DramaAnalysis

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Contents

Preface

This book is in alpha stadium and not ready. It pertains to the (currently unreleased) version 3.0 of the DramaAnalysis package. Use at your own risk.

Chapter 1

Introduction

1.1 Technical basics

1.1.1 Classes

1.1.2 Pipelining with `margrittr`

DramaAnalysis 3.0

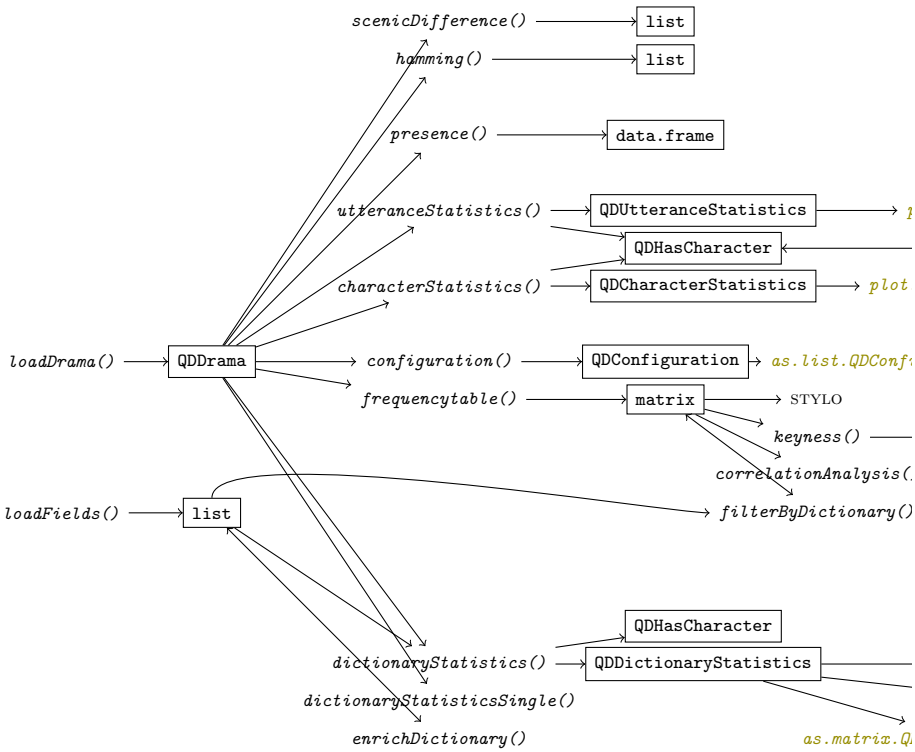


Figure 1.1: Some caption.

Chapter 2

Data

Before analysing any data, it needs to be imported, and converted into the proper structure. In QuaDramA, we process dramatic texts in multiple stages, described below.

2.1 Origin: TEI-XML

The base format that we use (and in which we put all our annotations) is an XML format known as TEI. This format is used by most researchers doing quantitatitve drama analysis. An excellent source for dramas in the proper format is DraCor, maintained by Frank Fischer.

While we are using GerDraCor as a basis, we have added linguistic annotations to a number of plays, and integrated more plays (e.g., translations) into the corpus. This corpus can be found here.

2.2 Preprocessed data

As a first step, we process all dramatic texts using our DramaNLP pipeline. The result of this processing is a set of CSV files for each play that contains the information in the play in a format suitable for analysis with R. This repository contains two plays in the format, which are also included in our R package.

Table 2.1: Different CSV files used in the analysis. *ID* is a placeholder for a unique identifier for the play

File	Description
ID.Metadata.csv	Meta data for the play (author, title, language, …)
ID.Characters.csv	Characters of the play, with some character specific information
ID.Entities.csv	All discourse entities (including characters, but also all other coreference chains)
ID.Mentions.csv	Mentions associated with characters
ID.StageDirections.csv	The stage directions of the play
ID.UtterancesWithTokens.csv	All character utterances of the play
ID.Segments.csv	Information about acts and scenes

2.3 Sample data

```
# Load Emilia Galotti
data(rksp.0)
```

```
# Load Miß Sara Sampson
data(rjmw.0)

text <- combine(rksp.0, rjmw.0)
```

Part I

Structure Analysis

We're assuming here that we have loaded some texts using `loadDrama()`, and that this text is stored as a `QDDrama`-object in the variable `text`. For demo purposes, we will use the two plays that are included in the R-package, Lessing's *Emilia Galotti* and *Miss Sara Sampson*. Both have been preprocessed by the DramaNLP pipeline.

Chapter 3

Who's talking how much?

First, we calculate summary statistics over all characters.

```
charStats <- characterStatistics(text)

charStats
```

```
##    corpus  drama        character tokens types utterances
## 1    test rjmw.0      sir_william   2056   698         23
## 2    test rjmw.0         waitwell   1826   568         41
## 3    test rjmw.0         der_wirt    324   177          7
## 4    test rjmw.0        mellefont   7981  1722        201
## 5    test rjmw.0           norton   1001   407         46
## 6    test rjmw.0            betty    482   223         20
## 7    test rjmw.0             sara   9121  1908        166
## 8    test rjmw.0          marwood   7225  1757        155
## 9    test rjmw.0           hannah    258   155         16
## 10   test rjmw.0     der_bediente     44    34          2
## 11   test rjmw.0         arabella    398   162         13
## 12   test rksp.0        der_prinz   5303  1257        157
## 13   test rksp.0 der_kammerdiener     42    34          6
## 14   test rksp.0            conti    764   325         24
## 15   test rksp.0        marinelli   5567  1324        221
## 16   test rksp.0     camillo_rota    106    62          6
## 17   test rksp.0  claudia_galotti   2098   657         73
```

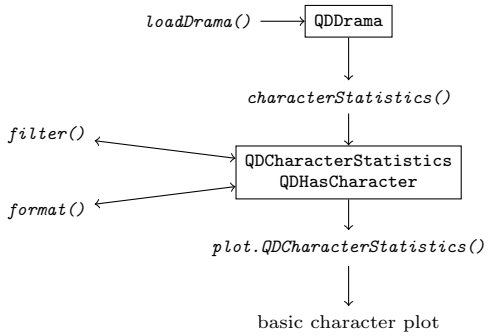


Figure 3.1: Relevant classes and functions in this chaper

```
## 18   test rksp.0            pirro    343   196         25
## 19   test rksp.0          odoardo   3248   891        108
## 20   test rksp.0           angelo    635   300         28
## 21   test rksp.0           emilia   2275   630         64
## 22   test rksp.0          appiani   1112   426         48
## 23   test rksp.0         battista    195   112         11
## 24   test rksp.0           orsina   2832   743         64
##    utteranceLengthMean utteranceLengthSd firstBegin lastEnd
## 1             89.39130        108.948169        597  170838
## 2             44.53659         59.768762        670  158683
## 3             46.28571         45.492543       3571    5695
## 4             39.70647         50.034972       5848  170188
## 5             21.76087         23.601492       6328  170456
## 6             24.10000         36.404019       9014  156987
## 7             54.94578         77.785973      11766  167457
## 8             46.61290         62.383230      28320  141005
## 9             16.12500         15.654073      28392   58523
## 10            22.00000         12.727922      31532   31790
## 11            30.61538         31.967532      43987   52559
## 12            33.77707         40.807176        426  136067
## 13             7.00000          5.215362       1149   24954
## 14            31.83333         40.141778       2654   12212
## 15            25.19005         29.844599      13147  134486
## 16            17.66667         19.179851      25577   26914
## 17            28.73973         29.131897      27006  112526
## 18            13.72000         10.663489      27113   50425
## 19            30.07407         40.399571      27385  135555
## 20            22.67857         19.573326      28777   64877
## 21            35.54688         48.159105      36769  134940
## 22            23.16667         25.525401      44452   56735
## 23            17.72727         32.948720      67465   86481
## 24            44.25000         50.977119      88094  112182
```

