

Lecture 2: Multivariate visualization

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Reminder of session 1

Univariate and bivariate plots

Plots and graphics usually are the starting point for statistical analysis. On the first session we have seen how to plot univariate covariate (e.g. histogram, boxplot, bar plot) and also bivariate analysis (such as scatter plot, or boxplots as a function of a categorical covariate).

What if you have more covariates? It would still be possible to observe data with a 3D plots¹, but one can hardly go beyond this analysis. Before going on, let's check that everyone is able to reproduce the plot below.

¹ Note that R does this well too. You can try the `gg3D` library

```
# Load library for plot
library(ggplot2)

# Load data set (be careful with the path)
immo <- read.csv("./2022.csv")

# Clean data
filter <- !is.na(immo$valeur_fonciere) &
  !is.na(immo$lot1_surface_carrez) &
  immo$valeur_fonciere < 1000000 &
  immo$lot1_surface_carrez < 80

immo <- immo[filter,]

# Plot data
ggplot(immo, aes(x = lot1_surface_carrez,
                 y = valeur_fonciere,
                 color = valeur_fonciere)) +
  geom_point() +
  geom_smooth(method = "lm", color = "purple") +
  xlab("Surface du premier lot en m2") +
  ylab("Valeur foncière") +
  theme_classic() +
  theme(legend.position = "none")
```

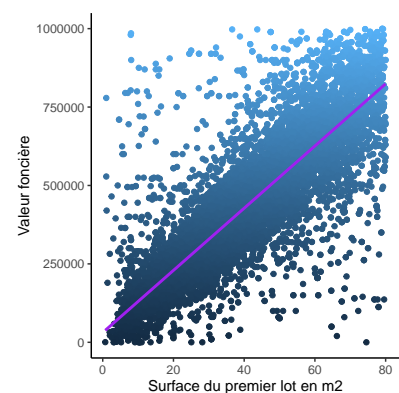


Figure 1: Open data for Paris housing price

Pipe operator in R

In the first lab, you were also asked to plot summary of data, such as the average price per year. Here, we will plot the average price of a flat depending on its number of living room. To do this, a first step is to compute this average price per group. This will allow us to present the pipe operator in R. The pipe operator is denoted `%>%` and corresponds to “chaining” several functions. It means that you invoke multiple method calls. As each method returns an object, you can actually allow the calls to be chained together in a single statement, without needing variables to store the intermediate results.

```
library(lubridate)

## Loading required package: timechange

##
## Attaching package: 'lubridate'

## The following objects are masked from 'package:base':
##
##   date, intersect, setdiff, union

immo$annee <- year(immo$date_mutation)

library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

summary.immo <- immo[immo$nombre_pieces_principales < 6,] %>%
  group_by(nombre_pieces_principales) %>%
  summarise(prix.moyen = mean(valeur_fonciere))

library(dplyr)

summarized.immo <- immo[immo$nombre_pieces_principales < 5,] %>%
  group_by(nombre_pieces_principales) %>%
  summarise(prix.moyen = mean(valeur_fonciere))
```

Then, we observe what has been produced.

```

head(summarized.immo[1:6,])

## # A tibble: 6 x 2
##   nombre_pieces_principales prix.moyen
##   <int> <dbl>
## 1 0 422752.
## 2 1 250833.
## 3 2 410749.
## 4 3 596760.
## 5 4 694314.
## 6 NA NA

library(ggplot2)
# Plot data
ggplot(summary.immo, aes(x = nombre_pieces_principales, y = prix.moyen)) +
  geom_point() +
  geom_line() +
  theme(legend.position = 'bottom') +
  xlab("Nombre de pièces principales") +
  ylab("Valeur foncière") +
  theme_bw()

```

Why is it interesting to visualize covariates jointly?

Let's look at a funny example. Imagine that we generate two variables X_1 and X_2 from normal distributions. We want these variables to be linked (correlated) and such that $X_j \sim \mathcal{N}(0, 1)$. The following chunk performs the simulation. You can take the output data frame and explore the data first with univariate analysis. And then with a bivariate plot. An outlier is in the dataset. Can we recover it?

```

# simulation -- don't need to understand what is going on here
library(MASS) # for simulations
Sigma <- matrix(c(1,0.8,1,0.8),2,2)
simulated_data <- mvrnorm(n = 500, mu = c(0,0), Sigma)
output <- data.frame(simulated_data)
names(output) <- c("X1", "X2")
output[501,] <- c("X1" = 2, "X2" = -2) # outlier step

ggplot(output, aes(x = X1)) +
  geom_histogram(bins = 20,
                fill = "blue",
                alpha = 0.6,
                color = "grey") +
  theme_classic()

```

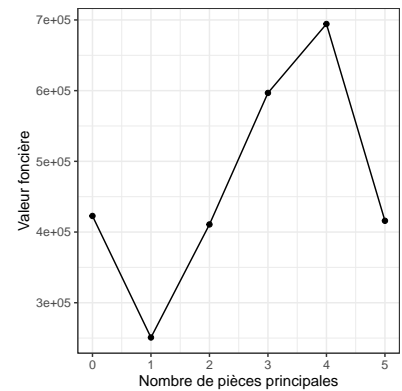


Figure 2: Aggregated data

```
ggplot(output, aes(x = X2)) +
  geom_histogram(bins = 20,
                fill = "magenta",
                alpha = 0.6,
                color = "grey") +
  theme_classic()
```

One can rather plot the two covariates at once.

```
ggplot(output, aes(x = X1, y = X2)) +
  geom_point() +
  theme_classic()
```

The outlier is clearly identifiable on this scatter plot, but not using only the boxplot or any univariate tool. This is to highlight that multivariate analysis will allow us to see high dimensional outliers.

Multivariate analysis will enable us to summarize highly dimensional data into a simpler 2D plot. This will rely on factorial analysis, where the aim is to summarize a large dataframe. The exact method chosen depends on the nature of the covariates. For example if all covariates are continuous, then Principal Component Analysis (PCA) can be used, but if the covariates are qualitative, then the method is rather correspondence analysis.

Principal Component Analysis (Work from home)

We recommend to watch the videos² from François Husson about PCA. Below we recall the main principles.

