### Correspondence Analysis

In practice

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#### Data

We consider the relationship between hair and eyes colors measured on 592 students in Delaware University.

```
mydata <- read.table("http://pbil.univ-lyon1.fr/R/donnees/snee74.txt",</pre>
    header = TRUE, stringsAsFactors = TRUE)
names(mydata)
## [1] "cheveux" "yeux"
                           "sexe"
head(mydata)
##
    cheveux
              yeux
                       sexe
## 1
       Noir Marron
                      Male
## 2 Blond Bleu Femelle
## 3 Noir Bleu
                      Male
## 4 Marron Marron Femelle
## 5 Roux Marron
                      Male
## 6 Marron
              Bleu
                      Male
```

#### Factors in R

A qualitative variables is stored as factor with different categories (levels)

```
is.factor(mydata$yeux)

## [1] TRUE

levels(mydata$yeux)

## [1] "Bleu" "Marron" "Noisette" "Vert"
```

#### Univariate analysis

• Compute the vector with the number of individuals for each eye color category and display the results on a plot.

## Contingency table

- Build the contingency table crossing hair and eyes colors. See ?table
- Compute row and columns marginal relative frequencies

## Bivariate analysis

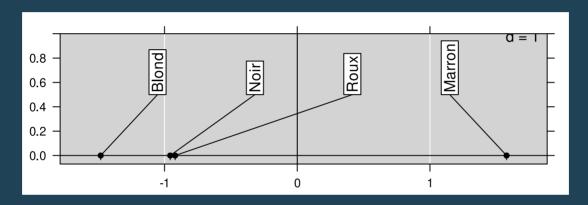
• Compute and interpret the results of a  $\chi^2$  test

### Bivariate analysis

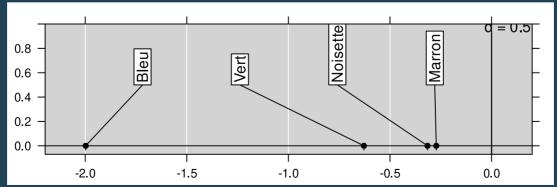
• Display and interpret the associations between categories with mosaicplot

## Scoring

Suppose that we can display hair colors by a score



what would be the best way to display eyes colors? (given the distribution of students)



# Weighted averaging

- Assign a random score to hair colors
- Compute the position for the eye color 'Vert'

## Compute scores for all eye colors

By repeating the same formulas

```
score.e <- sapply(1:4, function(x) sum(score.h * mytab[,
    x]/sum(mytab[, x])))</pre>
```

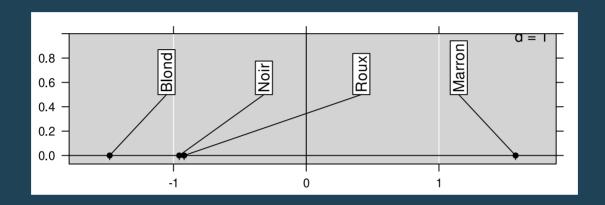
or using matrix algebra

```
t(prop.table(mytab, 2)) %*% score.h
```

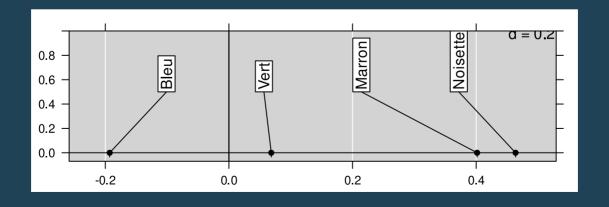
```
## [,1]
## Bleu -0.19286557
## Marron 0.40154742
## Noisette 0.46376597
## Vert 0.06851621
```

## Best representation

if



then



### Reciprocal representation

• Start from a random score for eyes colors to position hair colors by averaging

## Reciprocal averaging

- 1. Set a random score for columns
- 2. Use the column score to compute a score for rows by weighted averaging
- 3. Compute a new score for columns by weighted averaging of row score
- 4. Center and scale the two scores using row and column weights
- 5. Repeat steps 2-4 until convergence

### Correspondence analysis

The solution of the iterative algorithm can also be obtained by the diagonalization procedure provided by the dudi.coa function

```
library(ade4)
coa1 <- dudi.coa(unclass(mytab), scannf = FALSE)</pre>
```

• Check the link between  $\chi^2$  statistic and total inertia

#### Interpretation

- Compute a row score with unit norm (\$11) to obtain a column score by weighted averaging (\$co) with maximal variance (\$eig)
- Compute a row score with unit norm (\$c1) to obtain a column score by weighted averaging (\$li) with maximal variance (\$eig)
- Check both results

### Graphical representation

• Represent and interpret the results using the scatter function (check the method argument of scatter.coa)

#### **Inertia statistics**

- Use the inertia.dudi function to compute inertia statistics for rows and columns
- Represent absolute contributions using the plot function (see ? plot.inertia)