This already gives us a lot of information about the characters. In particular, the function `characterStatistics()` returns a table (of the types `QDCharacterStatistics`, `QDHasCharacter` and `data.frame`) with information about:

- the number of tokens a character speaks (`tokens`),
- the number of different tokens a character speaks (`types`),
- the number of utterances (`utterances`),
- the average length of the utterances (`utteranceLengthMean`),
- their standard deviation (`utteranceLengthSd`),
- the character position of the start of the first utterance (`firstBegin`), and
- the character position of the end of the last utterance (`lastEnd`).

The function `characterStatistics()` provides a number of options to control its exact behaviour. Entering `?characterStatistics` in the R console opens the documentation for the function with a description of all the options. We'll describe some frequently used options here as well:

- Punctuation: By default, all punctuation marks are counted as tokens. This behaviour can be changed by setting `filterPunctuation=TRUE`.
- Normalization: The values in the table above are all absolute values. When comparing to other texts, one is often interested in normalized values. If the option `normalize` is set to `TRUE`, all values will be normalised (if applicable).
- Segmentation: By default, the function extracts values for the entire play. With the option `segment`,

it is possible to extract statistics by act or scene, as shown in the example below. Except for the additional column `Act`, the columns in the table are the same as before.

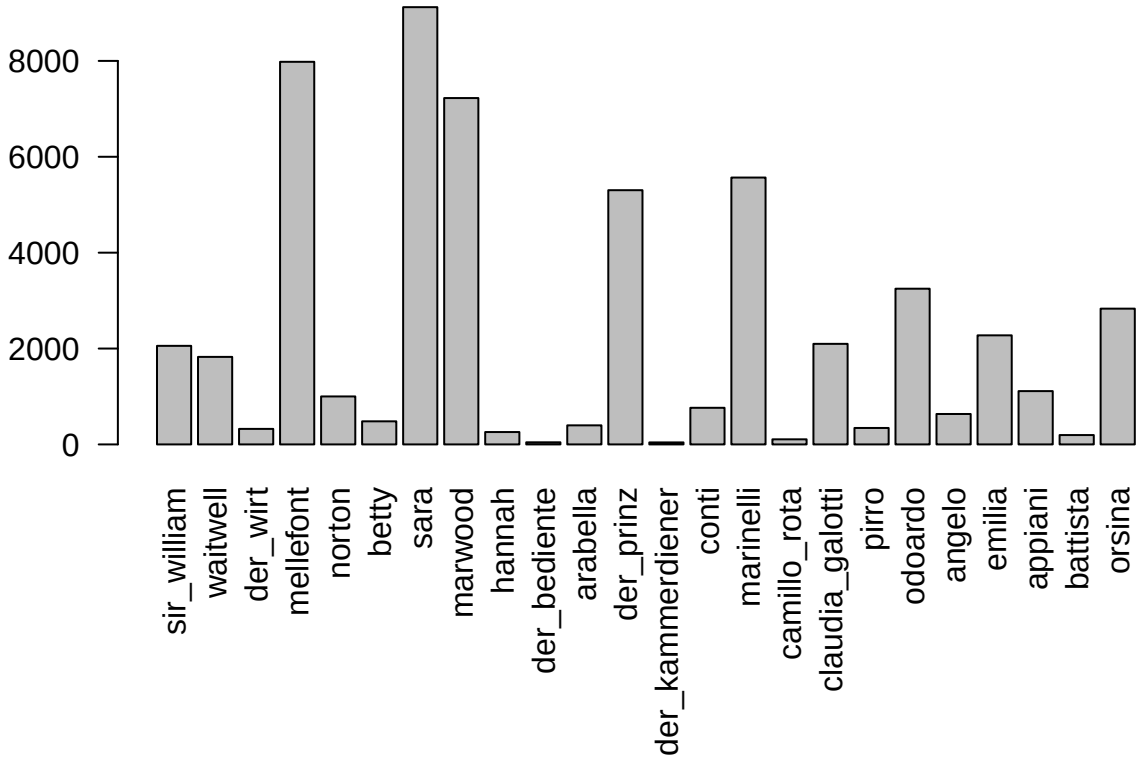
```
characterStatistics(rksp.0,
                    segment="Act")
```

```
##    corpus  drama Act        character tokens types utterances
## 1    test rksp.0   I        der_prinz   2740   795         71
## 2    test rksp.0   I der_kammerdiener     42    34          6
## 3    test rksp.0   I            conti    764   325         24
## 4    test rksp.0   I        marinelli   1038   412         31
## 5    test rksp.0   I     camillo_rota    106    62          6
## 6    test rksp.0  II  claudia_galotti   1252   477         50
## 7    test rksp.0  II            pirro    343   196         25
## 8    test rksp.0  II          odoardo    611   295         17
## 9    test rksp.0  II           angelo    385   214         19
## 10   test rksp.0  II           emilia   1166   417         28
## 11   test rksp.0  II          appiani   1112   426         48
## 12   test rksp.0  II        marinelli    527   255         28
## 13   test rksp.0 III        marinelli   1814   605         56
## 14   test rksp.0 III        der_prinz    945   398         29
## 15   test rksp.0 III           angelo    250   147          9
## 16   test rksp.0 III         battista    187   106          9
## 17   test rksp.0 III           emilia    413   175         16
## 18   test rksp.0 III  claudia_galotti    614   241         16
## 19   test rksp.0  IV        der_prinz    803   319         30
## 20   test rksp.0  IV        marinelli   1358   495         72
## 21   test rksp.0  IV         battista      8     7          2
## 22   test rksp.0  IV           orsina   2832   743         64
## 23   test rksp.0  IV          odoardo    706   275         30
## 24   test rksp.0  IV  claudia_galotti    232   125          7
## 25   test rksp.0   V        marinelli    830   363         34
## 26   test rksp.0   V        der_prinz    815   321         27
## 27   test rksp.0   V          odoardo   1931   620         61
## 28   test rksp.0   V           emilia    696   251         20
##    utteranceLengthMean utteranceLengthSd firstBegin lastEnd
## 1             38.59155         39.891130        426   26559
## 2              7.00000          5.215362       1149   24954
## 3             31.83333         40.141778       2654   12212
## 4             33.48387         34.234360      13147   23619
## 5             17.66667         19.179851      25577   26914
## 6             25.04000         28.927750      27006   56666
## 7             13.72000         10.663489      27113   50425
## 8             35.94118         41.800823      27385   36244
## 9             20.26316         19.674694      28777   32436
## 10            41.64286         63.826277      36769   48503
## 11            23.16667         25.525401      44452   56735
## 12            18.82143         17.346263      50431   55936
## 13            32.39286         35.245982      56826   79337
## 14            32.58621         48.327393      56914   72918
## 15            27.77778         19.466495      63132   64877
## 16            20.77778         36.033935      67465   75603
## 17            25.81250         27.352559      67517   79537
## 18            38.37500         28.765431      74777   79747
## 19            26.76667         38.699268      79838   95409
```

```
## 20          18.86111          20.750229      79972  103998
## 21           4.00000           2.828427      86363   86481
## 22          44.25000          50.977119      88094  112182
## 23          23.53333          18.528140     101533  112733
## 24          33.14286          29.952343     110037  112526
## 25          24.41176          36.802600     112824  134486
## 26          30.18519          37.235257     113752  136067
## 27          31.65574          47.382094     114998  135555
## 28          34.80000          34.284568     128910  134940
```

