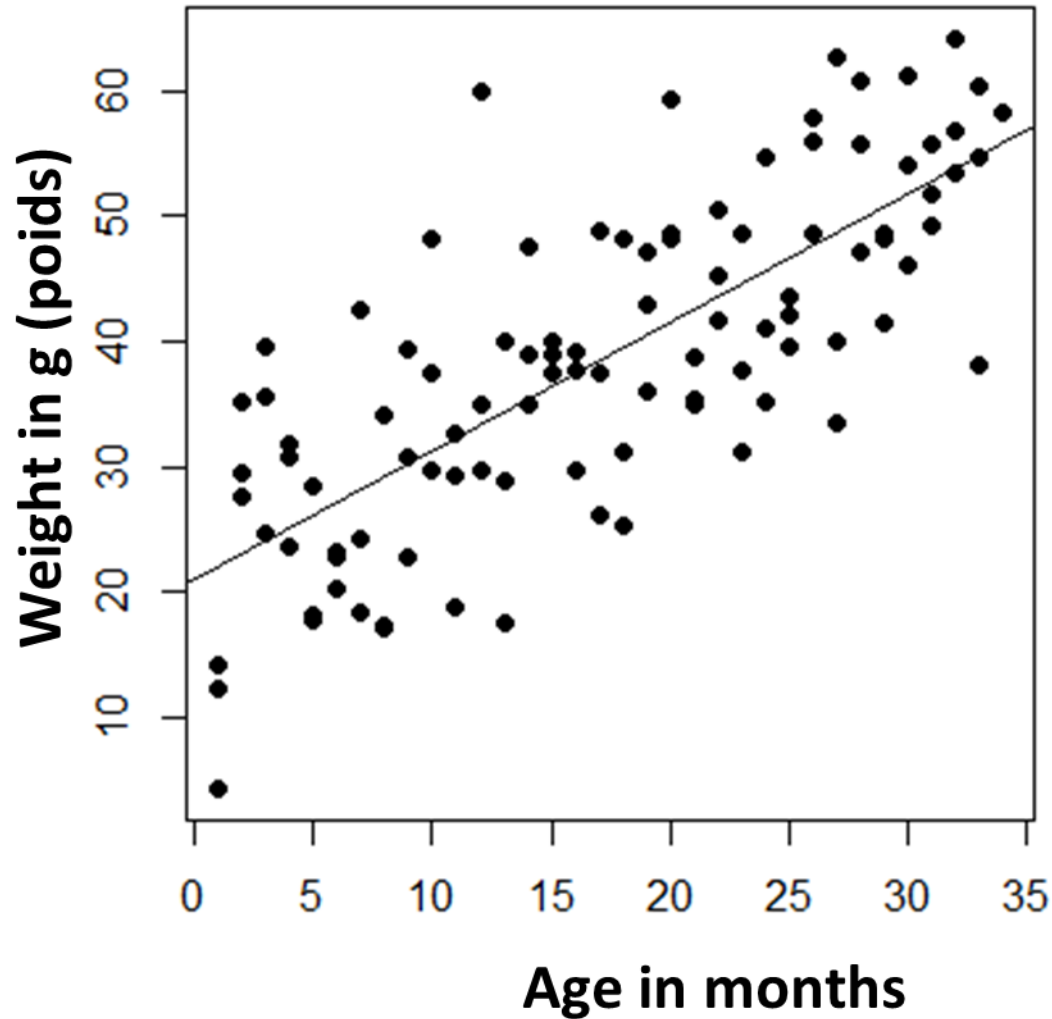
$$y = \alpha + \beta x + \varepsilon$$

$$\text{weight} = \alpha + \beta.\text{age} + \varepsilon$$

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



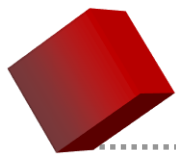
Lemur weight and determinants



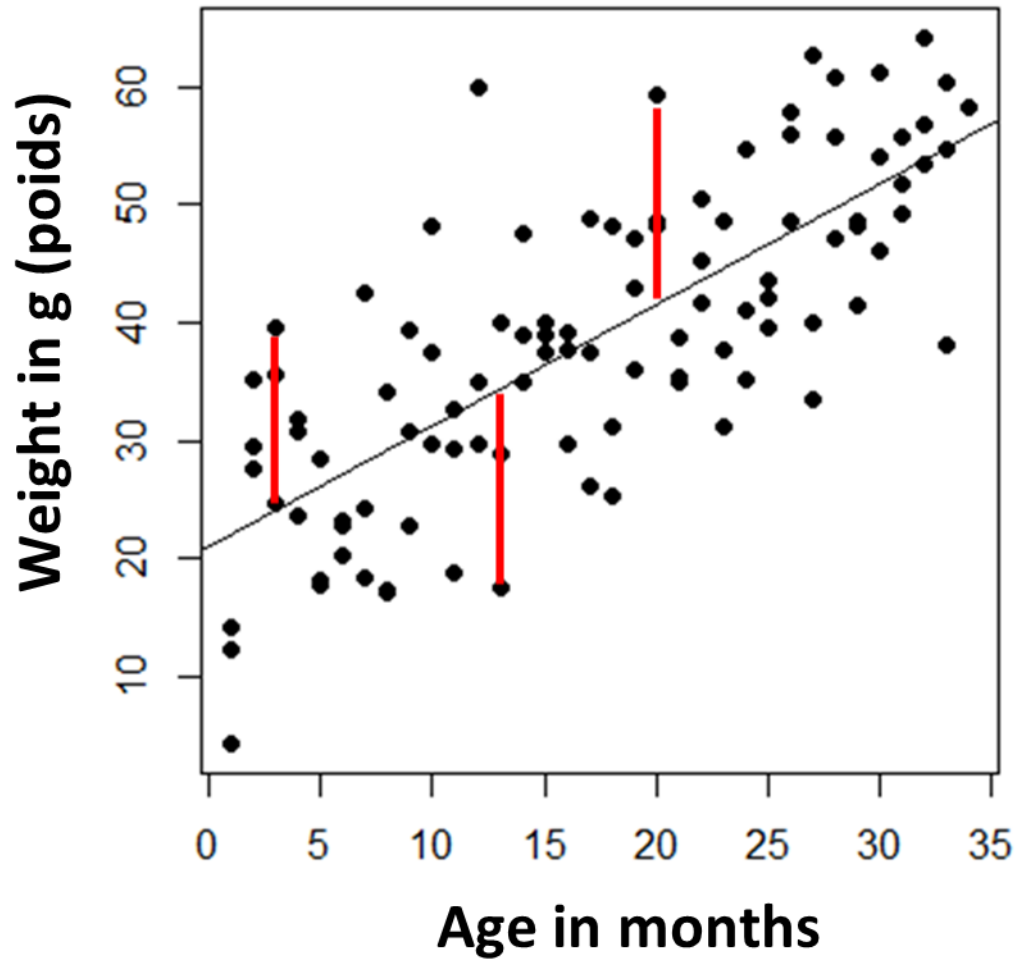
lm(weight ~ Age, data)

Formula : $y = f(x)$

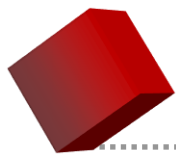
$$y = \alpha + \beta x + \varepsilon$$



Lemur weight and determinants

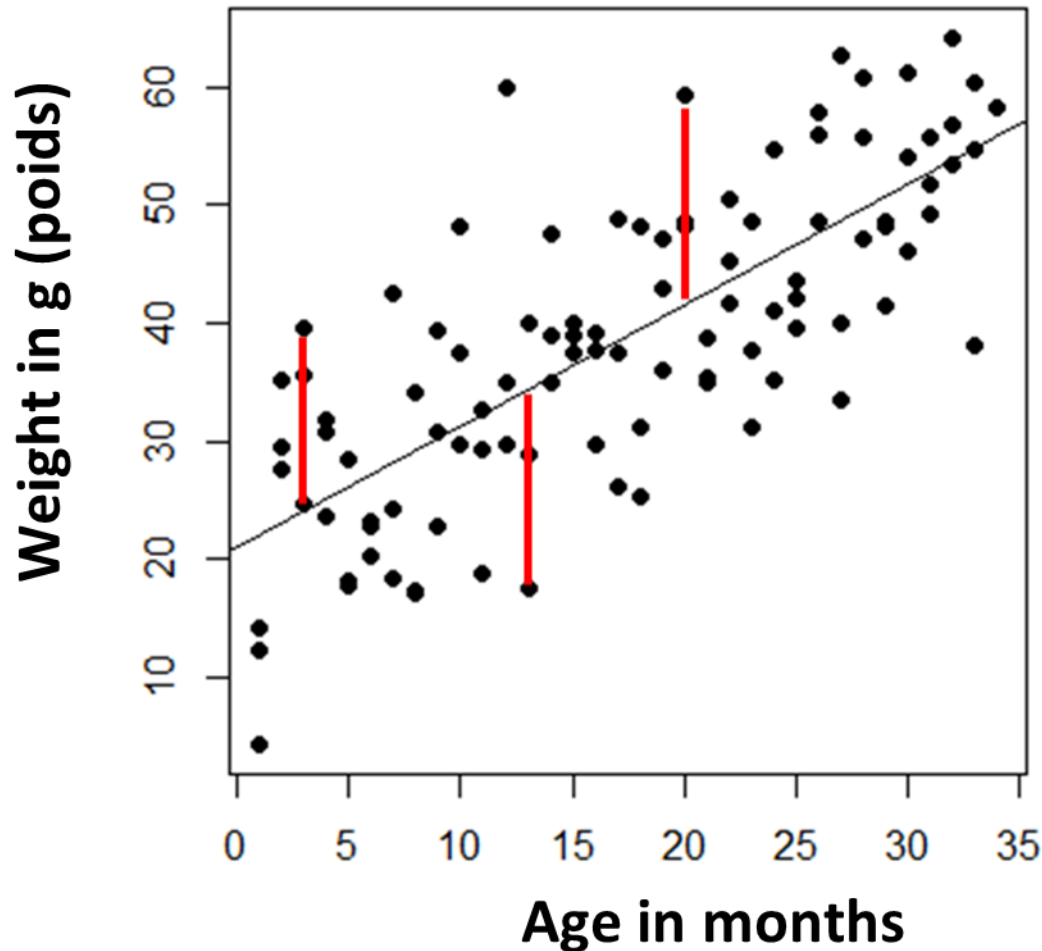


$$y = \alpha + \beta x + \epsilon$$

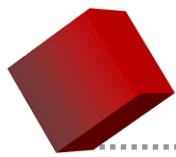


Lemur weight and determinants

The goal is to minimize the difference between what we predict and what we observe
(Méthode des moindres carrés ry reto an!)

