Correspondence analysis: introduction

Correspondence Analysis (CA) is a multivariate statistical technique used to analyze and visualize the relationships between categorical variables in a contingency table. It reduces the dimensionality of the data, representing the associations between rows and columns in a low-dimensional space, typically two dimensions, for an easier interpretation. The Chi-squared distance between rows or columns to highlight associations:

$$\chi^2 = \sum_{i=1}^n \sum_{j=1}^m \frac{(n_{ij} - Np_{ij})^2}{Np_{ij}}$$

where n_{ij} is the observed frequency and Np_{ij} is the expected frequency under the independence assumption.

'HairEyeColor' dataset

HairEyeColor: dataset of 592 observations x 3 variables.

Hair: qualitative variable: Black, Brown, Red, Blond Eye: qualitative variable: Brown, Blue, Hazel, Green

Sex: qualitative variable: Male, Female

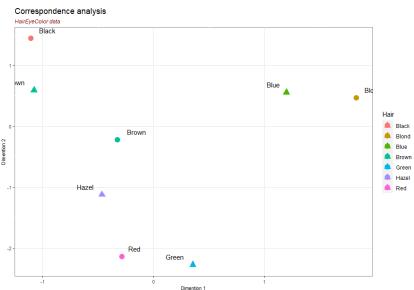
```
1 > head(HaiEyeColor)
2 > HaiEyeColor
3 > head(HairEveColor)
  . . Sex = Male
5
         Eye
7 Hair
          Brown Blue Hazel Green
    Black
            32
                 11
    Brown
            53
                   50
                               15
10
  Red
             10 10
11
                   30
  Blond
12
13
  . . Sex = Female
14
15
         Eye
16 Hair
          Brown Blue Hazel Green
17
    Black
              36
18
    Brown
              66
                   34
                                14
19
    Red
              16
20
    Rl ond
```

Summary

In order to perform a correspondence analysis, we typically call the function 'ca()' from the package 'ca' on a contingency table. The summary of our analysis is displayed below.

```
1 > ca result = ca(contingency table)
2 > ca_result
   Principal inertias (eigenvalues):
6 Value
             0.208773 0.022227 0.002598
7 Percentage 89.37% 9.52%
                              1.11%
   Rows:
10
              Black
                        Brown
                                    Red
                                           Blond
11 Mass
          0.182432 0.483108 0.119932 0.214527
12 ChiDist 0.551192 0.159461 0.354770 0.838397
13 Inertia 0.055425 0.012284
                              0.015095 0.150793
14 Dim. 1 -1.104277 -0.324463 -0.283473 1.828229
15 Dim. 2 1.440917 -0.219111 -2.144015 0.466706
16
17
  Columns:
18
              Brown
                        R111e
                                 Hazel
                                           Green
19 Mass
           0.371622 0.363176
                              0.157095
                                        0.108108
20 ChiDist 0.500487 0.553684
                              0.288654
                                        0.385727
21 Inertia 0.093086 0.111337
                              0.013089
                                        0.016085
22 Dim. 1
          -1.077128 1.198061 -0.465286
                                        0.354011
```

Plot of factors in main dimentions



Main observations

- Dimension Reduction: The relationships between hair color and eye color in a lower-dimensional space, here the first two on the plot.
- Association Visualization: Points close to each other in the plot indicate a stronger association between the corresponding hair and eye colors. For example, if "Black Hair" and "Brown Eyes" are close together (frequently observed together).
- Dimensional Interpretation: The axes (Dimension 1 and Dimension 2) represent the principal dimensions that capture the most variance in the data.
- Categorical Differentiation: The plot visually differentiates between hair and eye colors using different shapes and colors, making it easy to interpret the correspondence between categories.

References

An Introduction to Applied Multivariate Analysis with R, 2011, B. Everitt, T. Hothorn, Springer, e-ISBN 978-1-4419-9650-3

The R Project for Statistical Computing: https://www.r-project.org/