Of course, the values in the above table can be directly plotted:

```
charStats <- characterStatistics(text, normalize=FALSE)

par(mar=c(9,3,2,2)) # increase plot margins, so that the labels can be plotted

barplot(charStats$tokens,  # these are the values to be plotted
        names.arg = charStats$character, # the labels on the x-axis
        las=2 # rotate the labels on both axes
)
```

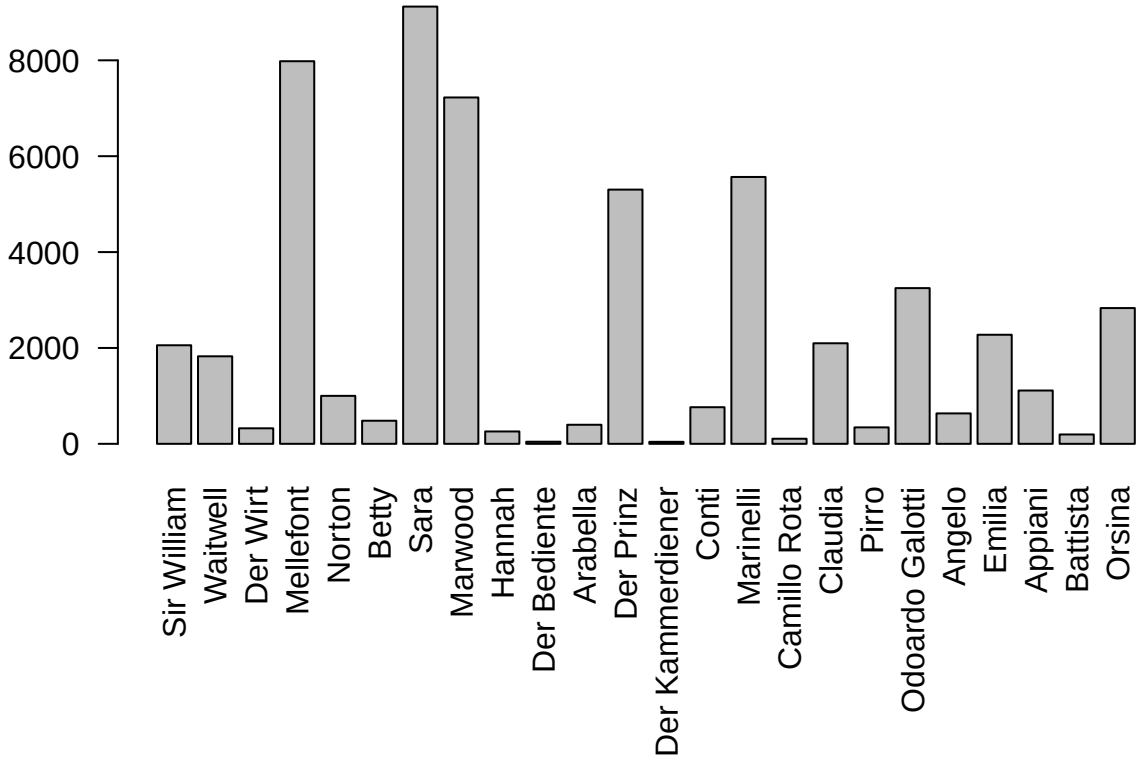


3.1 Character names instead of identifiers

By default, all our functions identify characters using technical ids, which may or may not be human-readable. Even if they are, it's usually a good idea to replace them with nice to read labels before publication. We therefore provide the function `format()`, which can be applied to any table that contains a column with character ids (i.e., any object of type `QDHasCharacter`).

```
charStats <- characterStatistics(text, normalize=FALSE)
```

```
charStats <- format(charStats, text) # Replace ids with names

par(mar=c(9,3,2,2)) # increase plot margins, so that the labels can be plotted

barplot(charStats$tokens,  # these are the values to be plotted
        names.arg = charStats$character, # the labels on the x-axis
        las=2 # rotate the labels on both axes
)
```

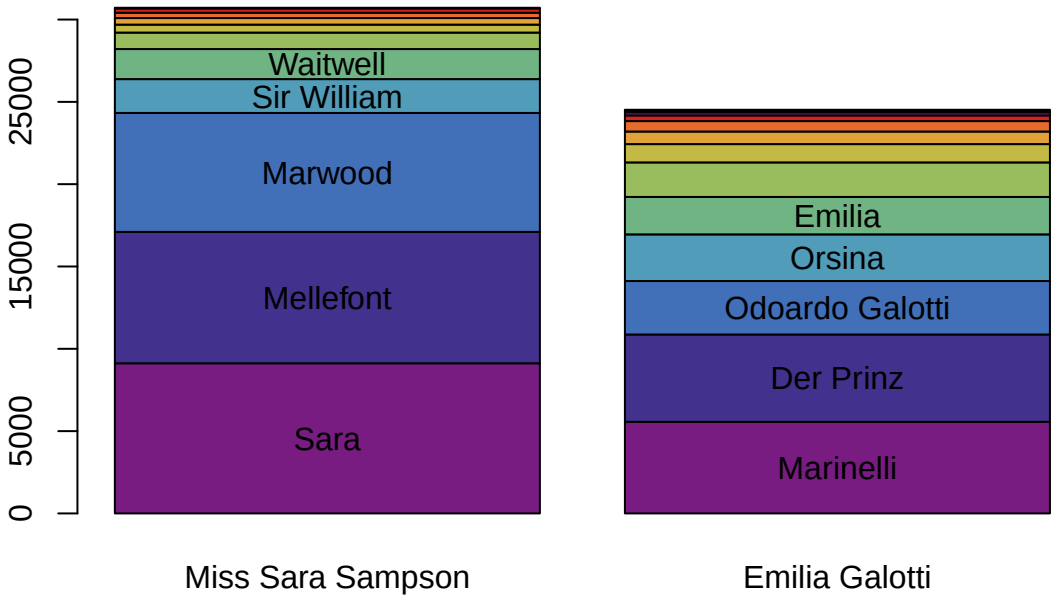


As can be seen above `format()` requires two arguments: The table in which we want to replace ids by characters, and the original drama object (that we got from calling the function `loadDrama()`).