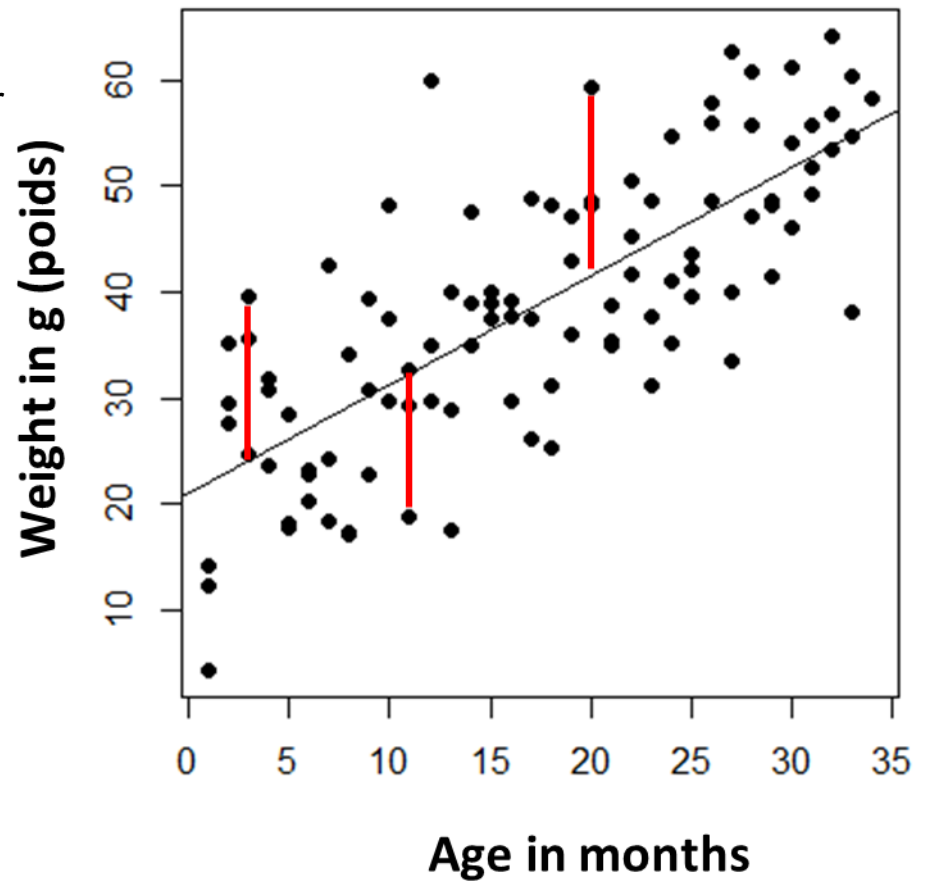


$$y = \alpha + \beta x + \epsilon$$



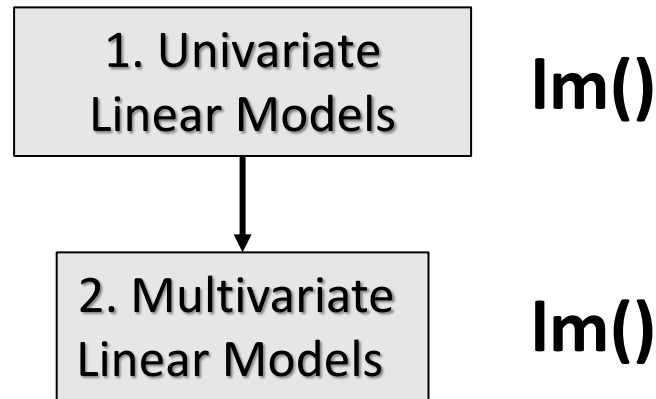
Simple linear regression

$$\text{weight} = 20 + 1.15 \times \text{Age (months)} + \text{Error}$$



A process is generally the result
of several others...

Mety hoe tsy ny mahabe ny taona ihany no mampavesatra ny lanja an!

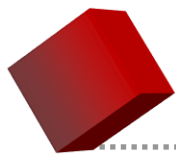


Considering the effect of **multiple explanatory variables** ($x_1, x_2 \dots x_n$) on a response variable (**y**) with a normal distribution

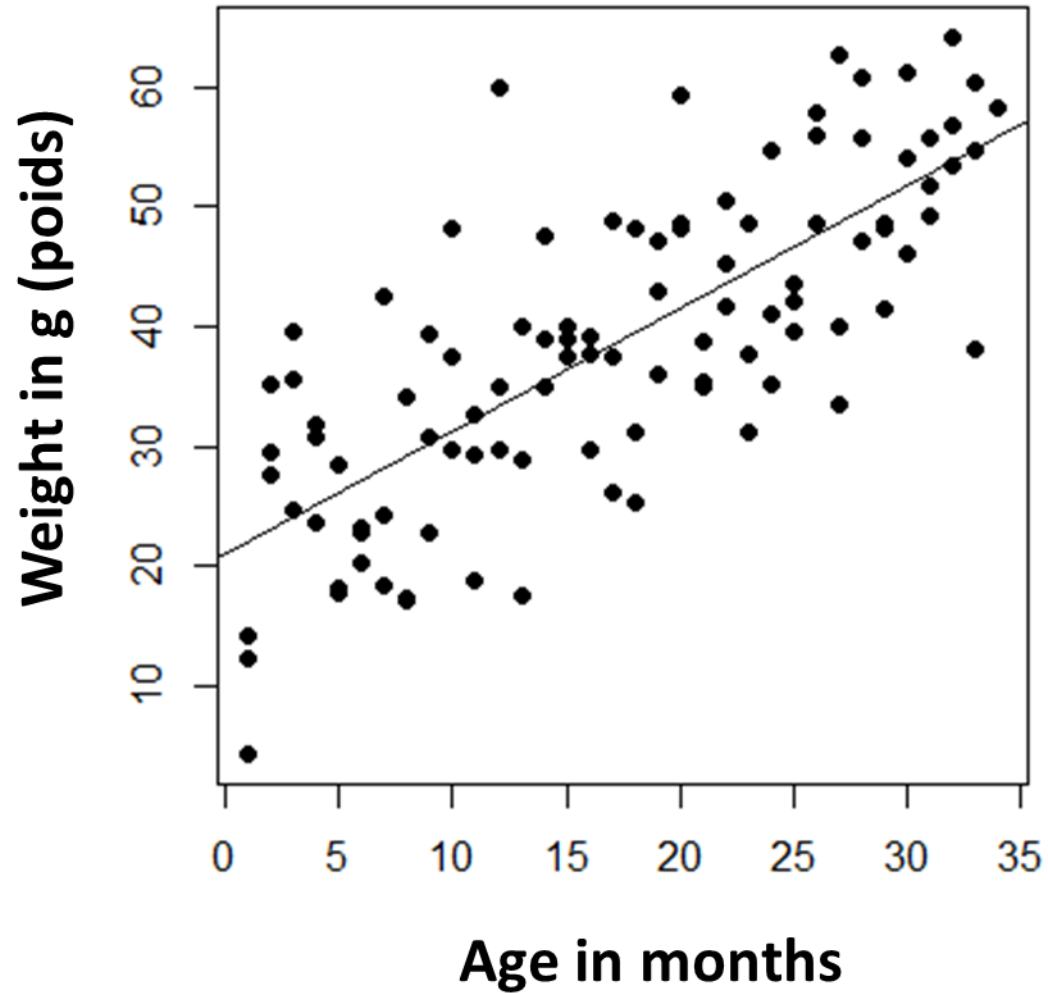
(Etudier « l'effet » de plusieurs variables explicatives sur une variable réponse ayant une distribution normale)

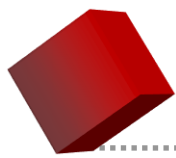
(Azo tsotsorina hoe fiakina ny "y" amin'ny "x" maromaro)

INTRODUCING MULTIVARIATE LINEAR MODELS



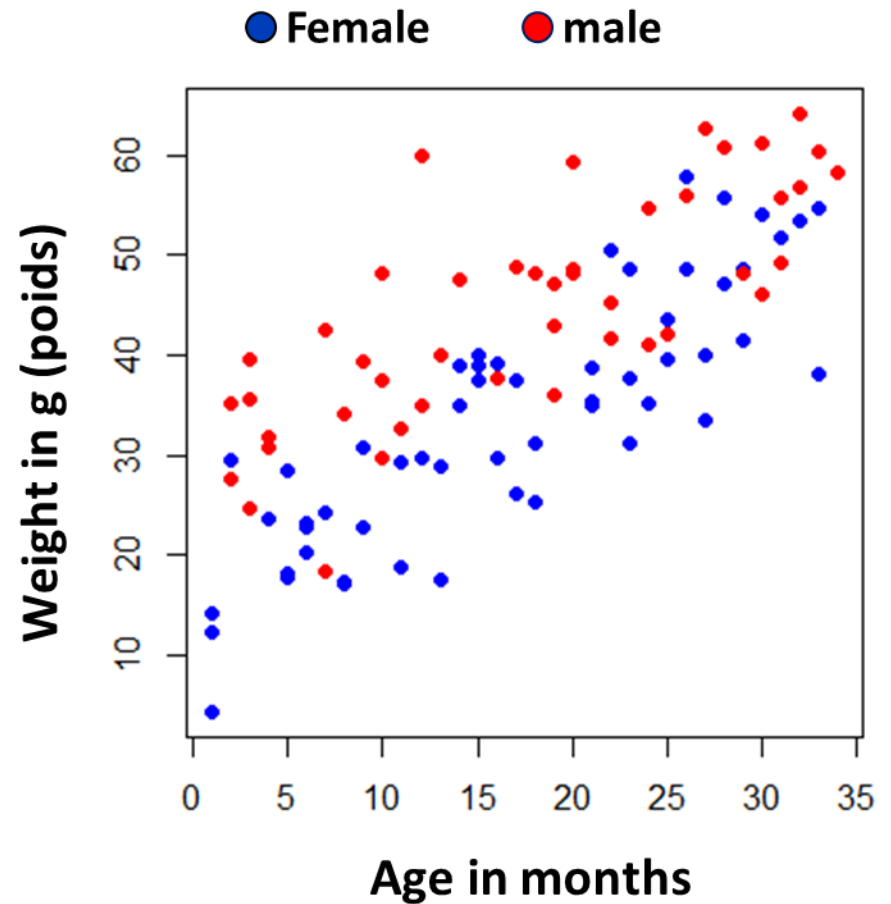
Lemur weight and determinants

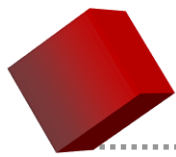




Lemur weight and determinants

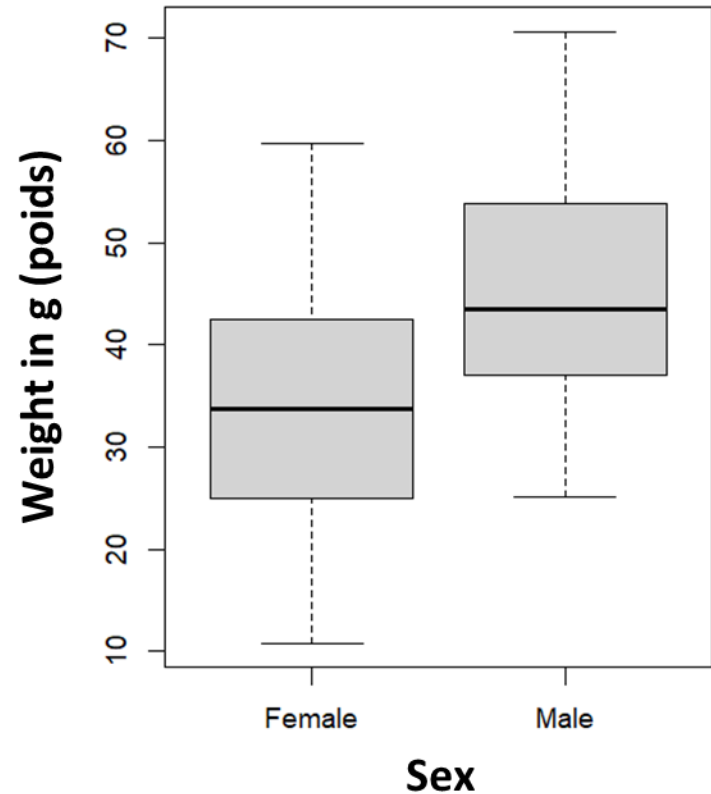
The effect of gender



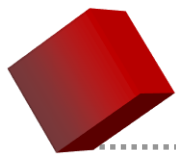


Lemur weight and determinants

The effect of gender



$$\text{Taille} = 15 + 1.15 \times \text{Age (months)} + 15 \times \text{Sexe (male)} + \text{Error}$$