- **Context** Principal Component Analysis (usually the shortname is PCA but you can also find ACP in French) focuses on typical data you can find in several domains: *observations* (or individus) in rows, and *variables* in column. Note that the PCA focuses on *quantitative* variables (for example age, or price, but not color or sex). For example we can study the average temperature depending on cities. In that case cities are rows, and in column the average temperature per month.
- **Typical question an ACP answers** A typical question you may ask on your data is: how much the different observations are close to one another considering the variables? (remember that everything you will conclude depends on these variables that you added in your initial model) You can also see PCA as a way to find a low-dimensional representation that captures the “essence” of high-dimensional data

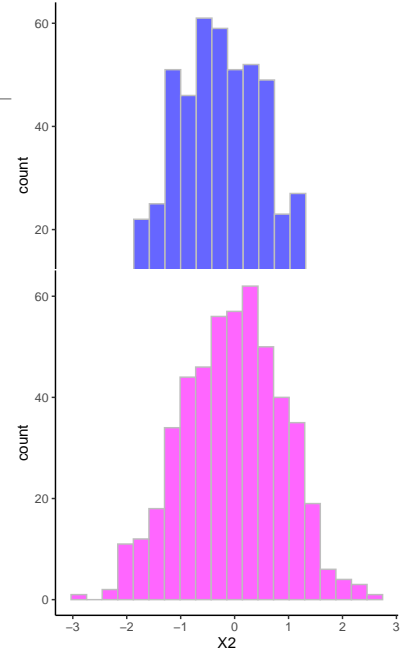


Figure 4: X2

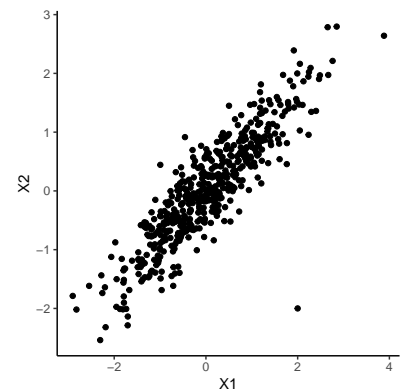


Figure 5: X1 and X2 on a scatter plot

² Here is the link.

- **What can you interpret from data?** The PCA will group similar individuals together. Information are also learned on variables, with the correlated variables (meaning that you have a linear link between two variables), and also which variables synthetize the most the observations, or which variables bring different information.

Correspondance Analysis (CA)

Typical situation is when you have two qualitative covariates, and in particular a data counting how many times the occurrences co-occur in the data. CA proposes you to visualize how the two covariates are associated.

A few historical information: - First applications in the 60's by Jean-Paul Bensécri (a whole French community on these kind of analysis) - One of the first application is on the characters from the play *Phèdre*.

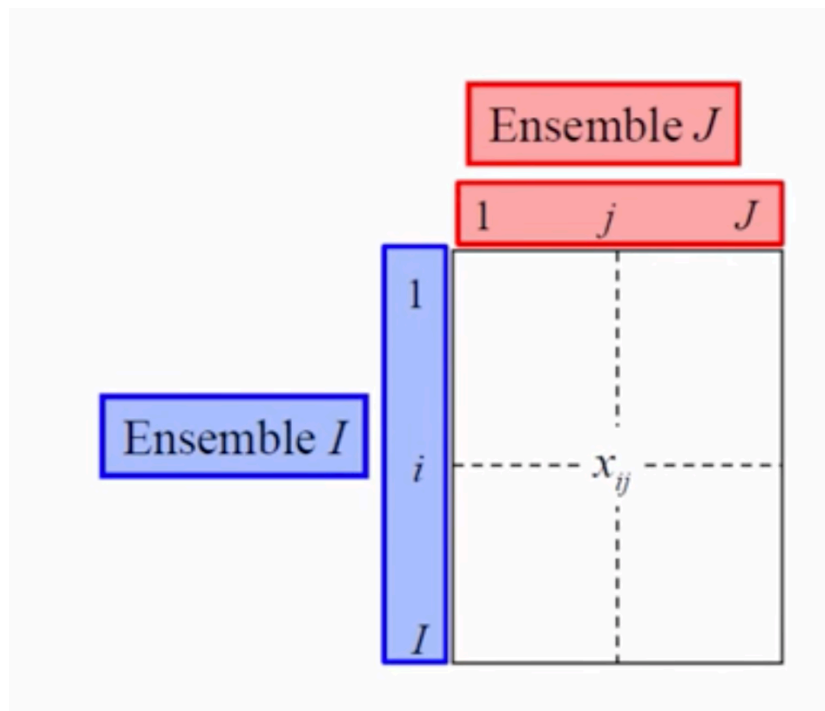


Figure 6: Typical data used: contingency table

Principle

For a concrete example, let's count the number of nobel prize in each domain for the height countries of G8. Is there a specialty depending on the countries? Below we show the example.

- $x_{i,j}$ corresponding to the number of individuals with both characteristics i from column I and j from column J (see Figure).
- Here n , was all the nobel prizes, and the count data are summarizing all these.
- Note that, $\sum_{i=1}^8 \sum_{j=1}^7 x_{i,j} = n$.

	Chimie	Econ.	Lettres	Médecine	Paix	Physique	Math
Allemagne	24	1	8	18	5	24	1
Canada	4	3	2	4	1	4	1
France	8	3	11	12	10	9	11
GB	23	6	7	26	11	20	4
Italie	1	1	6	5	1	5	1
Japon	6	0	2	3	1	11	3

From this contingency table, it is possible to go to the probability table, noting that

$$f_{i,j} = \frac{x_{i,j}}{n}.$$

The correspondance analysis will work on this table. For those who have been doing a lecture in statistics or probabilities, one has the joint probability to observe both i and j simulatenously,

$$\mathbb{P}[X_i = i, X_j = j] = f_{i,j}.$$

We will also look at marginal probabilities, that are,

$$f_i = \sum_{j=1}^J f_{i,j}$$

and

$$f_j = \sum_{i=1}^I f_{i,j}.$$

Now, the key idea is to observe how much each column (or row) is different than the marginal probability. Two events are said to be independent if

$$\mathbb{P}[A, B] = \mathbb{P}[A] \cdot \mathbb{P}[B].$$

“The joint probability is equal to the product of the marginal probabilities.”

Now, the idea is to say that data are not independent if the observed joint probabilities $f_{i,j}$ (observed) are different than the product of the marginal probabilities $f_i \cdot f_j$ (i.e. independence model). Maybe

this reminds you the χ^2 test to compare the observed values with the theoretical values.

$$\chi_{\text{obs}}^2 = \sum_{i=1}^1 \sum_{j=1}^J \frac{(\text{obs. num.} - \text{theor. num.})^2}{\text{theor. num.}} = \sum_{i=1}^1 \sum_{j=1}^J \frac{(n_{f_{ij}} - n_{f_i} f_j)^2}{n_{f_i} f_j} = n\Phi^2.$$

The higher Φ^2 , the higher the deviation from independence. In other words Φ^2 is the strength of the relation shipt (it does not depend on n). In this class, we do not focus on whether the link is statistically different from 0. But we use it to plot the data and understand the link.