3.2 Stacked bar plot

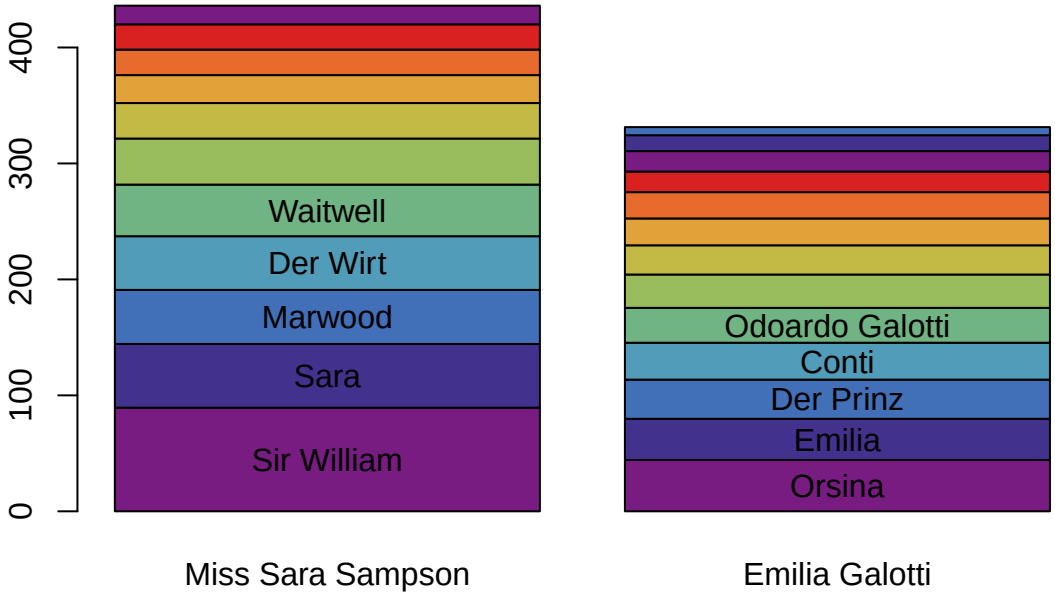
The plot shown above is quite wide, and some aspects (like the ranking in terms of spoken tokens), is hard to see. We often use another way of visualizing this, which can be used if you supply the `QDCharacterStatistics` directly into the `barplot()` function. In combination with the `margrittr` pipes, we can call it like this (enter `?barplot.QDCharacterStatistics` for details on the special barplot function):

```
characterStatistics(text, normalize=FALSE) %>%
  format(text) %>%
  barplot(names.arg=c("Miss Sara Sampson", "Emilia Galotti"))
```



By default, the `barplot.QDCharacterStatistics()` function visualises the number of tokens spoken by the characters (and ranks the characters accordingly). This can be changed by supplying the option `column`, and specifying another value.

```
characterStatistics(text, normalize=FALSE) %>%
  format(text) %>%
  barplot(names.arg=c("Miss Sara Sampson", "Emilia Galotti"),
          column = "utteranceLengthMean") # show mean utterance length
```



This picture looks quite different! Other interesting columns to experiment with are `types` and `utteranceLengthSd`.

Chapter 4

Who's talking how often?

So far, we have counted words for characters. Now we will turn to utterances, and their properties.

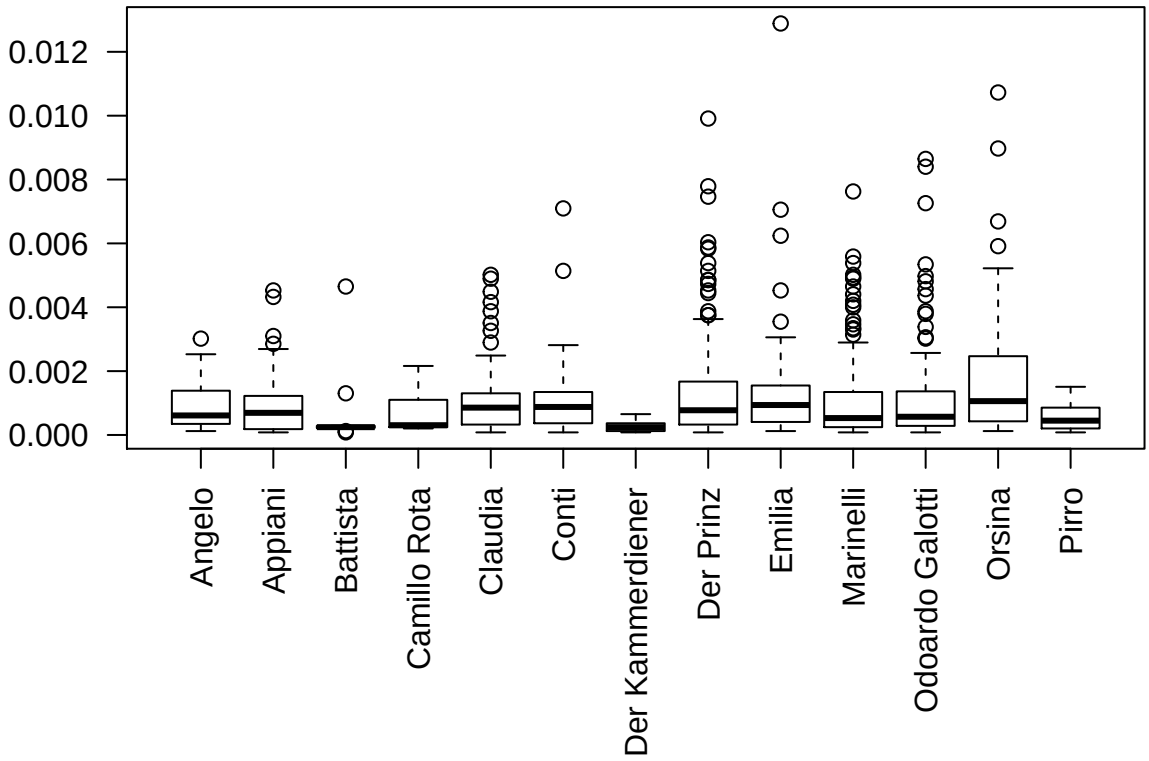
First, we will use the function `utteranceStatistics()` to extract quantitative information about utterances:

```
utteranceStatistics(rksp.0)
```

```
##    corpus  drama        character utteranceBegin utteranceLength
## 1    test rksp.0        der_prinz            426    3.874388e-03
## 2    test rksp.0 der_kammerdiener           1149    8.156607e-05
## 3    test rksp.0        der_prinz           1178    2.895595e-03
## 4    test rksp.0 der_kammerdiener           1526    6.525285e-04
## 5    test rksp.0        der_prinz           1655    2.854812e-04
## 6    test rksp.0 der_kammerdiener           1697    1.631321e-04
## 7    test rksp.0        der_prinz           1739    1.019576e-03
## 8    test rksp.0 der_kammerdiener           1857    3.262643e-04
## 9    test rksp.0        der_prinz           1919    2.528548e-03
## 10   test rksp.0 der_kammerdiener           2299    3.670473e-04
```