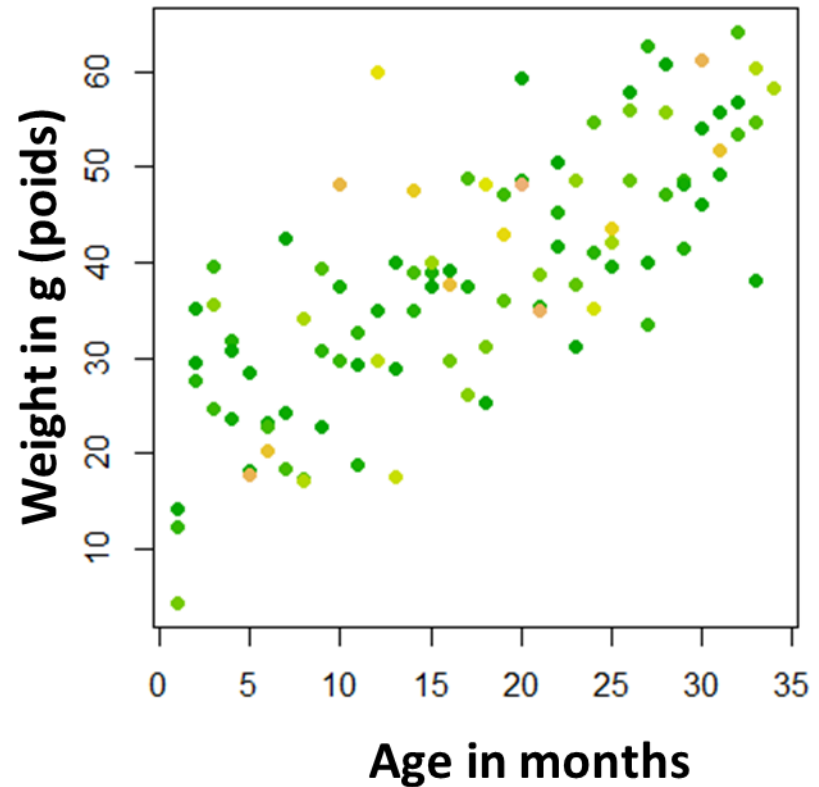


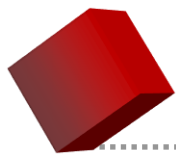
Lemur weight and determinants

The effect of parasites



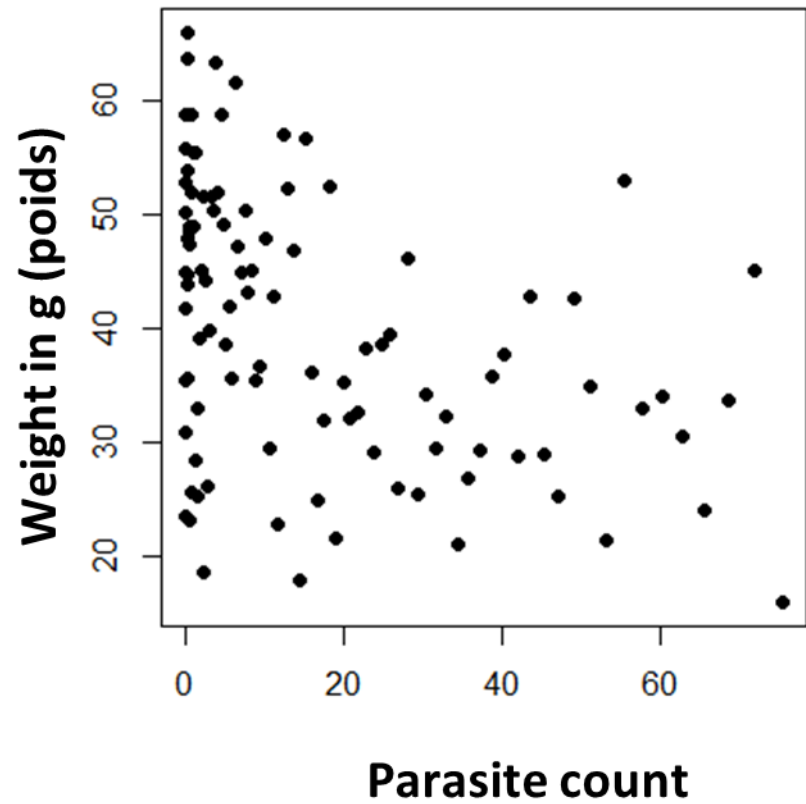
Green: low parasite burden
Yellow: high parasite burden

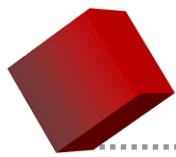




Lemur weight and determinants

The effect of parasites



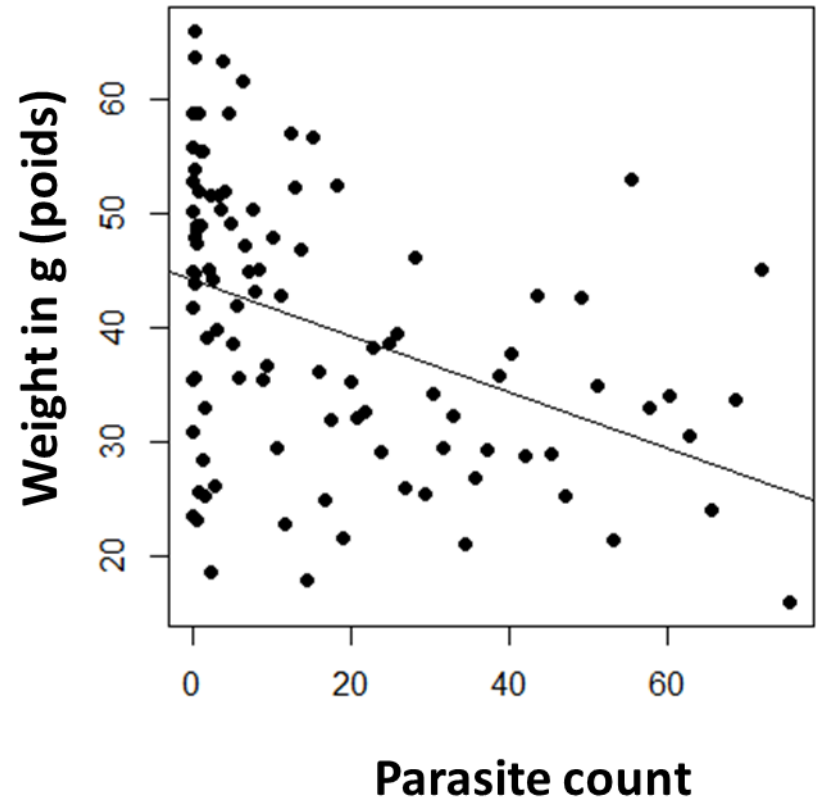


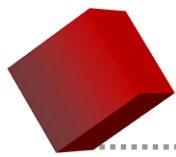
Lemur weight and determinants

The effect of parasites

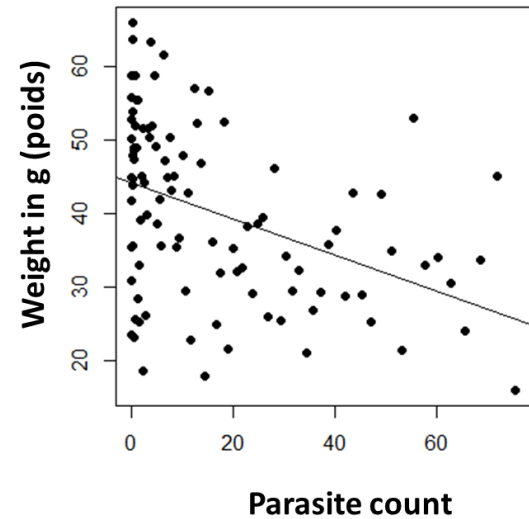
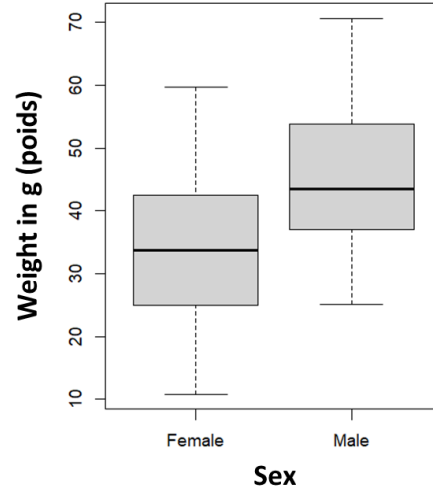
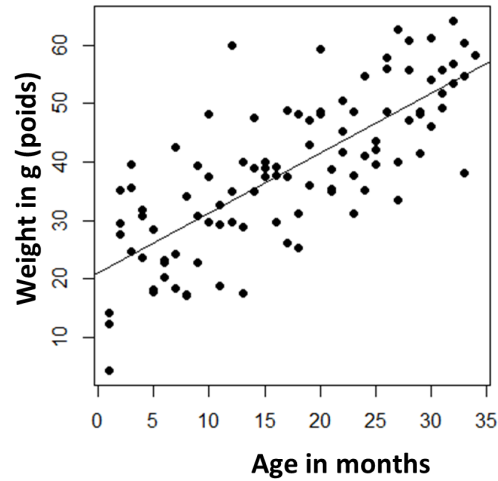


$$\text{Taille} = 45 - 0.3 \times \text{Nb Parasites} + \text{Error}$$

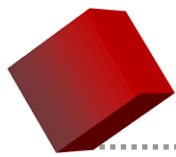




Lemur weight and determinants

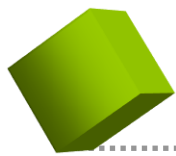


`lm(taille~age+sexe+GIparasites, data)`



Multiple linear regression

- Linear regression with multiple explanatory variables
- To describe the relationship between
 - The response variable, y
 - The explanatory variables, $x = (x_1, x_2, \dots, x_n)$
- The model: $y = \alpha + \beta_1 * x_1 + \dots + \beta_n * x_n + \varepsilon$
with $\varepsilon \sim N(0, \sigma^2)$
- We generally select the model that best fits the data (best explains observed patterns) with the smallest number of variables

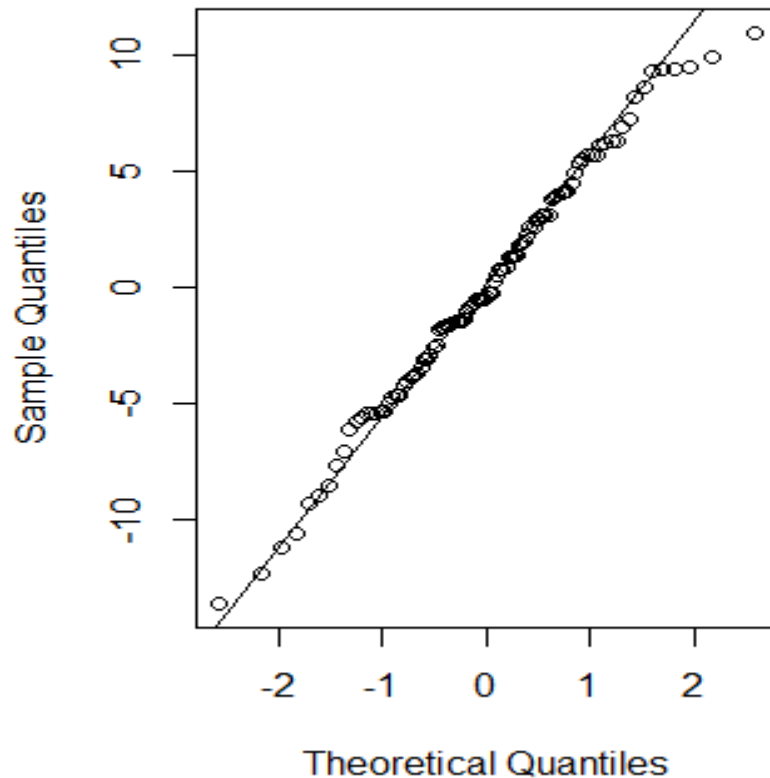


Model validation (case of linear regression $y \sim N(0,1)$)