In other words, we are comparing each column profile, with its marginal one. This may seem a bit abstract, so let's look at our running example. First, this is the frequency table.

	Chimie	Econ.	Lettres	Médecine	Paix	Physique	Math
Allemagne	0.1983471	0.0082645	0.0661157	0.1487603	0.0413223	0.1983471	0.0082645
Canada	0.0330579	0.0247934	0.0165289	0.0330579	0.0082645	0.0330579	0.0082645
France	0.0661157	0.0247934	0.0909091	0.0991736	0.0826446	0.0743802	0.0909091
GB	0.1900826	0.0495868	0.0578512	0.2148760	0.0909091	0.1652893	0.0330579
Italie	0.0082645	0.0082645	0.0495868	0.0413223	0.0082645	0.0413223	0.0082645
Japon	0.0495868	0.0000000	0.0165289	0.0247934	0.0082645	0.0909091	0.0247934

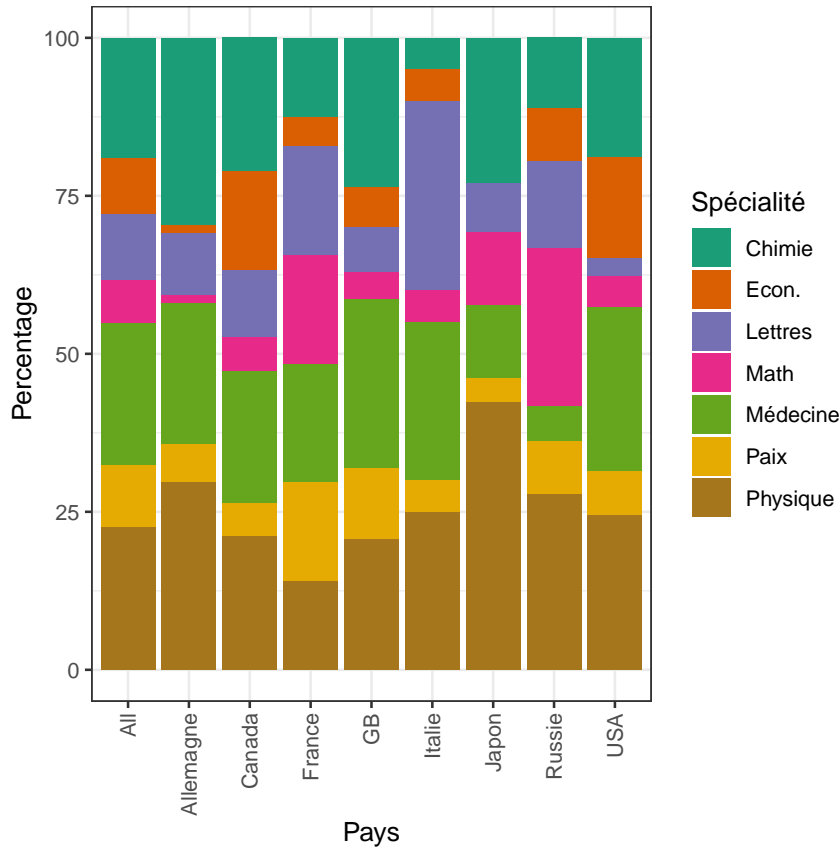
Now, if we are willing to observe marginal versus conditional distribution.

```
library(ggplot2)
```

```
library(tidyr)
```

```
nobel.freq %>%
```

```
  pivot_longer(cols = c("Chimie", "Econ.", "Lettres", "Médecine", "Paix", "Physique", "Math"), names_to = "Spécialité") +
  ggplot(aes(x = Pays, y = Percentage, fill = Spécialité, group = Spécialité)) +
  geom_bar(stat = "identity") +
  theme_bw() +
  scale_fill_brewer(palette="Dark2") +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



See how Italy has a relative more important number in letters. This is the contrary for the USA. The goal is to compare the distribution of Nobel Prize winners.

Each country has a profile. Denoting i the country, for each country we have

$$\left(\frac{f_{i,1}}{f_i}, \frac{f_{i,2}}{f_i}, \dots, \frac{f_{i,J}}{f_i} \right).$$

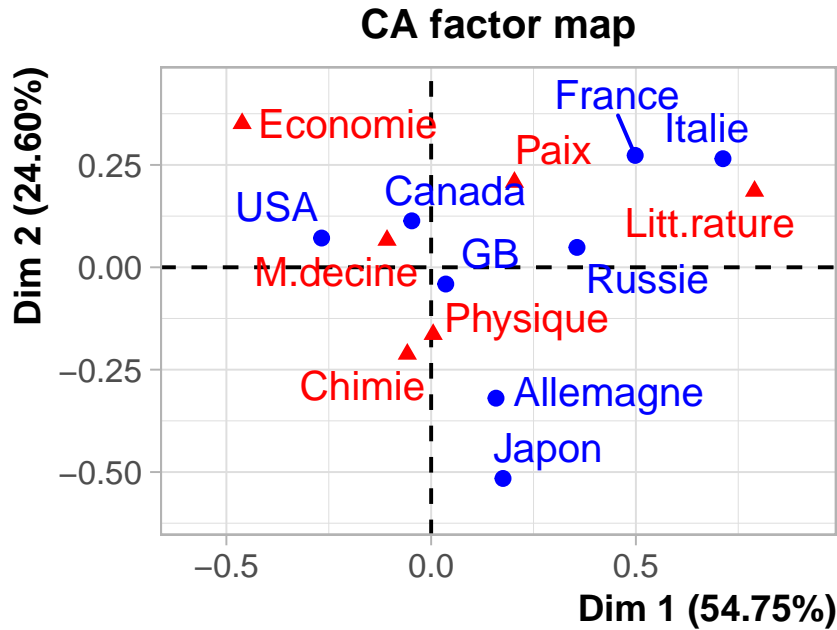
Then, in this space we compute the distance between each vector to the mean vector. In other words, each country is represented by a vector in J dimensional space (i.e. the number of majors).

$$d_{\chi^2}^2(i, i') = \sum_{j=1}^J \frac{1}{f_{.j}} \left(\frac{f_{ij}}{f_i} - \frac{f_{i'j}}{f_{i'}} \right)^2.$$

The same can be done in the other direction.

If there is independence, all the vectors are more or less confounded with the mean vector. You can imagine a cloud of points and that all the points are very close to the center of gravity. It can be shown that the inertia of the cloud (i.e. how much it spreads) is linked with Φ^2 . Also, it can be shown that rows and columns have the same role.

Then, the process is the same than the PCA, finding plan on which the cloud is the most dispersed.



- UK is super close to the center of gravity (look back to the previous plot);
- Italy and France seems really different;
- Red points are spread out and we could put the assumption that the first axis contrasts science and other categories, while the second axis contrasts natural science with economic science.

Application on Majors and studies

The example of this part are data from the French universities, and in particular how many students are in which specialty, degree (L3, M2, PhD), and their gender. Typical questions we will answer are “Are they major in which students are similar or different?” “Is there an association between the major and the gender?” and so on.

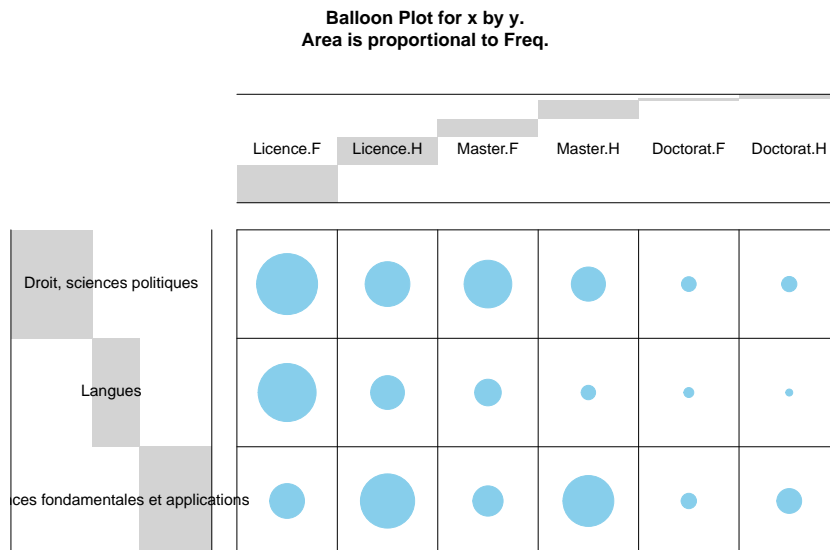
```
univ <- read.csv("./universite.csv", header = TRUE, row.names = 1, skip = 0, sep = ";")
head(univ)
```

##	Licence.F	Licence.H	Master.F	Master.H
## Droit, sciences politiques	69373	37317	42371	21693
## Sciences économiques, gestion	38387	37157	29466	26929
## Administration économique et sociale	18574	12388	4183	2884
## Lettres, sciences du langage, arts	48691	17850	17672	5853
## Langues	62736	21291	13186	3874
## Sciences humaines et sociales	94346	41050	43016	20447
##	Doctorat.F	Doctorat.H	Total.F	Total.H