This creates a table that is very long, which is why we only show the first 10 rows here. The table contains one row for each utterance, and information about the speaker of the utterance, its length (measured in tokens) and its starting position (character position). We can now inspect the variance in utterance length:

```
ustat <- utteranceStatistics(rksp.0) %>%
  format(rksp.0)   # use names instead of character ids (see above)

par(mar=c(9,4,2,2)) # increase margin

boxplot(utteranceLength ~ character, # what do we want to correlate
        data=ustat,
        las = 2 # rotate axis labels
      )
```

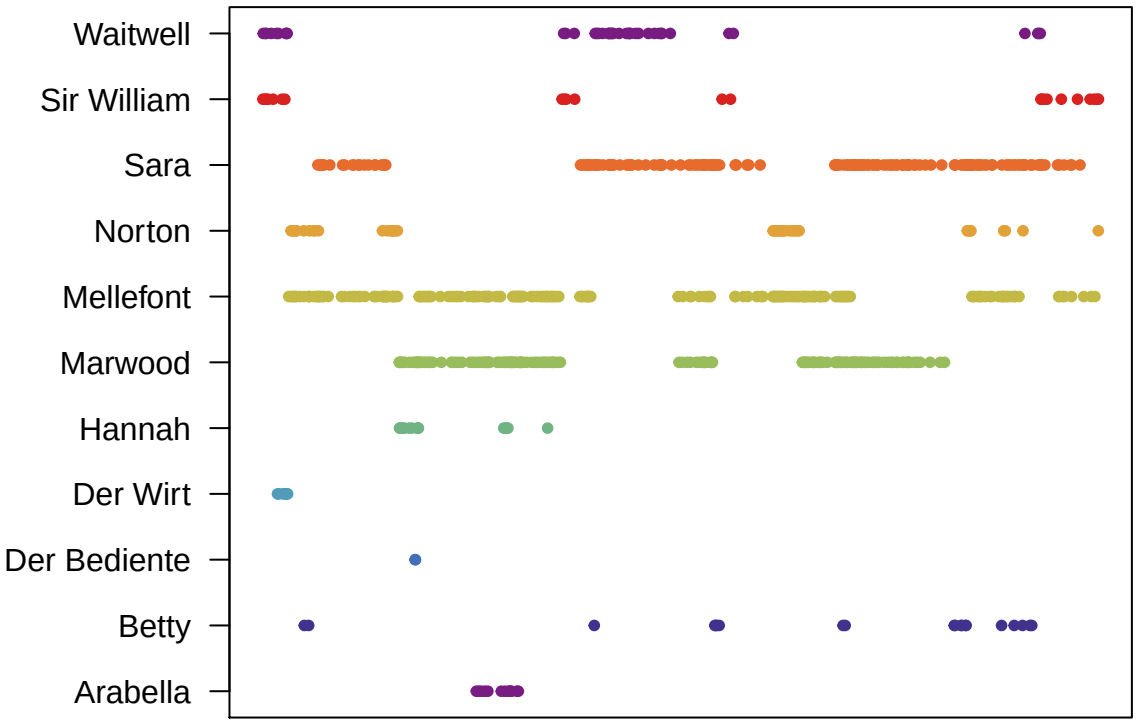


This uses the regular `boxplot()` function, enter `?boxplot` for documentation. `utteranceLength ~ character` is called a formula in R and (in this case) expresses that we want to look at the column `utteranceLength`, grouped by the column `character`. The boxplot is a useful way to grasp the dispersion of a set of values (in this case: the lengths of all utterances by a character).

4.1 When are characters talking?

While the above displays the *length* of utterances, we can also display the position of utterances (remember the column `utteranceBegin`?). The following snippet visualizes when characters are talking, this time for Lessings Miss Sara Sampson:

```
par(mar=c(2,7,2,2))

utteranceStatistics(rjmw.0) %>%
  format(rjmw.0) %>%    # character names instead of ids
  plot()                # calling plot.QDUtteranceStatistics()
```

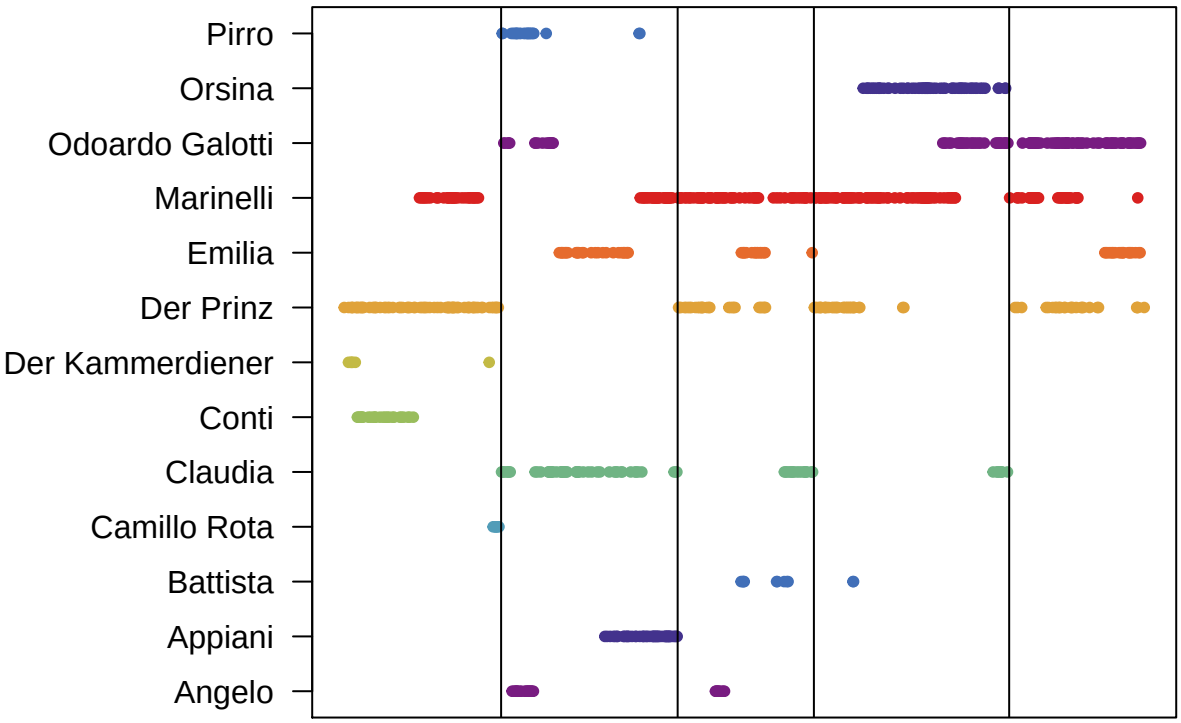


Each dot in this plot represents one utterance, the x-axis is measured in character positions. This is not really intuitive, but the flow from left to right represents the flow of the text. More technically, we again apply the function `format()` to display character names instead of character ids. This is the same function as above, just applied to a different table. It can be applied to any table of the type `QDHasCharacter`. Secondly, the call to the function `plot()` gets rerouted to the function `plot.QDUtteranceStatistics()`, because the object we supply as argument is of the type `QDUtteranceStatistics`. Information about this function can be retrieved by entering `?plot.QDUtteranceStatistics`.