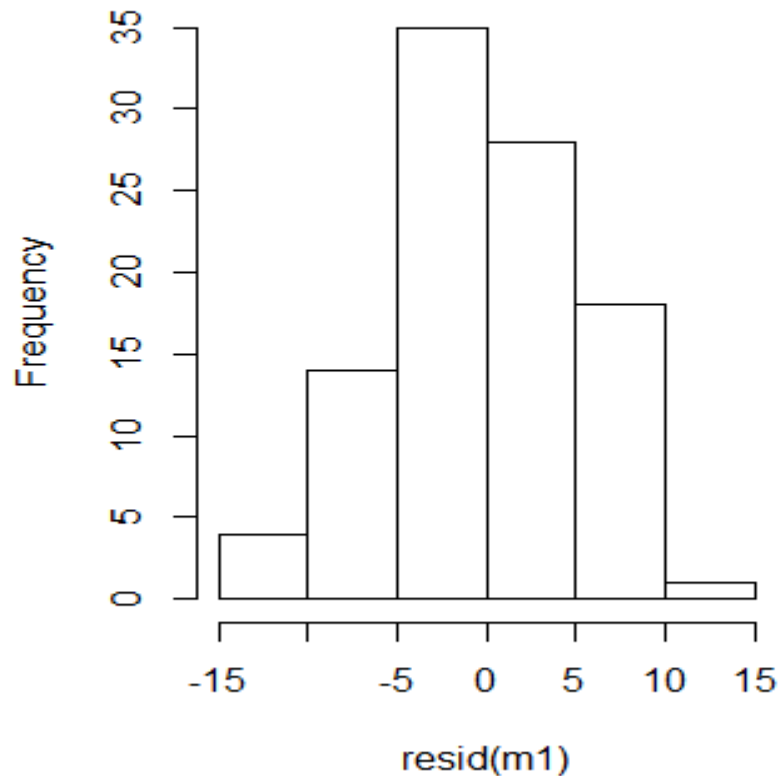
- Check that model assumptions have not been violated

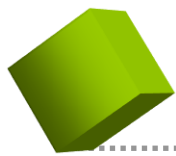
Normality of residuals

Normal Q-Q Plot



Histogram of resid(m1)

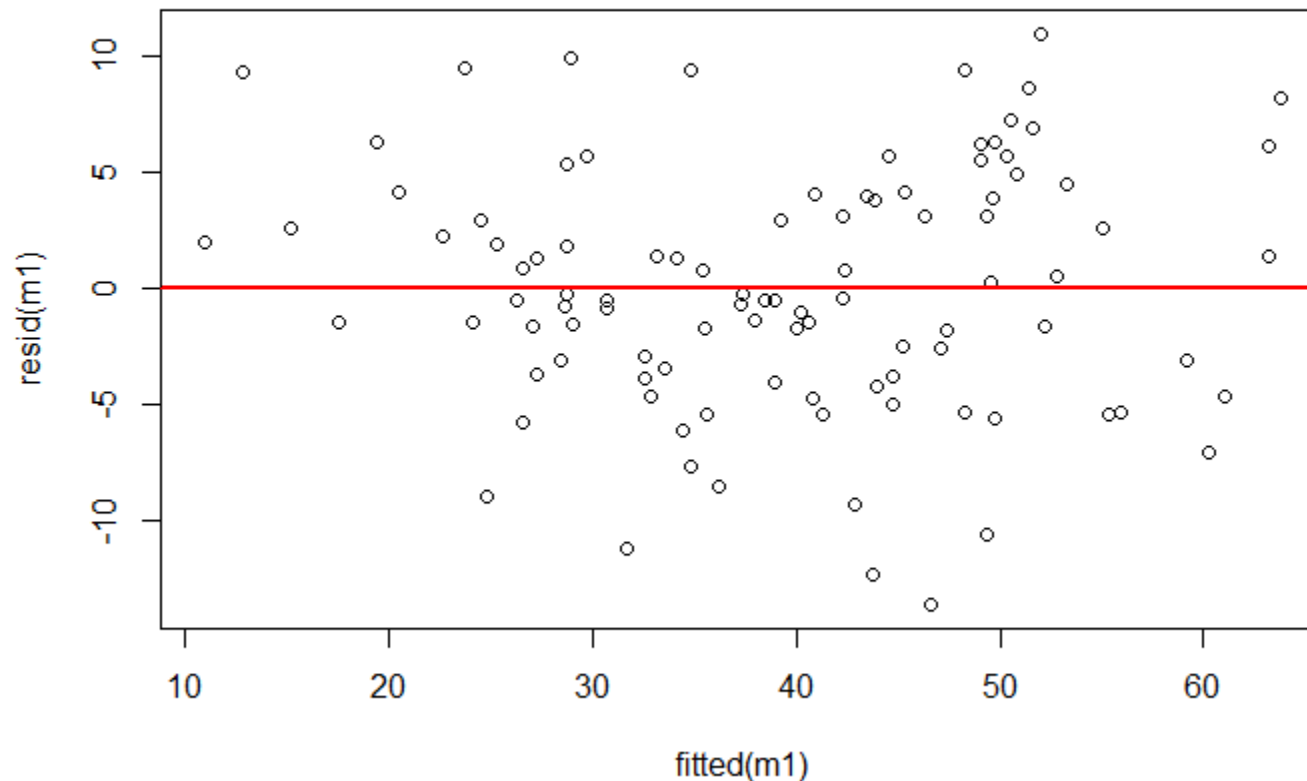




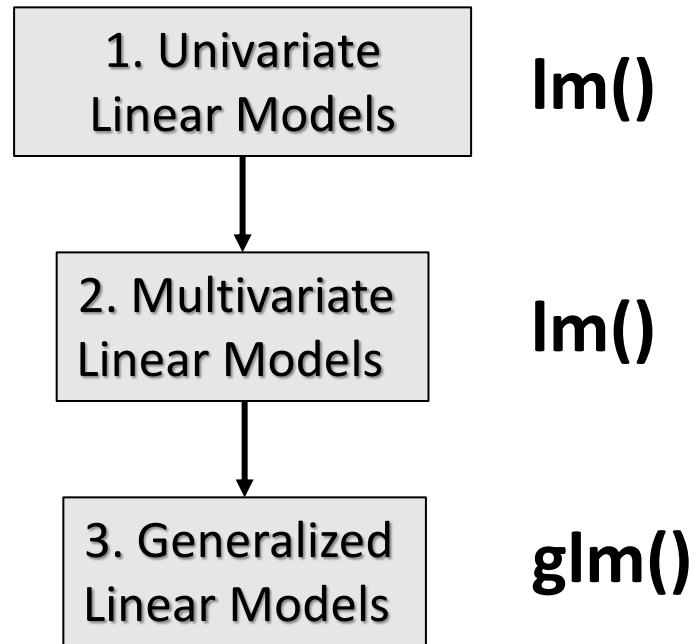
Model validation (case of linear regression $y \sim N(0,1)$)

- Check that model assumptions have not been violated

Homogeneity of residuals



Unfortunately, not all things in
life are normal...

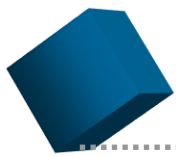


Case when the response variable do **not follow a Normal distribution**

(Dans le cas ou la variable réponse ne suit pas une distribution normale)

Dia ahoana raha tsy manaraka ny “distribution normale” ny “variable response”?

INTRODUCING GENERALIZED LINEAR MODELS

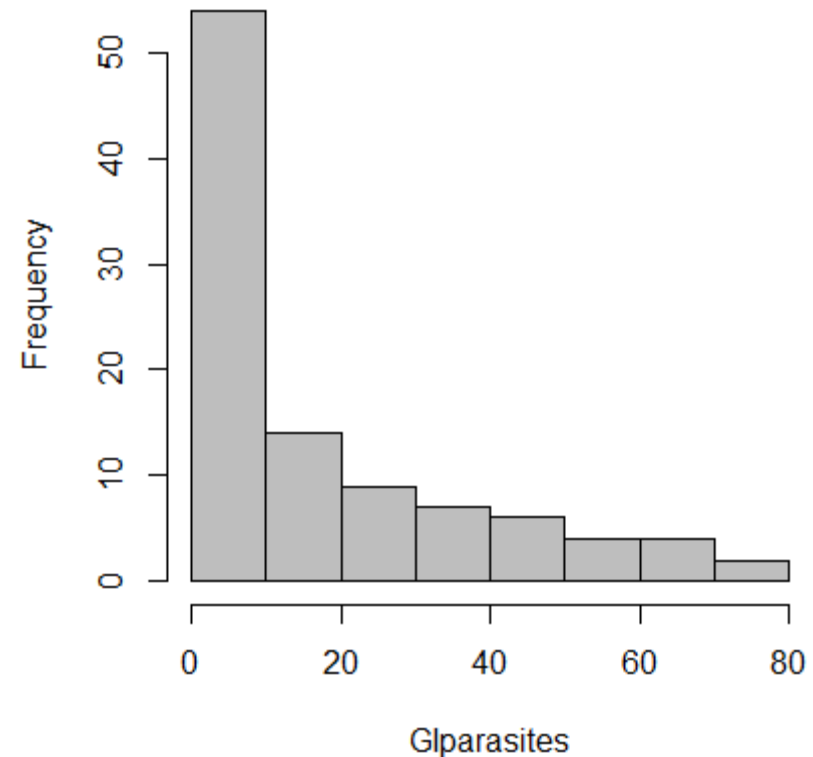


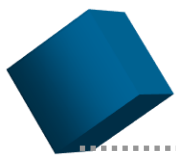
Count data



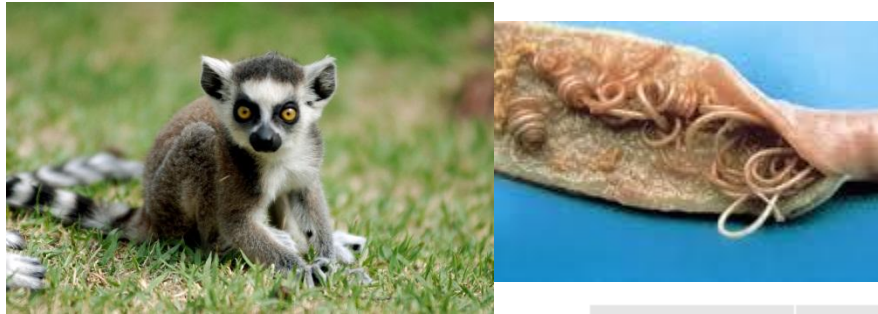
Histogram of Giparasites

- Cannot be negative
- Discrete values
- The lower the values, the « less normal » they generally are.
- Examples:
 - Number of individuals of a species X
 - Number of people with a disease X