```
## Droit, sciences politiques          4029      4342 115773 63352
## Sciences économiques, gestion      1983      2552 69836 66638
## Administration économique et sociale 0          0 22757 15272
## Lettres, sciences du langage, arts  4531      2401 70894 26104
## Langues                             1839       907 77761 26072
## Sciences humaines et sociales      7787      6972 145149 68469
##                                     Licence Master Doctorat Total
## Droit, sciences politiques          106690 64064 8371 179125
## Sciences économiques, gestion      75544 56395 4535 136474
## Administration économique et sociale 30962 7067 0 38029
## Lettres, sciences du langage, arts  66541 23525 6932 96998
## Langues                             84027 17060 2746 103833
## Sciences humaines et sociales      135396 63463 14759 213618
```

Be careful as some columns are summing other columns. Here we focus on the variable with gender and level. Before launching the CA, one can still have another view of the data.

```
library("gplots")
# 1. convert the data as a table
dt <- as.table(as.matrix(univ[c(1,5,8),1:6]))
# 2. Graph
balloonplot(t(dt), xlab="", ylab="",
            label = FALSE, show.margins = FALSE)
```



```
library(FactoMineR)
analysis.ca <- CA(univ, col.sup = 7:12, graph = FALSE)
```

The object `analysis.ca` contains all the results, and automatically it output one plot if `graph = TRUE`.

```
# summary only for the first 2 dimensions
```

```
summary(analysis.ca, ncp = 2, dim = 2, nb.dec = 1, nbelements = 2)
```

```
##
```

```
## Call:
```

```
## CA(X = univ, col.sup = 7:12, graph = FALSE)
```

```
##
```

```
## The chi square of independence between the two variables is equal to 170789.2 (p-value = 0 ).
```

```
##
```

```
## Eigenvalues
```

```
##           Dim.1 Dim.2 Dim.3 Dim.4 Dim.5
## Variance      0.1  0.0  0.0  0.0  0.0
## % of var.     70.7 15.5 10.9  2.6  0.2
## Cumulative % of var. 70.7 86.2 97.1 99.8 100.0
```

```
##
```

```
## Rows (the 2 first)
```

```
##                                     Iner*1000  Dim.1
## Droit, sciences politiques         |      5.7 | -0.1
## Sciences économiques, gestion     |      9.8 |  0.2
##                                     ctr cos2  Dim.2
## Droit, sciences politiques         |  1.4 0.3 |  0.1
## Sciences économiques, gestion     |  3.9 0.5 |  0.0
##                                     ctr cos2
## Droit, sciences politiques         |  2.9 0.1 |
## Sciences économiques, gestion     |  0.1 0.0 |
```

```
##
```

```
## Columns (the 2 first)
```

```
##                                     Iner*1000  Dim.1
## Licence.F                          |     48.3 | -0.4
## Licence.H                          |     24.3 |  0.2
##                                     ctr cos2  Dim.2
## Licence.F                          | 39.7 1.0 |  0.0
## Licence.H                          | 11.5 0.6 | -0.2
##                                     ctr cos2
## Licence.F                          |  2.3 0.0 |
## Licence.H                          | 37.5 0.4 |
```

```
##
```

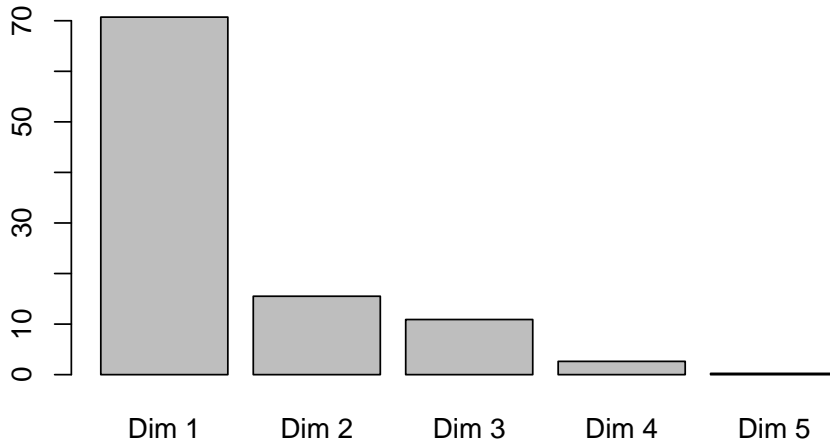
```
## Supplementary columns (the 2 first)
```