4.2 Adding act boundaries

Now it would be useful to include information on act/scene boundaries in this plot. This can be done by supplying the original drama object as a second argument to the `plot()` function:

```
par(mar=c(2,8,2,2))
utteranceStatistics(rksp.0) %>%
  format(rksp.0) %>%
  plot(rksp.0) # adding the `QDDrama` object here creates the act boundaries.
```



Please note that the information contained in this plot is very similar to the information in configuration matrices.

Chapter 5

Configuration

This section does not refer to the configuration of the R-package, but to the literary analysis concept configuration matrix (Pfister, 1988).

5.1 Matrices

Configuration matrices can be extracted with the function `configuration()`. As usual, entering `?configuration` provies more detailed information about the options the function provides.

```
configuration(rksp.0)
```

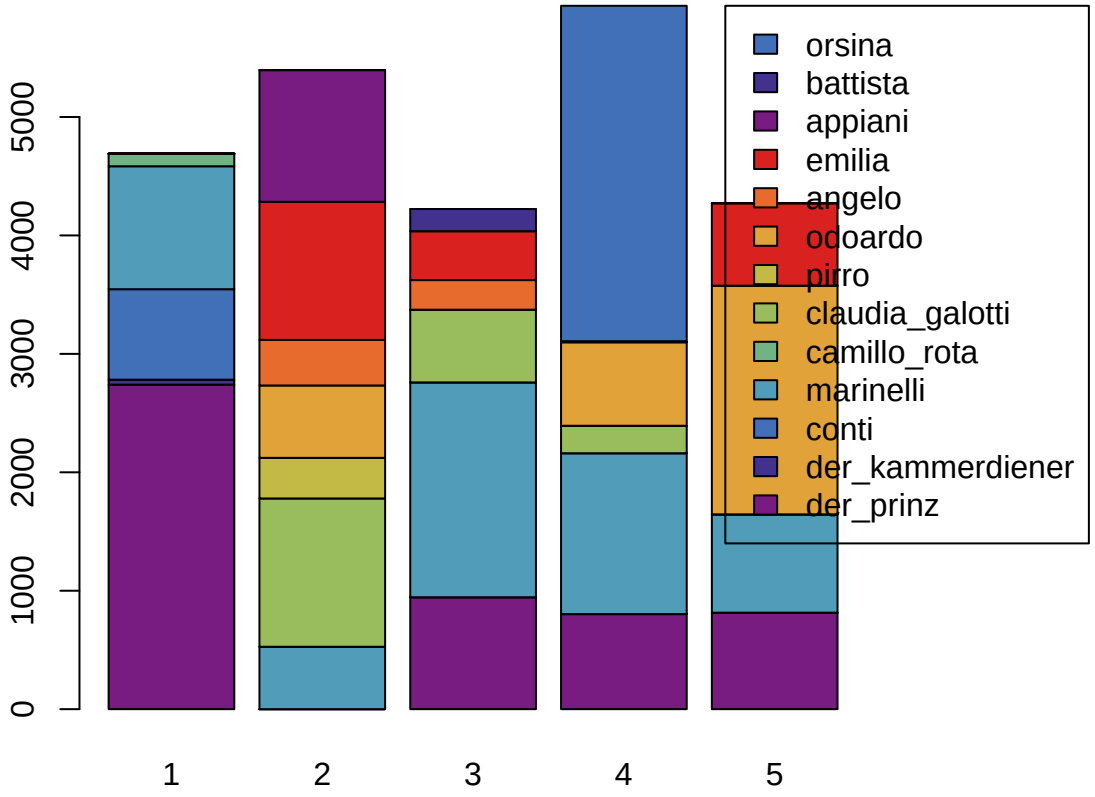
```
##    corpus  drama        character    1    2    3    4    5
## 1    test rksp.0        der_prinz 2740    0  945  803  815
## 2    test rksp.0 der_kammerdiener   42    0    0    0    0
## 3    test rksp.0            conti  764    0    0    0    0
## 4    test rksp.0        marinelli 1038  527 1814 1358  830
## 5    test rksp.0     camillo_rota  106    0    0    0    0
## 6    test rksp.0  claudia_galotti    0 1252  614  232    0
## 7    test rksp.0            pirro    0  343    0    0    0
## 8    test rksp.0          odoardo    0  611    0  706 1931
## 9    test rksp.0           angelo    0  385  250    0    0
## 10   test rksp.0           emilia    0 1166  413    0  696
## 11   test rksp.0          appiani    0 1112    0    0    0
## 16   test rksp.0         battista    0    0  187    8    0
## 22   test rksp.0           orsina    0    0    0 2832    0
```

This creates a basic configuration matrix, but instead of just containing the presence or absence of a figure, it contains the number of spoken tokens for each act for each character. This information is in fact similar to what we can extract with `characterStatistics(rksp.0, segment="Act")`, but in a different form and structure. The above table contains a lot of information that can be visualised.

We first need to extract the numeric content from the above table. This can easily be done with the function `as.matrix()` (this will, in detail, be rerouted to the function `as.matrix.QDConfiguration()`, which knows what part of the table needs to be in the matrix). A matrix containing only numbers can easily be plotted using the regular `barplot()` function, as shown below.

```
c <- configuration(rksp.0)

mat <- as.matrix(c)
```

```
par(mar=c(2,2,2,10))

barplot(mat,
        legend.text = c$character,  # legend text
        args.legend = list(x=ncol(mat)+3,      # legend x position
                           y=max(colSums(mat)) # legend y pos
                         ),
        col=qd.colors)
```