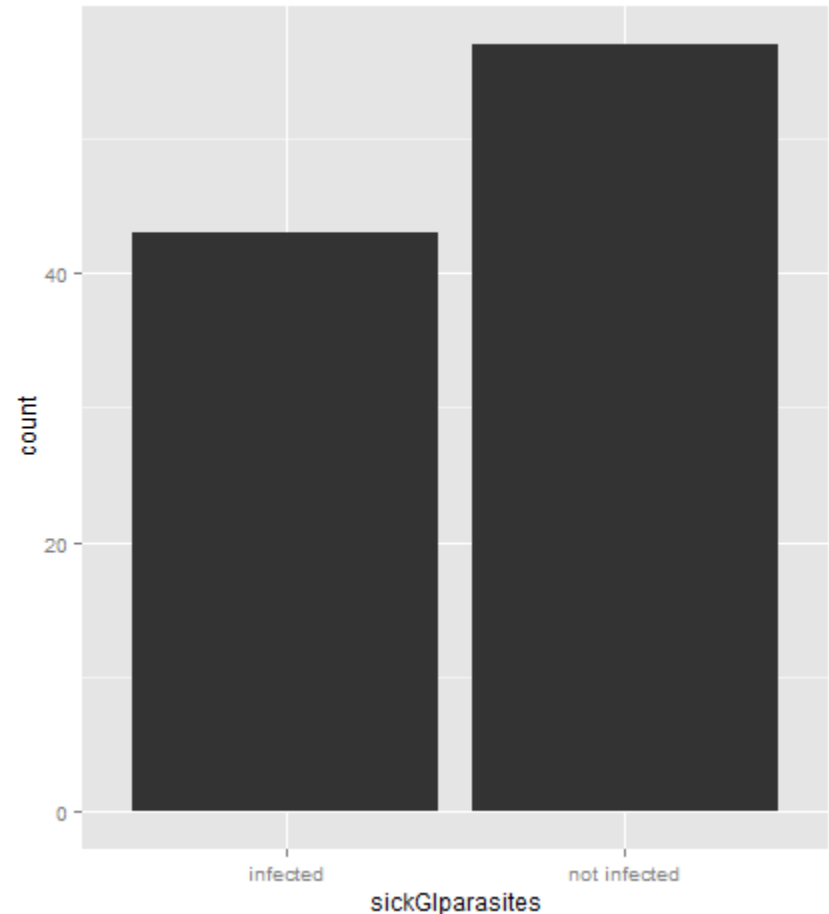


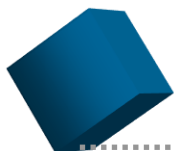


Binary data (events)



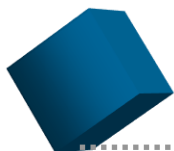
- Values either 1 or 0 (either happened or not happened)
- The outcome variable is the number of successes /failures
- Examples:
 - Presence of a species X
 - Presence of a disease X





Limitations of linear models

- In this type of situations, general linear models are not appropriate because:
 - The range of Y is restricted (e.g. binary, count)
 - The variance of Y depends on the mean
- **Generalized linear models** extend the linear model framework to address both of these issues by using a **linear predictor** and a **link function**



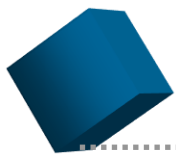
Generalized linear modeling

- One generalization of multiple linear regression. Response, y , predictor variables $x_1, x_2, \dots$. The distribution of Y depends on the X 's through a single linear function, the “linear predictor”

$$\nu = \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_p x_p$$

- A link function describes how the mean $E(Y) = \mu$, depends on the linear predictor ν .

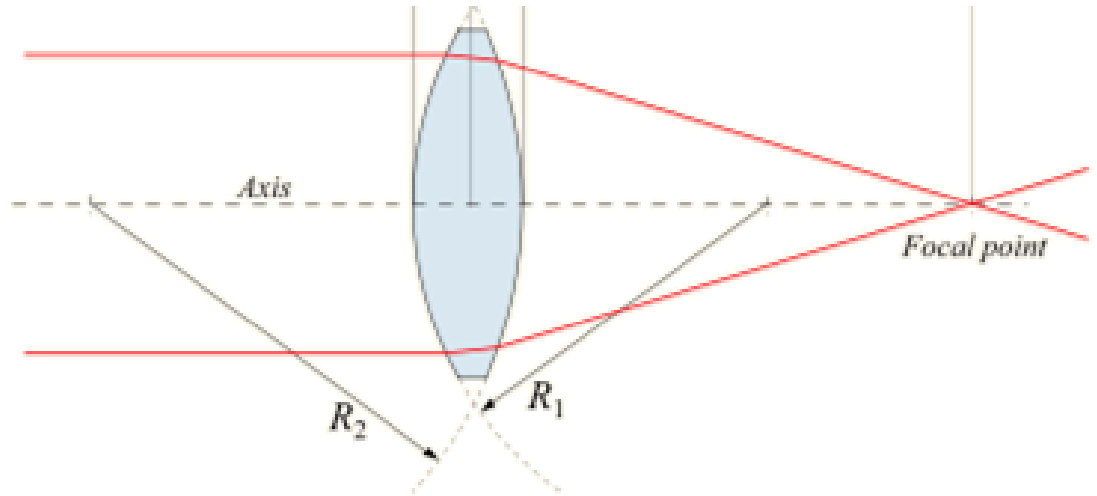
$$\mu = m(\nu), \quad \nu = m^{-1}(\mu) = l(\mu)$$



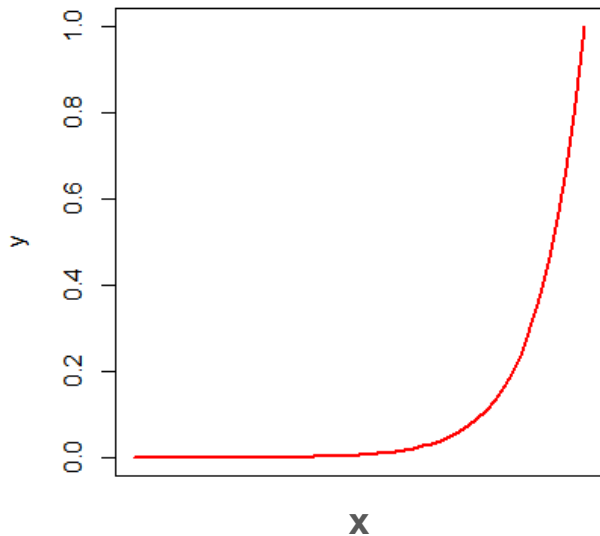
Generalized linear modeling

Most common families and links

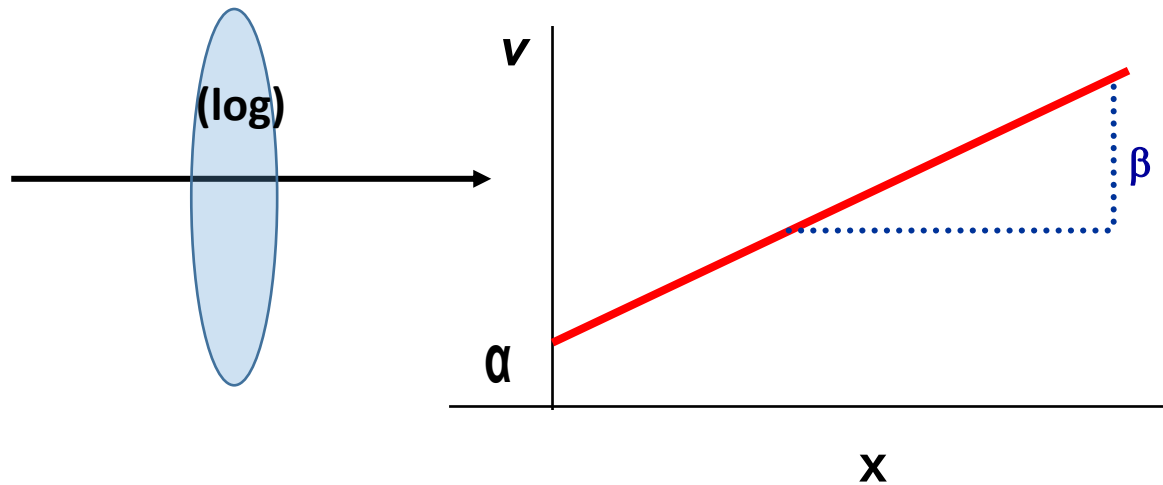
- Gaussian: identity
- Poisson: log
- Binomial: logit
- Negative binomial: log