```
##                                     Dim.1 cos2
## Total.F                             | -0.3 1.0 |
## Total.H                             |  0.4 1.0 |
##                                     Dim.2 cos2
## Total.F                             |  0.0 0.0 |
## Total.H                             | -0.1 0.0 |
```

As you can see, the independence test is rejected:

The chi square of independence between the two variables is equal to 170789.2 (p-value = 0). So one can conclude on the existence of associations between some majors and the level-gender covariates.

```
barplot(analysis.ca$eig[,2],names=paste("Dim",1:nrow(analysis.ca$eig)))
```



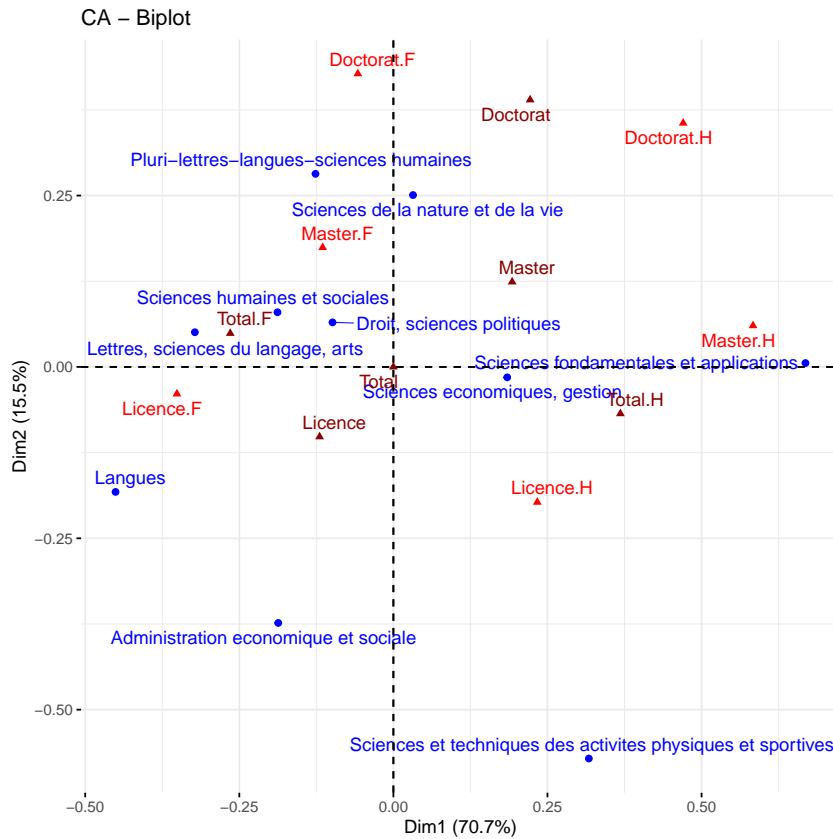
Looking at the percentage of inertia, one can say that the three first dimensions summarizes 97% of the total inertie (almost equal to variability), so only analyzing those three dimensions is enough.

We use another library (but this is not mandatory of course) to plot the results.

```
library("factoextra")
```

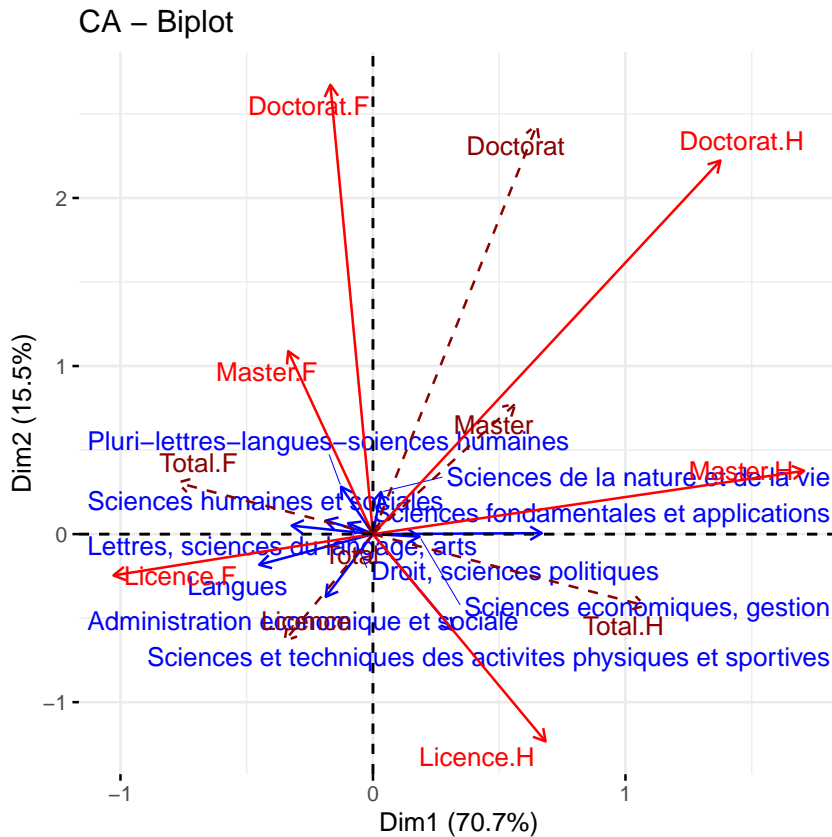
By the way, this package can also allow you to visualize the data before the analysis.

```
# repel= TRUE to avoid text overlapping (slow if many point)
fviz_ca_biplot(analysis.ca, repel = TRUE)
```



You can also draw what is called an *asymmetric biplot*.