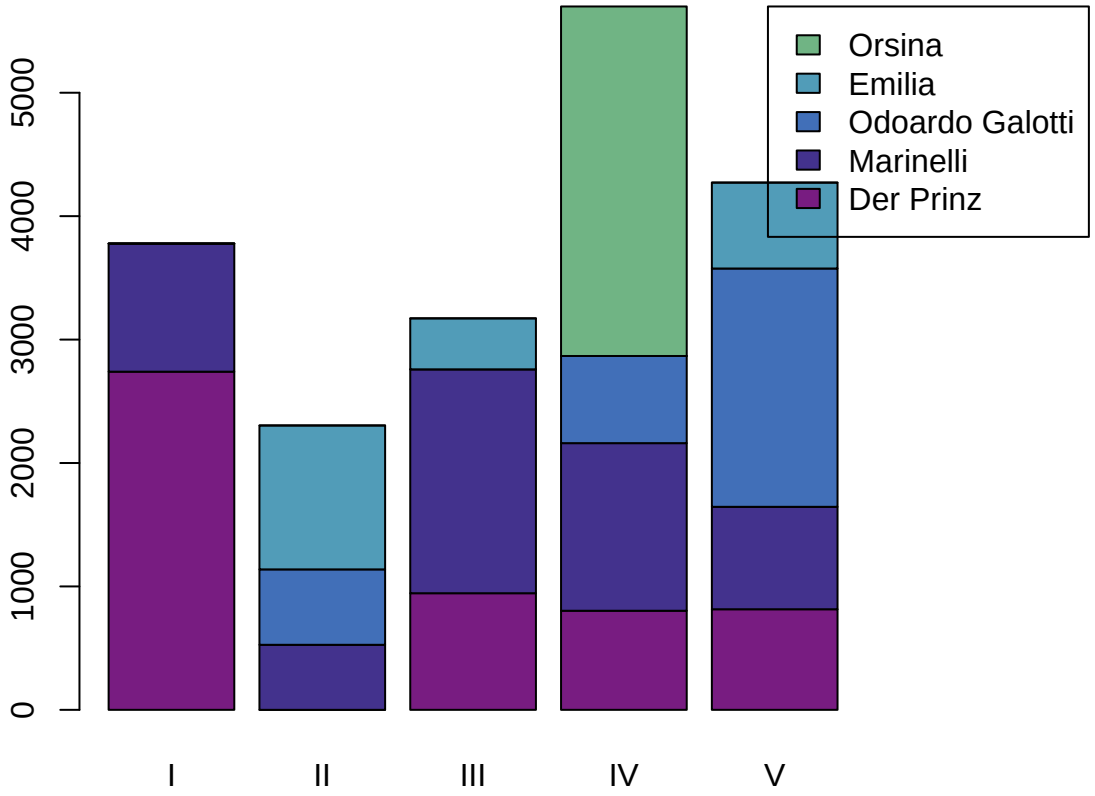


5.1.1 Filtering unimportant characters

This is informative, but doesn't look very nice and some colors are difficult to associate with characters because colors are repeating. We will therefore use the function `filter()`, which is very similar to `format()`: It can be applied to any object of the type `QDHasCharacter` and removes rows according to certain criteria. In this case, we filter every character except the five most talkative ones. As usual, see `?filter.QDHasCharacter` to see other options.

```
c <- configuration(rksp.0) %>%
  filter(rksp.0, threshold = 5) %>%
  format(rksp.0)

mat <- as.matrix(c)

par(mar=c(2,2,2,10))

barplot(mat,
        names.arg = as.roman(1:ncol(mat)),   # roman act numbers
        legend.text = c$character,  # legend text
        args.legend = list(x=ncol(mat)+3,   # legend x position
```

```
                        y=max(colSums(mat)) # legend y pos
                    ),
       col=qd.colors)
```



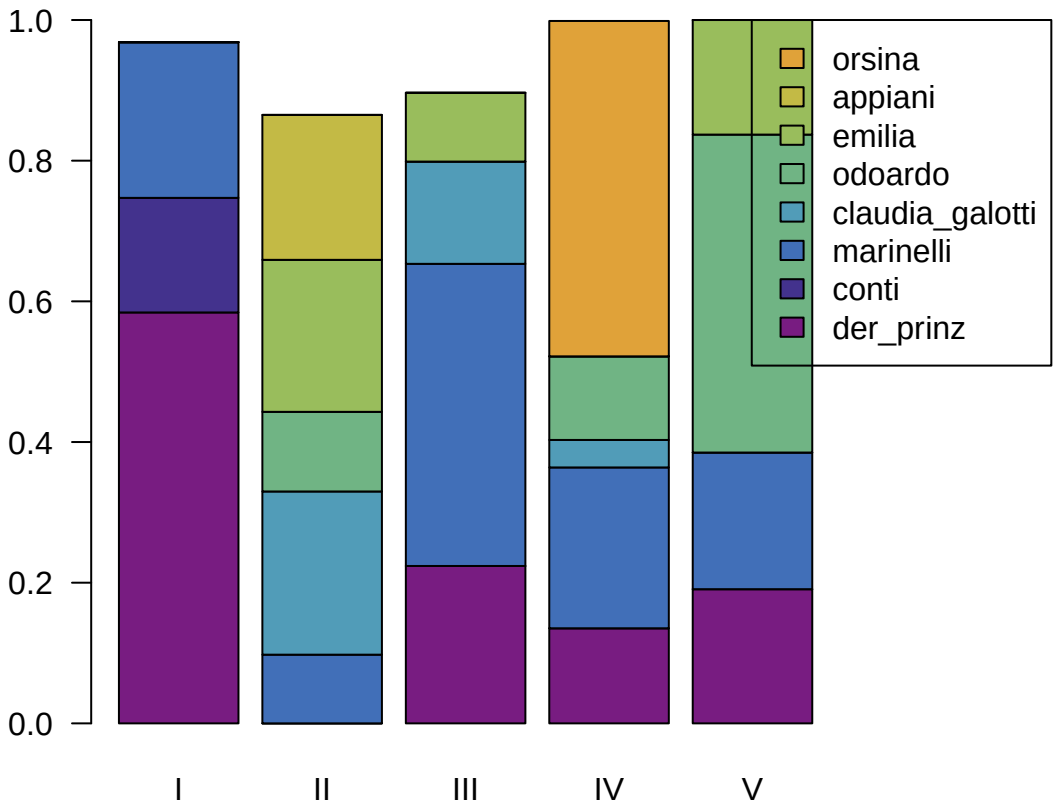
Of course, the speech of characters that do not speak often is now removed, leaving only a portion of each act covered in the plot.

5.1.2 Normalization

Since each act has a different length, it is often useful to normalize each block, according to the total number of spoken tokens. This way, we can display the relative active presence of each character in each act. In combination with the filtering we did before, however, we need to be careful: If the scale with the filtered matrix, a certain portion of the character speech gets lost. The following snippet shows how to scale using the original matrix, but still only include the top eight characters into the matrix.

```
c <- configuration(rksp.0)    # getting the full configuration (with all characters)
mat_orig <- as.matrix(c)      # extracting the matrix

c_filtered <- c %>% filter(rksp.0, threshold=8) # filtering, so that only the top 8 characters remain
mat_filtered <- as.matrix(c_filtered)  # extracting the filtered matrix

# scaling, using values from the unfiltered matrix
mat <- scale(mat_filtered,
             center=FALSE,
             scale=colSums(mat_orig))

# plot, as above
par(mar=c(2,3,2,10))
barplot(mat,
```

```
        names.arg = as.roman(1:ncol(mat)),  # roman act numbers
        # legend
        legend.text=c_filtered$character,            # set legend text
        args.legend = list(x=ncol(mat)+3,      # legend x position
                           y=max(colSums(mat)) # legend y pos
                         ),
        # visual
        las=1,        # rotate labels on y axis
        col=qd.colors # provide colors
)
```