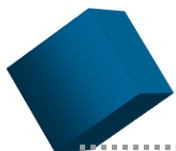


$$Y = e^{(\alpha + \beta x)}$$



$$V = \alpha + \beta x$$

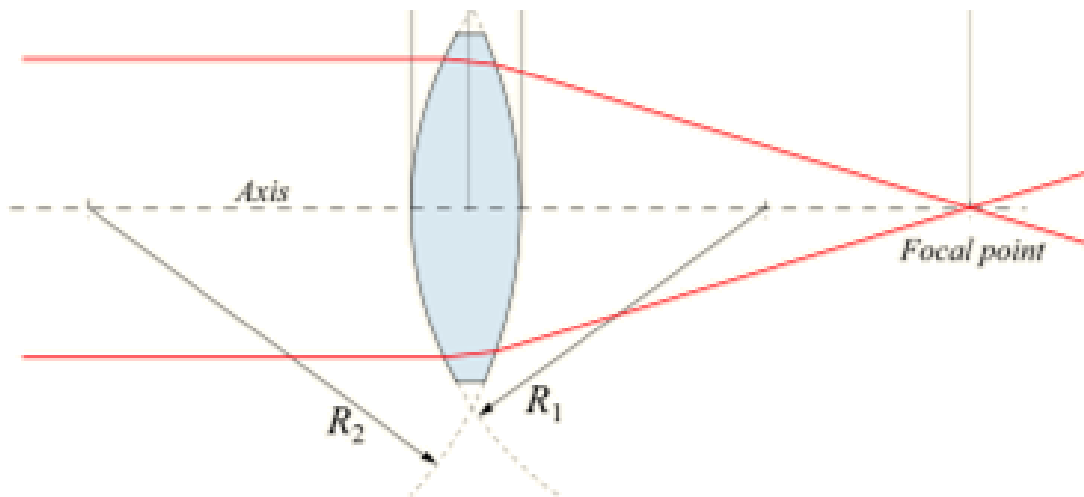




Generalized linear modeling

Most common families and links

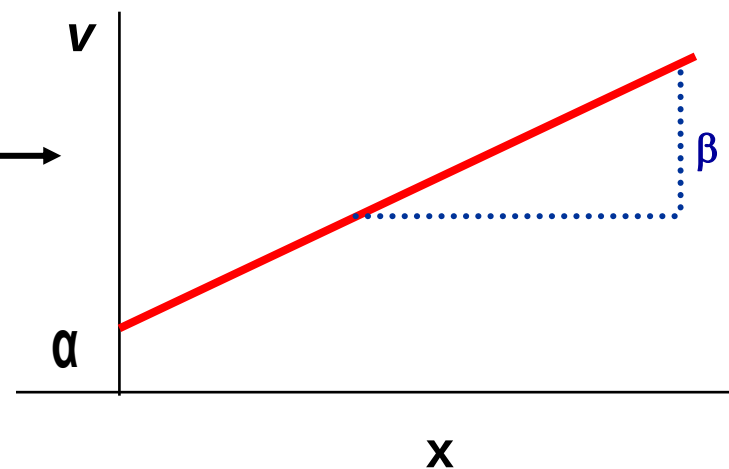
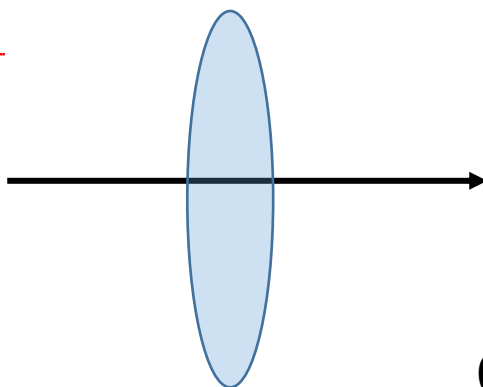
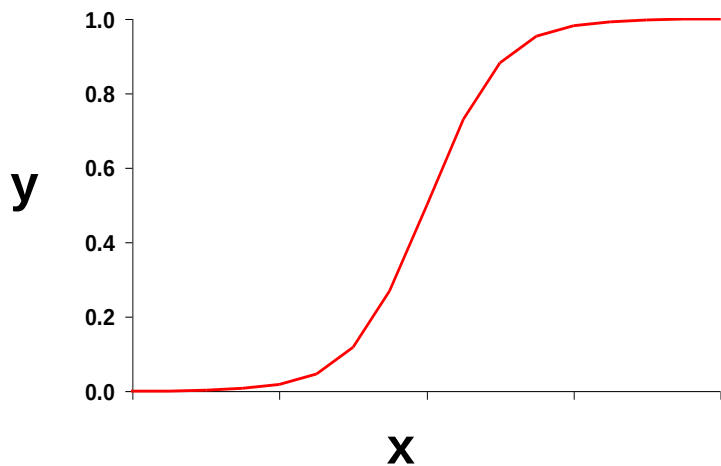
- Gaussian: identity
- Poisson: log
- Binomial: logit
- Negative binomial: log

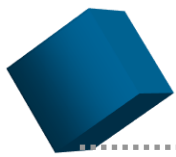


$$P(y|x) = \frac{e^{\alpha + \beta x}}{1 + e^{\alpha + \beta x}}$$

(logit)

$$V = \alpha + \beta x$$





Generalized linear modeling

Most common families and links

- Gaussian: identity
- Poisson: log
- Binomial: logit
- Negative binomial: log

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



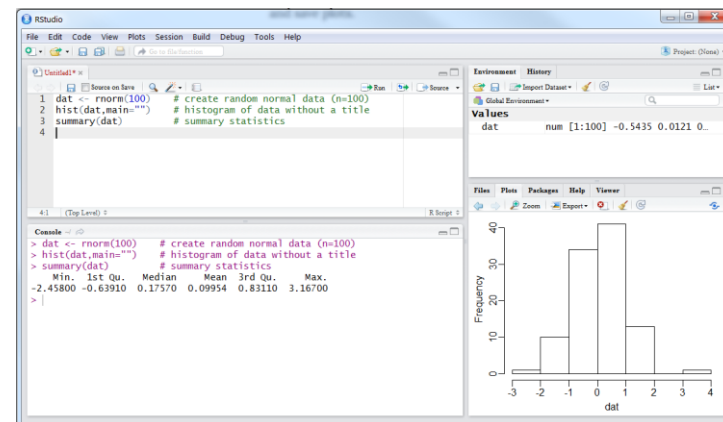
1. Univariate
Linear Models



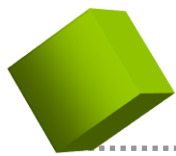
2. Multivariate
Linear Models



3. Generalized
Linear Models



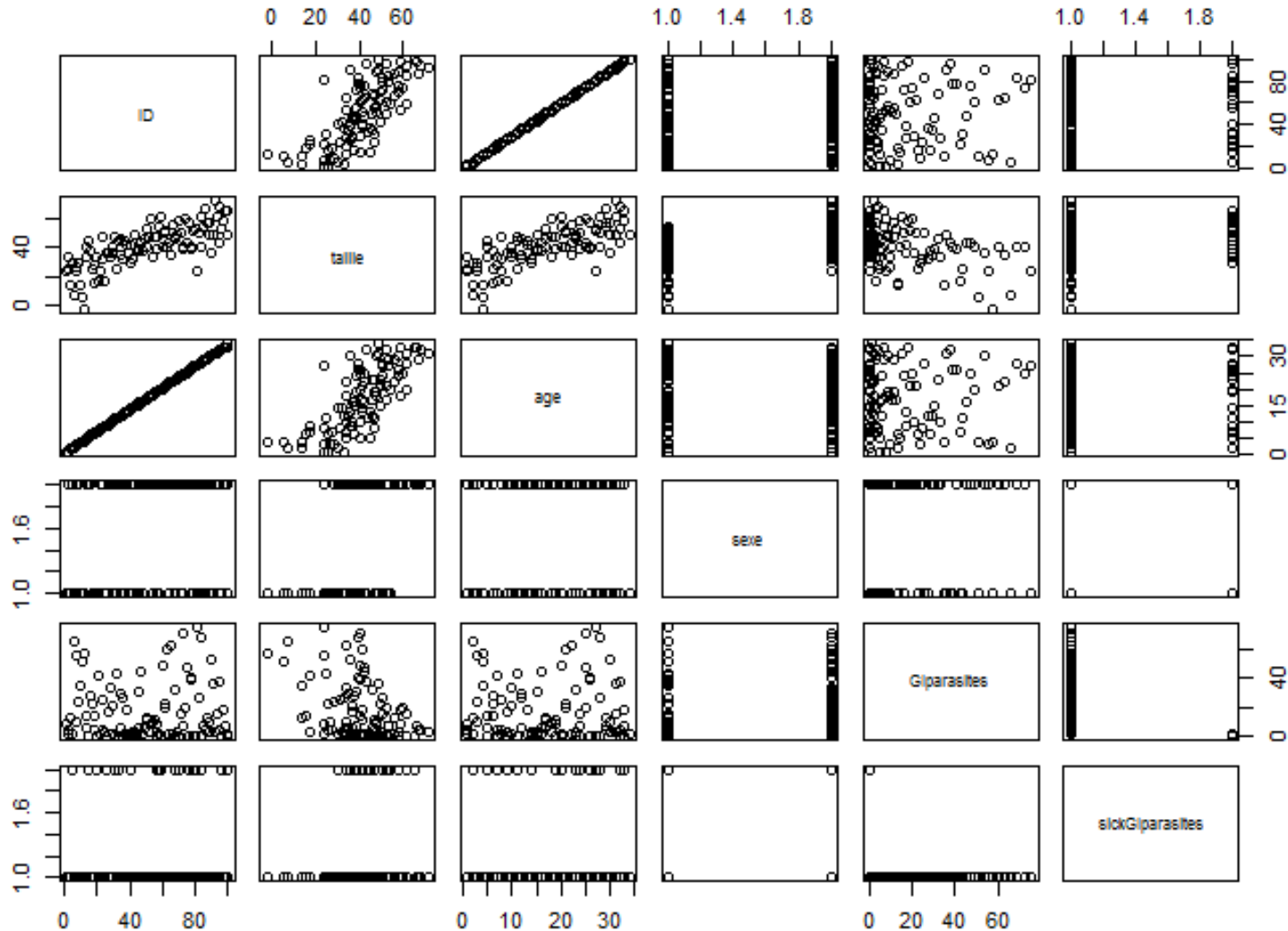
STEPS IN DEVELOPMENT OF STATISTICAL MODELS (TUTORIAL)

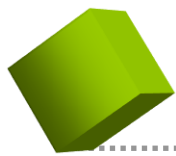


Database construction and descriptive analyses

- Relationships between the variables

`pairs(mydata)`

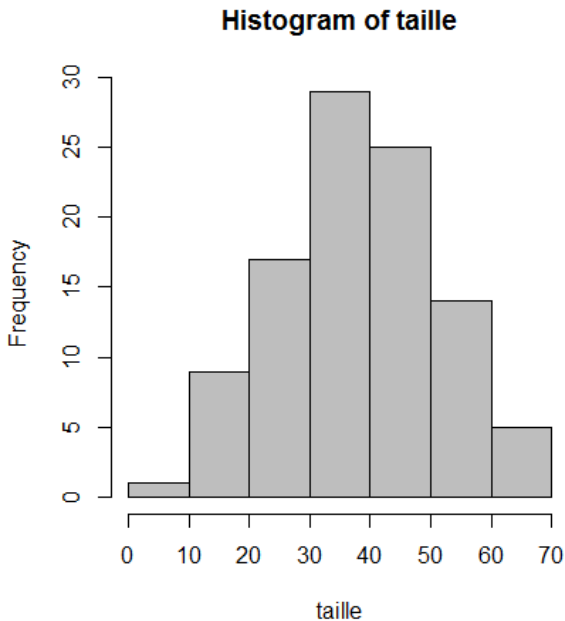




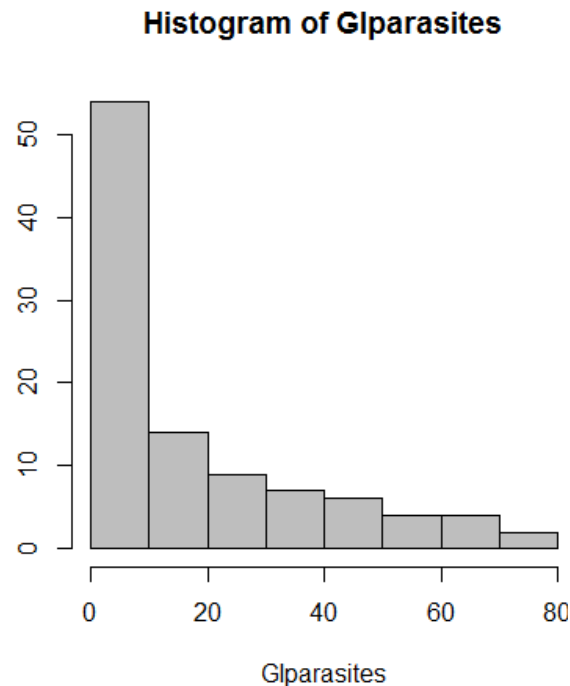
Database construction and descriptive analyses

- Distribution of the response variable
- Distribution of the explanatory variables