```
fviz_ca_biplot(analysis.ca,
               map = "rowprincipal", arrow = c(TRUE, TRUE),
               repel = TRUE)
```



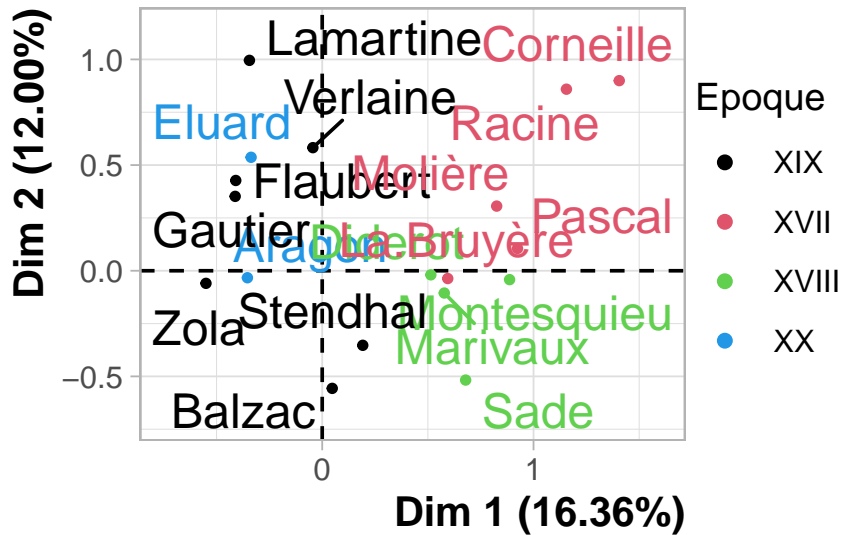
- Recall that two majors are close if they attract similar profiles (here gender and studies length)
- Langues, Lettres, Science du Language: attract women in licence.
- Women and men seems to be separated along the first axis, men on the right, women on the left. Second dimension is more related to studies length: from licence at the bottom and PhD in the upper part.
- Major on the left are mostly occupied by women, and on the right by men.
- It is not always easy to interpret the axis (here it seems possible): in general, you can focus on which entities are close or not.

Correspondance analysis and text data analysis

```
books = read.table("./litterature.csv", header=TRUE, row.names=1, sep=";", check.names=FALSE, quote="\")
res.ca = CA(books, quanti.sup=1, quali.sup=2:3, graph = F)

plot(res.ca,
      invis = c("col", "quali"),
      hab=2, cex=1.2,
      title="", cex.axis=1.2,
```

```
cex.lab=1.2,
palette=palette(c("black","green3","blue","darkred","orange")),shadow=TRUE)
```



```
summary(res.ca)
```

```
##
## Call:
## CA(X = books, quanti.sup = 1, quali.sup = 2:3, graph = F)
##
## The chi square of independence between the two variables is equal to 2768153 (p-value = 0 ).
##
## Eigenvalues
##          Dim.1  Dim.2  Dim.3  Dim.4  Dim.5  Dim.6  Dim.7
## Variance      0.285  0.209  0.190  0.174  0.163  0.145  0.116
## % of var.     16.359  11.997  10.912  9.961  9.369  8.334  6.685
## Cumulative % of var. 16.359  28.356  39.268  49.230  58.598  66.932  73.617
##          Dim.8  Dim.9  Dim.10  Dim.11  Dim.12  Dim.13  Dim.14
## Variance      0.108  0.082  0.063  0.052  0.042  0.037  0.027
## % of var.      6.187  4.708  3.612  3.011  2.392  2.136  1.545
## Cumulative % of var. 79.804  84.512  88.124  91.135  93.528  95.664  97.209
##          Dim.15  Dim.16  Dim.17
## Variance      0.023  0.019  0.007
## % of var.      1.302  1.068  0.421
## Cumulative % of var. 98.511  99.579  100.000
##
## Rows (the 10 first)
##          Iner*1000  Dim.1  ctr  cos2  Dim.2  ctr
## Aragon            | 148.513 | -0.354  6.620  0.127 | -0.033  0.079
## Balzac            | 138.225 |  0.047  0.144  0.003 | -0.557  27.062
## Corneille         | 160.564 |  1.406  26.364  0.468 |  0.900  14.724
```

```

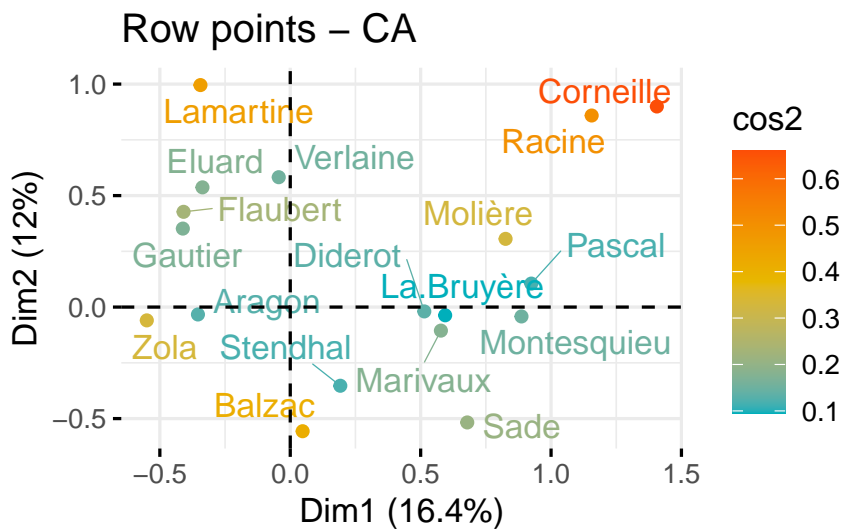
## Diderot          |  52.930 |  0.514  2.458  0.132 | -0.019  0.005
## Eluard           |  46.859 | -0.337  0.822  0.050 |  0.537  2.853
## Flaubert         |  68.150 | -0.409  2.632  0.110 |  0.427  3.914
## Gautier          | 115.002 | -0.412  3.945  0.098 |  0.352  3.929
## La.Bruyère       |  26.605 |  0.594  0.879  0.094 | -0.037  0.005
## Lamartine        | 153.156 | -0.344  2.634  0.049 |  0.996 30.028
## Marivaux         |  84.755 |  0.578  5.150  0.173 | -0.106  0.234
##
##              cos2    Dim.3    ctr    cos2
## Aragon         0.001 |  0.119  1.122  0.014 |
## Balzac         0.409 | -0.330 10.465  0.144 |
## Corneille      0.192 | -0.661  8.731  0.103 |
## Diderot        0.000 |  0.116  0.187  0.007 |
## Eluard         0.127 |  0.214  0.500  0.020 |
## Flaubert       0.120 |  0.162  0.621  0.017 |
## Gautier        0.071 |  0.236  1.941  0.032 |
## La.Bruyère     0.000 |  0.121  0.055  0.004 |
## Lamartine      0.410 |  0.312  3.250  0.040 |
## Marivaux       0.006 | -0.225  1.172  0.026 |
##
## Columns (the 10 first)
##
##              Iner*1000    Dim.1    ctr    cos2    Dim.2    ctr    cos2
## accord          |  0.913 |  0.571  0.039  0.123 |  0.349  0.020  0.046 |
## affaire         |  1.566 |  0.089  0.011  0.021 | -0.461  0.412  0.550 |
## âge            |  0.287 |  0.049  0.002  0.018 | -0.068  0.005  0.033 |
## ah              |  0.777 | -0.663  0.021  0.078 | -0.073  0.000  0.001 |
## air             |  1.387 | -0.324  0.290  0.596 | -0.078  0.023  0.035 |
## allemagne      |  1.221 | -0.434  0.017  0.039 | -0.074  0.001  0.001 |
## allemand       |  1.689 | -0.663  0.046  0.078 | -0.073  0.001  0.001 |
## amant          |  1.876 |  0.637  0.297  0.452 |  0.036  0.001  0.001 |
## âme            |  3.739 |  0.417  0.372  0.284 |  0.350  0.359  0.201 |
## ami            |  1.136 |  0.164  0.057  0.144 | -0.260  0.197  0.362 |
##
##              Dim.3    ctr    cos2
## accord        -0.223  0.009  0.019 |
## affaire       -0.254  0.137  0.166 |
## âge           0.168  0.031  0.202 |
## ah            0.273  0.005  0.013 |
## air          -0.080  0.026  0.036 |
## allemagne     0.314  0.013  0.021 |
## allemand      0.273  0.012  0.013 |
## amant        -0.359  0.142  0.144 |
## âme          -0.032  0.003  0.002 |
## ami           0.046  0.007  0.011 |
##
## Supplementary continuous variable

```



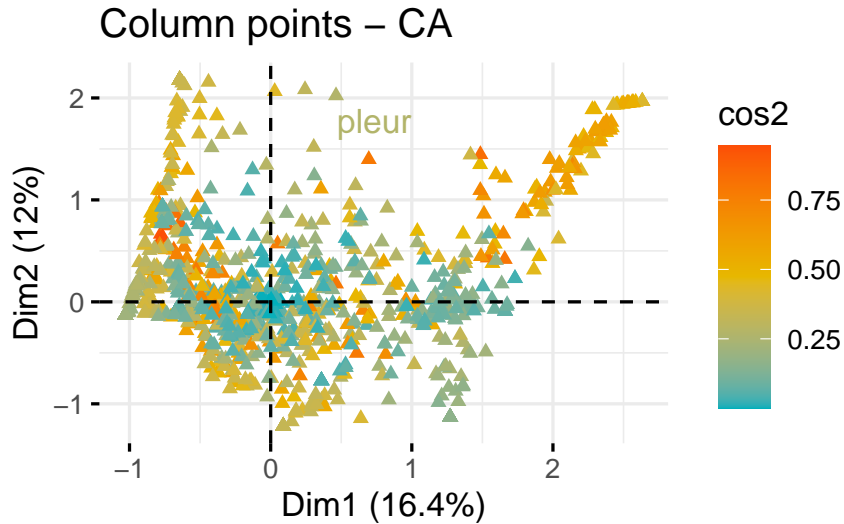
```
##          Dim.1  cos2   Dim.2  cos2   Dim.3  cos2
## Décès      | -0.863  0.744 | -0.125  0.016 |  0.143  0.020 |
##
## Supplementary categorical variables
##          Dim.1  cos2  v.test   Dim.2  cos2  v.test
## Epoque.XIX   | -0.234  0.406 -352.769 | -0.043  0.014 -65.062 |
## Epoque.XVII  |  1.110  0.633 441.534 |  0.584  0.175 232.322 |
## Epoque.XVIII |  0.655  0.442 345.097 | -0.226  0.053 -119.064 |
## Epoque.XX    | -0.352  0.148 -201.647 |  0.036  0.002  20.444 |
## Courant.Classicisme |  1.110  0.633 441.534 |  0.584  0.175 232.322 |
## Courant.Lumières |  0.655  0.442 345.097 | -0.226  0.053 -119.064 |
## Courant.Naturalisme | -0.550  0.328 -308.118 | -0.060  0.004 -33.409 |
## Courant.Réalisme |  0.008  0.000   6.522 | -0.361  0.364 -290.476 |
## Courant.Romantisme | -0.379  0.132 -184.274 |  0.666  0.408 324.154 |
## Courant.Surréalisme | -0.352  0.148 -201.647 |  0.036  0.002  20.444 |
##          Dim.3  cos2  v.test
## Epoque.XIX   -0.112  0.093 -169.013 |
## Epoque.XVII  -0.400  0.082 -159.092 |
## Epoque.XVIII  0.537  0.296 282.736 |
## Epoque.XX     0.131  0.020  74.800 |
## Courant.Classicisme -0.400  0.082 -159.092 |
## Courant.Lumières  0.537  0.296 282.736 |
## Courant.Naturalisme -0.216  0.050 -120.744 |
## Courant.Réalisme  -0.227  0.144 -182.761 |
## Courant.Romantisme  0.273  0.069 132.923 |
## Courant.Surréalisme  0.131  0.020  74.800 |
```

```
fviz_ca_row(res.ca, col.row = "cos2",
            gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
            repel = TRUE)
```

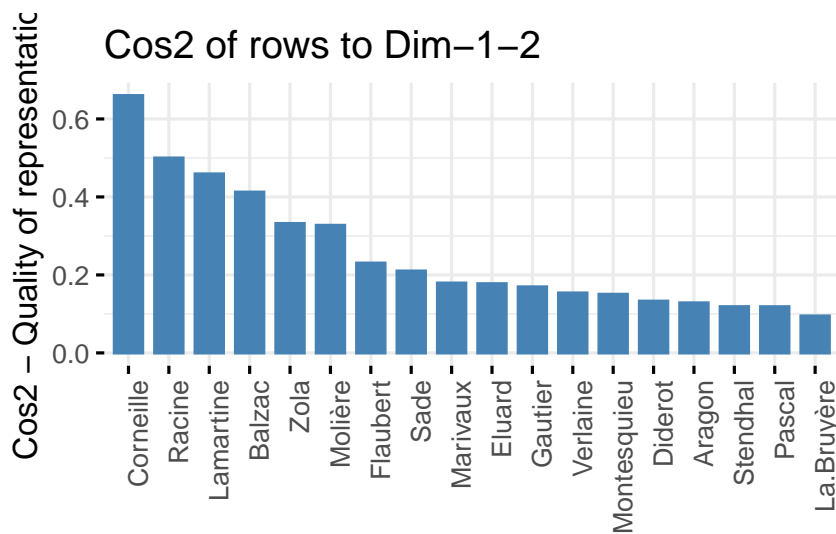