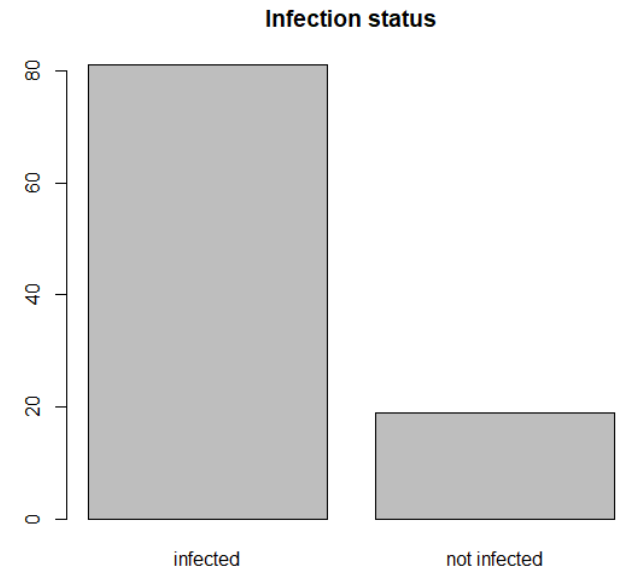
`hist(mydata$var)`

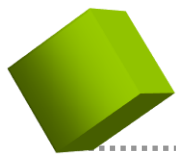


`hist(mydata$var)`



`plot(mydata$var)`



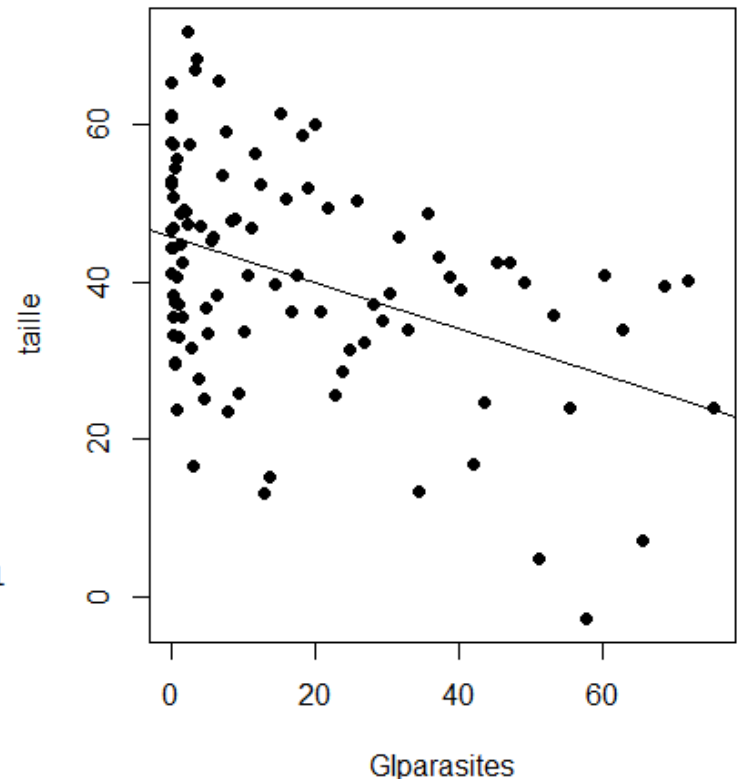


Univariate analyses

- Quantify the strength of the relationship between the response variable and each explanatory variable
- Test the significance of the relationship between the response variable and each explanatory variable

```
Model1 = lm(taille~Giparases, data=mydata)  
summary (Model1)
```

```
call:  
lm(formula = taille ~ Giparases)  
  
Residuals:  
    Min       1Q   Median       3Q      Max   
-31.605  -8.351   1.113   9.901  26.528   
  
Coefficients:  
            Estimate Std. Error t value Pr(>|t|)      
(Intercept)  45.8267    1.7154  26.714 < 2e-16 ***  
Giparases    -0.2927    0.0651  -4.495 1.91e-05 ***  
---  
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
  
Residual standard error: 13.07 on 98 degrees of freedom  
Multiple R-squared:  0.171,    Adjusted R-squared:  0.1625   
F-statistic: 20.21 on 1 and 98 DF,  p-value: 1.906e-05
```





Multivariate analyses

- Quantify the relationship between the response variable and a set of explanatory variables

```
Model1 = lm(taille~age+sexe+GIparasites, data=mydata)
summary (m1)
```

```
call:
lm(formula = taille ~ age + sexe + GIparasites, data = mydata)

Residuals:
    Min       1Q   Median       3Q      Max
-16.9962  -2.6011  -0.1584   3.7331  12.0600

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)  21.94145    1.28143   17.12  <2e-16 ***
age           1.02365    0.05584   18.33  <2e-16 ***
sexeMale     10.88561    1.09295    9.96  <2e-16 ***
GIparasites  -0.29930    0.02652  -11.28  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

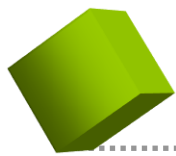
Residual standard error: 5.323 on 96 degrees of freedom
Multiple R-squared:  0.8653,    Adjusted R-squared:  0.8611
F-statistic: 205.5 on 3 and 96 DF,  p-value: < 2.2e-16
```

- Select the set of predictors that best explains the response variable (backwards, forward, stepwise)

```
drop1 (m1)
```

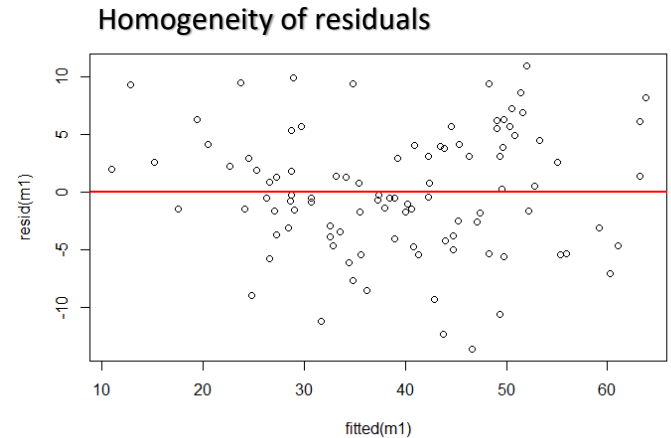
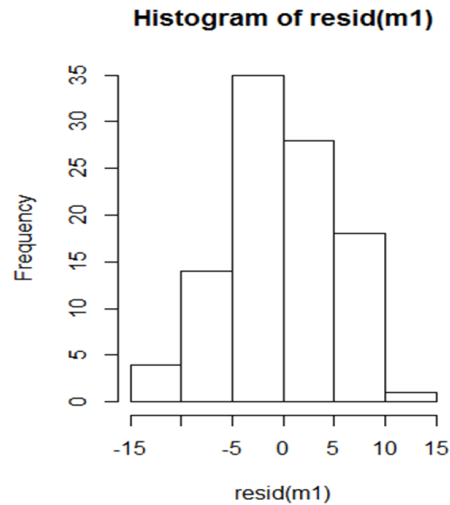
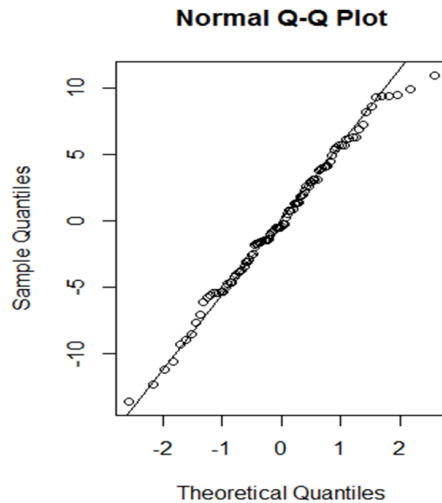
```
add1 (m1)
```

```
step (m1)
```



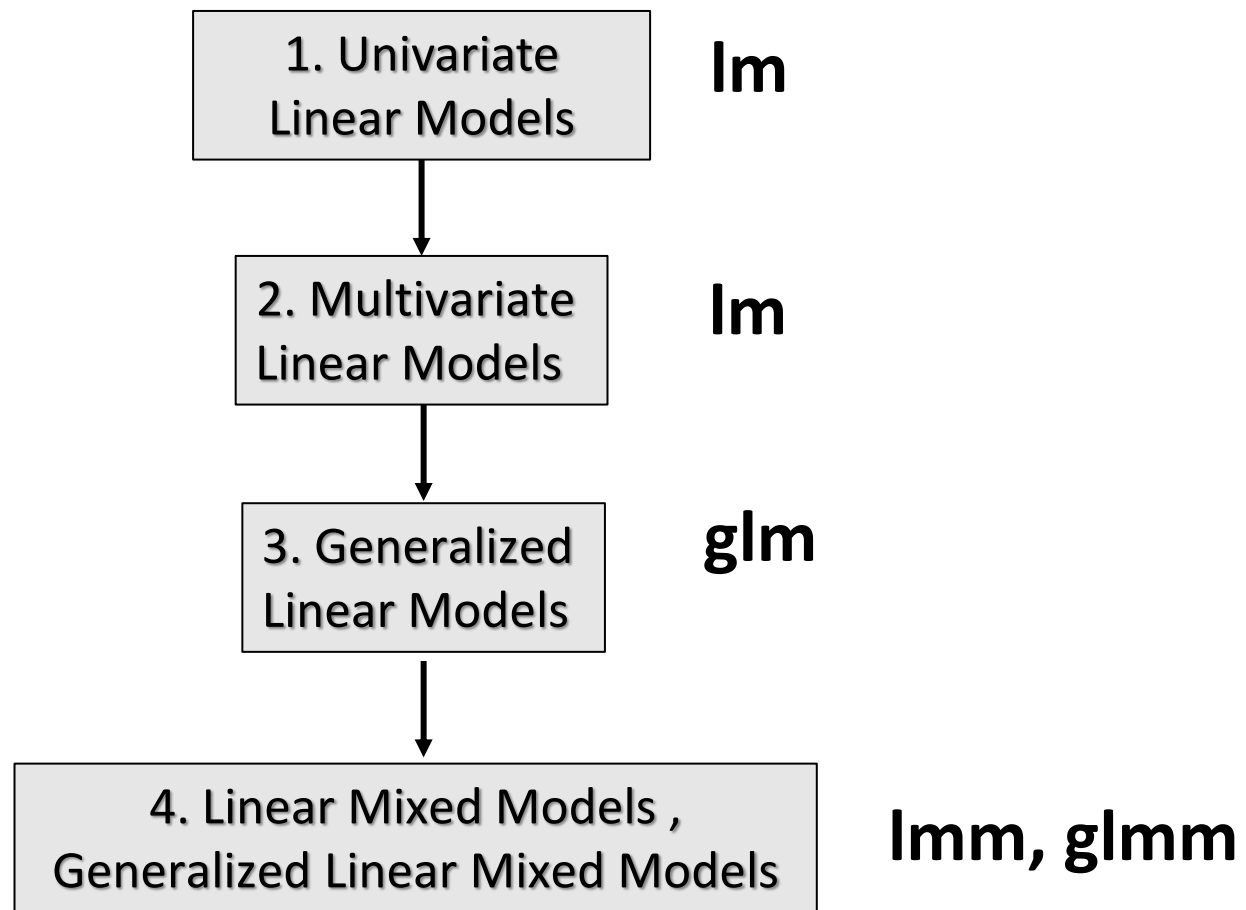
Model validation (case of linear regression $y \sim N(0,1)$)