```
fviz_ca_col(res.ca, col.col = "cos2",
            gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
            repel = TRUE)
```

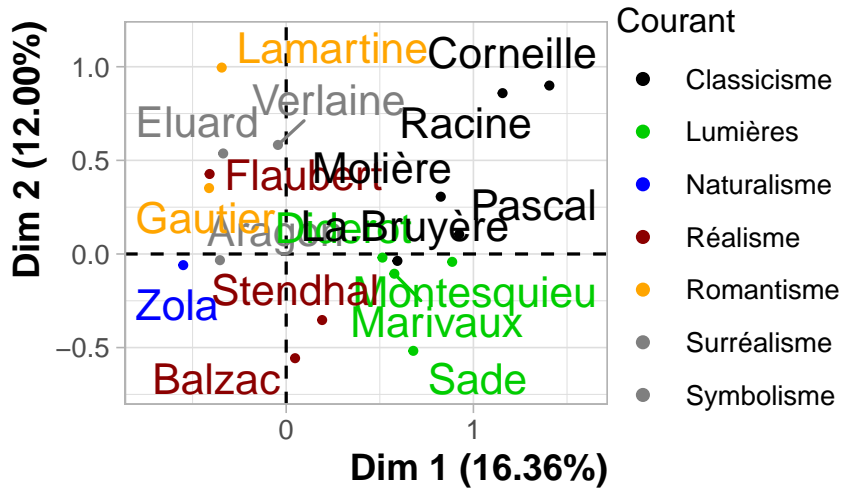
```
## Warning: ggrepel: 974 unlabeled data points (too many overlaps). Consider
## increasing max.overlaps
```



```
fviz_cos2(res.ca, choice = "row", axes = 1:2, xtickslab.rt = 90)
```



```
plot(res.ca,
     invis=c("col", "quali"), hab=3,
     cex=1.2, title="",
     cex.axis=1.2, shadow=TRUE, palette=palette(c("black", "darkred", "orange", "lightblue", "blue", "green3
```



```
res.ca.faster = CA(books[, 1:200], quanti.sup=1, quali.sup=2:3, graph = F)
```

```
### classif
```

```
res.hcpc = HCPC(res.ca.faster, nb.clust=-1, graph=FALSE, consol=FALSE)
```

```
plot(res.hcpc, choice="tree", palette=palette(c("black","green3","blue","darkred","orange","red","grey"
```

```
## Warning in graphics::plotHclust(n1, merge, height, order(x$order), hang, :
```

```
## "palette" is not a graphical parameter
```

```
## Warning in graphics::plotHclust(n1, merge, height, order(x$order), hang, :
```

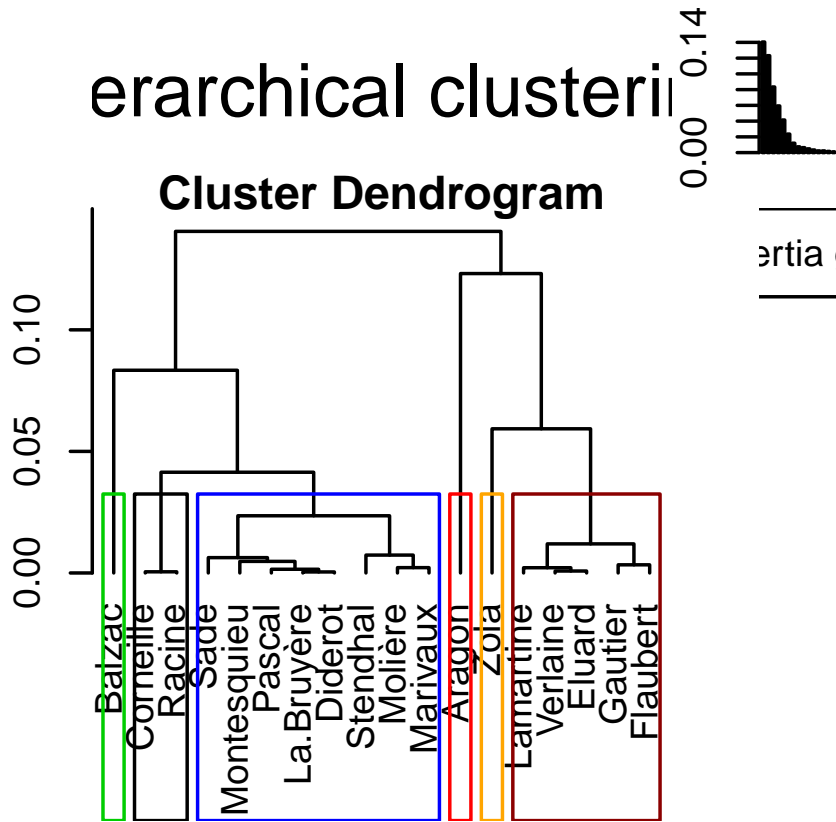
```
## "palette" is not a graphical parameter
```

```
## Warning in axis(2, at = pretty(range(height)), ...): "palette" is not a
```

```
## graphical parameter
```

```
## Warning in title(main = main, sub = sub, xlab = xlab, ylab = ylab, ...):
```

```
## "palette" is not a graphical parameter
```



```
# bb$row$coord[,1]==-bb$row$coord[,1]
# bb$col$coord[,1]==-bb$col$coord[,1]
# bb$quali.sup$coord[,1]==-bb$quali.sup$coord[,1]
# plot(bb, invis=c("col", "quali"), hab=4, cex=1.2, title="", cex.axis=1.2, cex.lab=1.2, palette=palette(c("y
```

```
res.hcpc$desc.ind$para
```

```
## Cluster: 1
##   Racine Corneille
## 0.1234185 0.1234185
## -----
## Cluster: 2
##   Balzac
##     0
## -----
## Cluster: 3
## La. Bruyère   Diderot   Marivaux   Pascal   Stendhal
## 0.1227552   0.2079247   0.3548788   0.3965549   0.4569940
## -----
## Cluster: 4
## Lamartine   Eluard   Flaubert   Gautier   Verlaine
## 0.2383384   0.3165550   0.3313708   0.4123533   0.4149375
```

```
## -----  
## Cluster: 5  
## Zola  
## 0  
## -----  
## Cluster: 6  
## Aragon  
## 0
```