- Check that model assumptions have not been violated



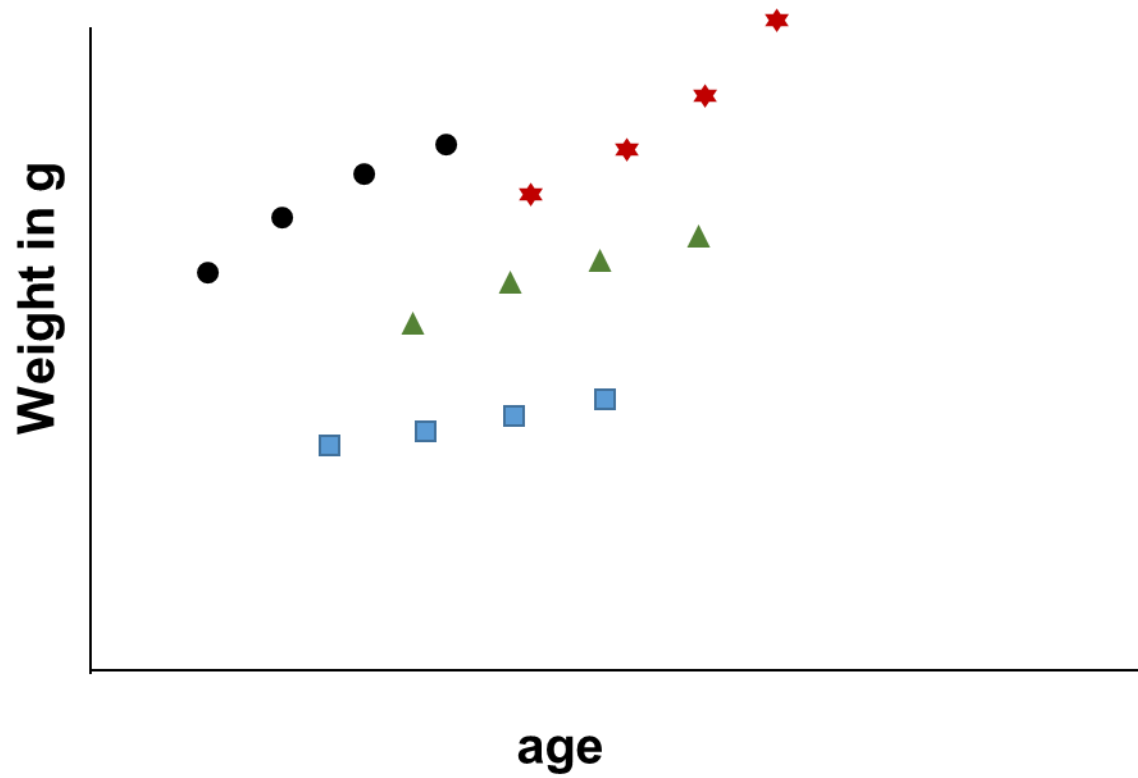
Assumption and limitation of lm, glm

**all observations are considered independent,
or they are often nested**



INTRODUCTION TO GENERALIZED LINEAR MIXED MODELS

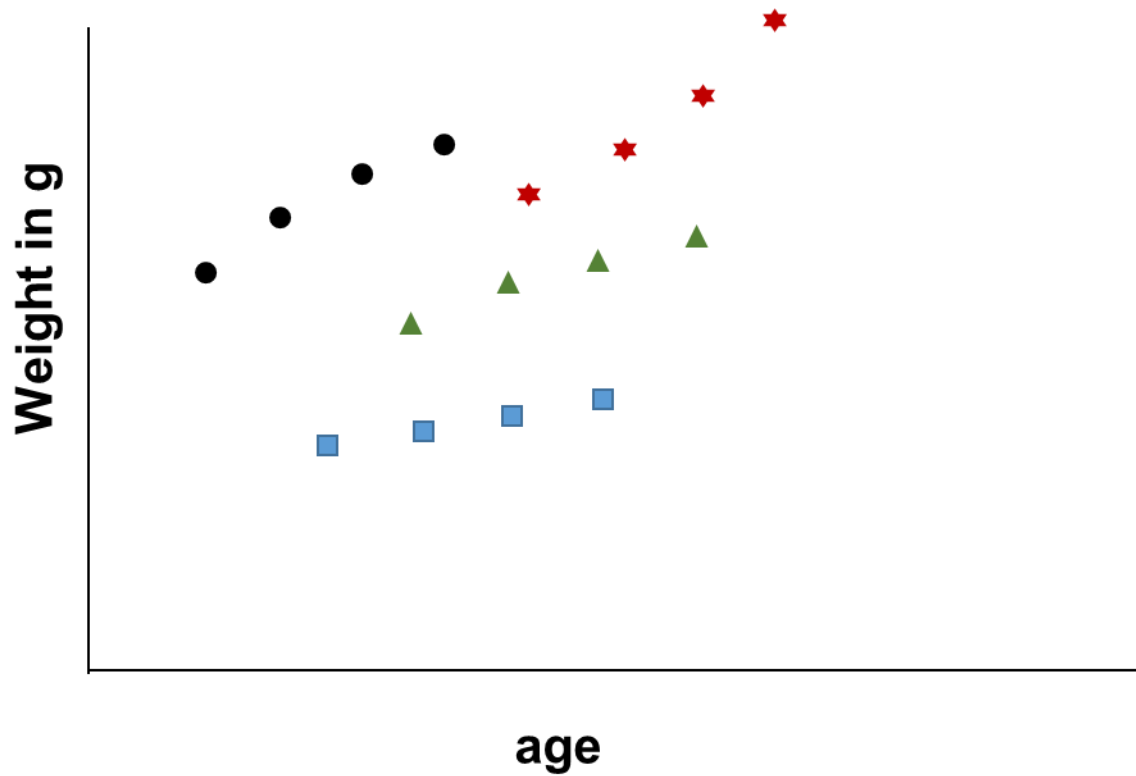
What if...?



Identity (IDs)



What if...?

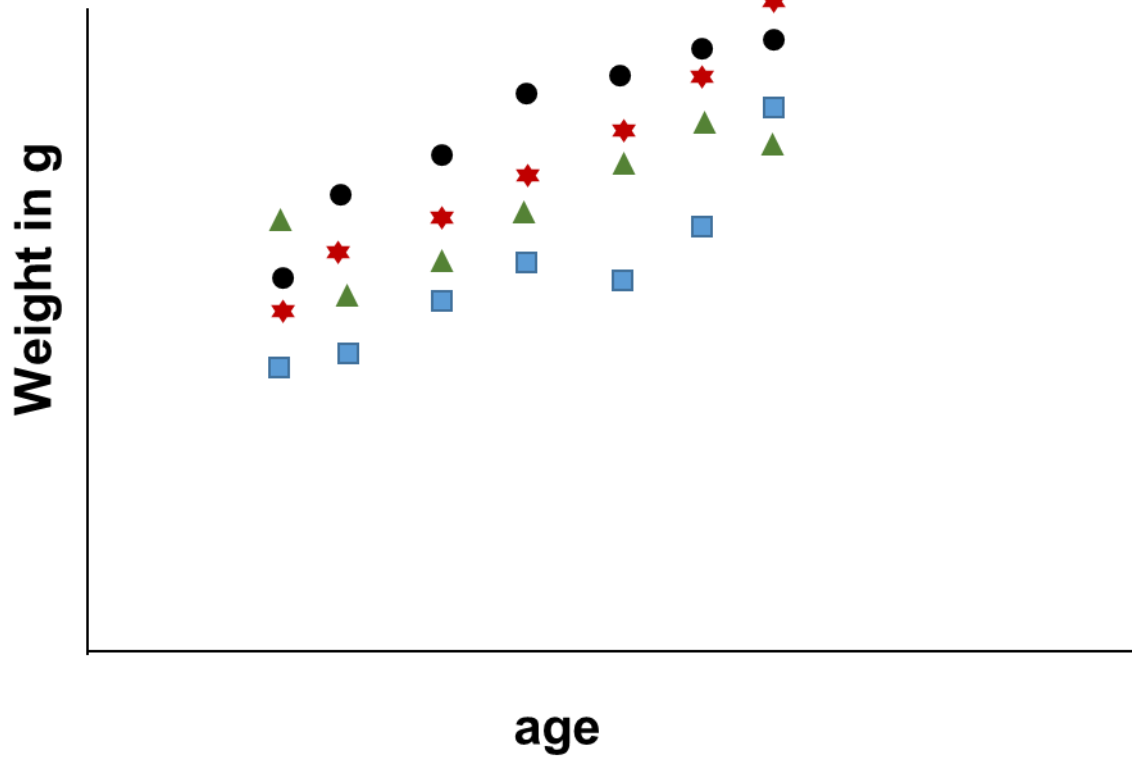


Repeated measure of same individual

Identity (IDs)



Or...?



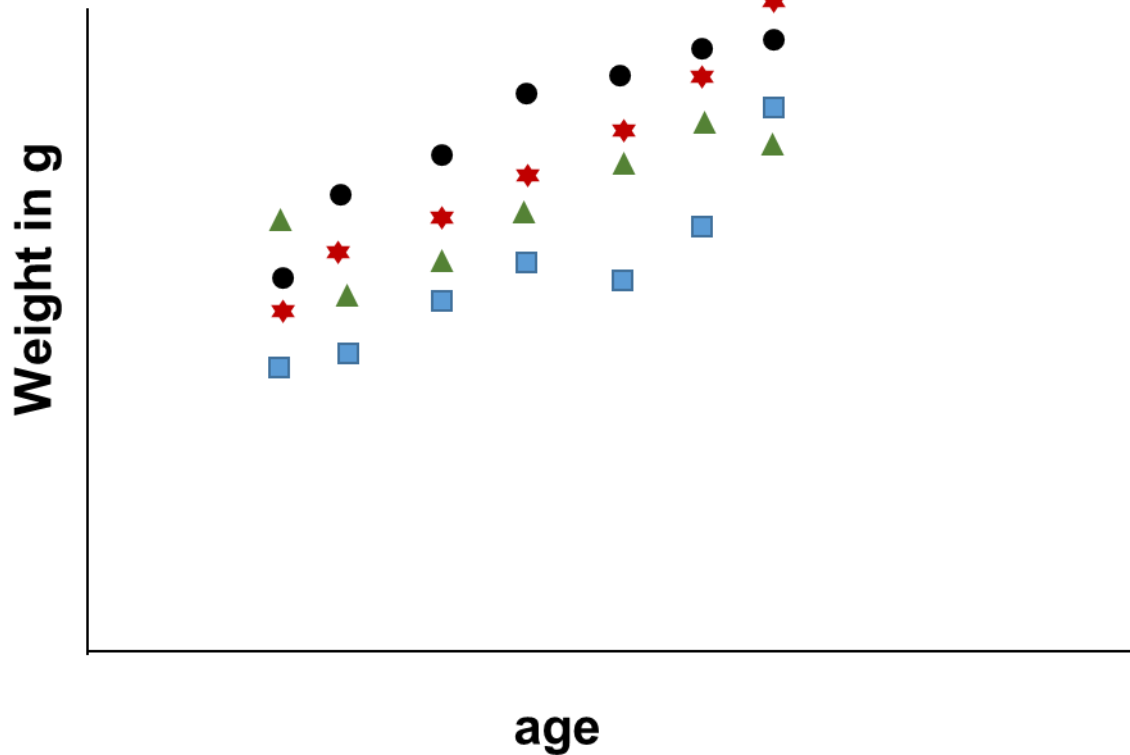
Site

- Andasibe
- ★ Ambohipo
- ▲ 67Ha
- Ambohipo

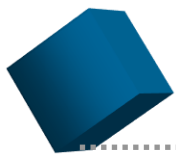
Or...?

Site

- Andasibe
- ★ Ambohipo
- ▲ 67Ha
- Ambohipo



Independent but clustered



Why use LMM, GLMM?

All explanatory variable we have seen in previous models are called variable that have fix effect

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

- Repeated measures
- Temporal correlation
- **Nested data**

$$y = \beta_i X_i + b_j Z_i + \varepsilon$$



Fixed Effects *Random Effects*

Random intercept (clustered measures)

$$y = (\alpha + a_i) + \beta_i X_i + \varepsilon$$

Random Effects *Fixed Effects*

The R function to fit a linear mixed model is lmer() and glmer() for generalized which uses the form

```
fitted.model <- lmer(formula, data=data.frame)
```

Formula : weight ~ age + 1 | sites

```
fitted.model <- glmer(formula, family="model family", data=data.frame)
```

Formula : weight ~ age + 1 | sites

Introduction to Linear Regression

Thank you very much!

Misaotra betsaka!

Hafaliana Christian Ranaivoson

Ecology and Evolution (E&E), University of Chicago
Mention Zoologie et Biodiversité Animale (MZBA), University of Antananarivo
Association Ekipa Fanihy (Efa), Madagascar.



Adapted from
Andrés Garchitorena, PhD (E2M2, 2024)

E²M² Workshop
Andasibe, May 2025