5.2 Copresence

Configuration matrices are also often used to get an overview of who is copresent on stage. This can also be achieved using the function `configuration()`. First, we create a configuration matrix that only represents presence or absence of a figure (and we switch to scenes). Obviously, the resulting matrix has many more columns, we include only the first 10 below.

```
configuration(rksp.0, onlyPresence = TRUE, segment="Scene")
```

```
##    corpus  drama        character     1     2     3     4     5     6
## 1    test rksp.0       der_prinz  TRUE  TRUE  TRUE  TRUE  TRUE  TRUE
## 2    test rksp.0 der_kammerdiener  TRUE FALSE FALSE FALSE FALSE FALSE
## 4    test rksp.0            conti FALSE  TRUE FALSE  TRUE FALSE FALSE
## 9    test rksp.0       marinelli FALSE FALSE FALSE FALSE FALSE  TRUE
## 14   test rksp.0    camillo_rota FALSE FALSE FALSE FALSE FALSE FALSE
## 15   test rksp.0  claudia_galotti FALSE FALSE FALSE FALSE FALSE FALSE
## 16   test rksp.0           pirro FALSE FALSE FALSE FALSE FALSE FALSE
```

```
## 17   test rksp.0          odoardo FALSE FALSE FALSE FALSE FALSE FALSE
## 20   test rksp.0           angelo FALSE FALSE FALSE FALSE FALSE FALSE
## 25   test rksp.0           emilia FALSE FALSE FALSE FALSE FALSE FALSE
## 27   test rksp.0          appiani FALSE FALSE FALSE FALSE FALSE FALSE
## 47   test rksp.0         battista FALSE FALSE FALSE FALSE FALSE FALSE
## 65   test rksp.0           orsina FALSE FALSE FALSE FALSE FALSE FALSE
##        7
## 1   TRUE
## 2   TRUE
## 4  FALSE
## 9  FALSE
## 14 FALSE
## 15 FALSE
## 16 FALSE
## 17 FALSE
## 20 FALSE
## 25 FALSE
## 27 FALSE
## 47 FALSE
## 65 FALSE
```

Creating a co-occurrence matrix is a simple matter of matrix multiplication, and we already know how to create a matrix.

```
# extract the configuration
c <- configuration(rksp.0, onlyPresence = TRUE, segment="Scene")

# extract a matrix
mat <- as.matrix(c)

# multiply the matrix with its inverse
# this creates the copresence matrix
copresence <- mat %*% t(mat)

# add character names
rownames(copresence) <- c$character
colnames(copresence) <- c$character

copresence
```

```
##                  der_prinz der_kammerdiener conti marinelli camillo_rota
## der_prinz               17                2     2         9            1
## der_kammerdiener         2                2     0         0            0
## conti                    2                0     2         0            0
## marinelli                9                0     0        19            0
## camillo_rota             1                0     0         0            1
## claudia_galotti          0                0     0         3            0
## pirro                    0                0     0         1            0
## odoardo                  2                0     0         4            0
## angelo                   0                0     0         1            0
## emilia                   2                0     0         4            0
## appiani                  0                0     0         2            0
## battista                 1                0     0         3            0
## orsina                   1                0     0         3            0
##                  claudia_galotti pirro odoardo angelo emilia appiani
```

```
## der_prinz                    0     0       2      0      2       0
## der_kammerdiener             0     0       0      0      0       0
## conti                        0     0       0      0      0       0
## marinelli                    3     1       4      1      4       2
## camillo_rota                 0     0       0      0      0       0
## claudia_galotti             13     3       3      0      3       4
## pirro                        3     4       1      1      0       1
## odoardo                      3     1      12      0      2       0
## angelo                       0     1       0      2      0       0
## emilia                       3     0       2      0      7       1
## appiani                      4     1       0      0      1       5
## battista                     2     0       0      0      1       0
## orsina                       1     0       3      0      0       0
##                   battista orsina
## der_prinz                1      1
## der_kammerdiener         0      0
## conti                    0      0
## marinelli                3      3
## camillo_rota             0      0
## claudia_galotti          2      1
## pirro                    0      0
## odoardo                  0      3
## angelo                   0      0
## emilia                   1      0
## appiani                  0      0
## battista                 4      0
## orsina                   0      6
```

The resulting copresence matrix shows the number of scenes in which two characters are both present. The diagonal shows the number of scenes in which a character is present in total (because each character is always copresent with itself, so to speak).

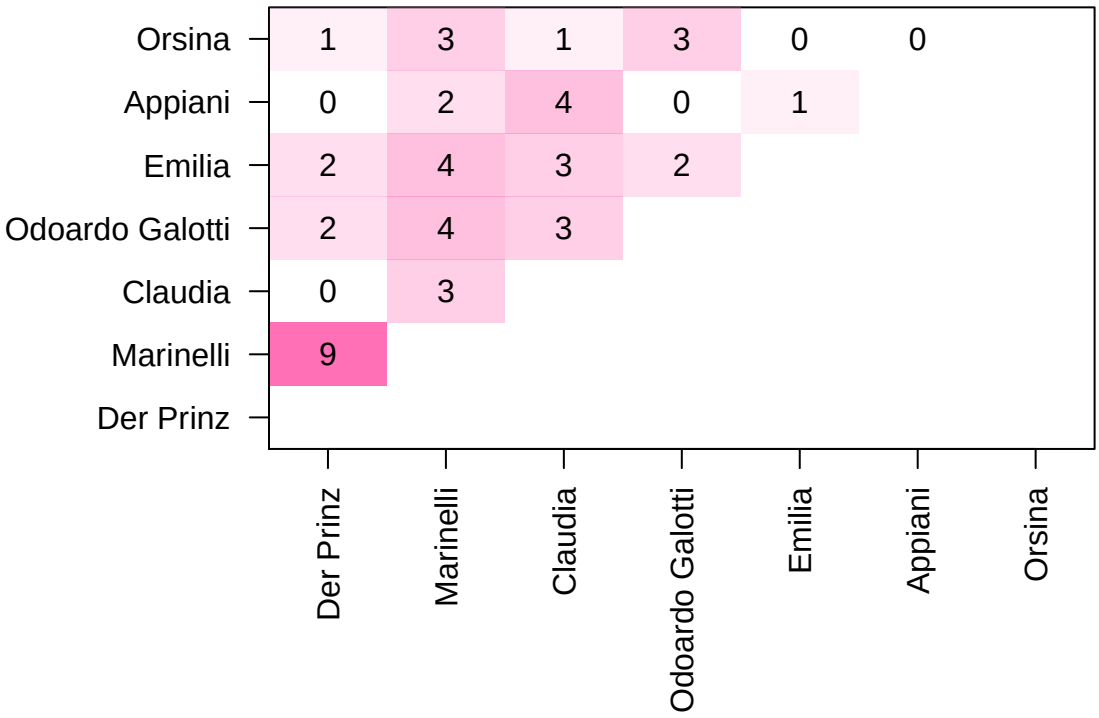
There are multiple ways to visualise this copresence. One option is a heat map, as shown below.

5.2.1 As Heatmap

The copresence can be visualised in a simple heat map. We first focus on the lower triangle and also remove the diagonal values. The actual plotting is a bit more complicated in this case, because we are just using the values in the copresence matrix as pixel intensities in the plot. Also, the axes need to be suppressed first, and can be added later with the proper names of the characters. If needed the code can also be used to include labels into the heat map.

```
c <- configuration(rksp.0, onlyPresence = TRUE, segment="Scene") %>%
  filter(rksp.0, threshold = 7) %>%
  format(rksp.0)

# extract a matrix
mat <- as.matrix(c)

# multiply the matrix with its inverse
# this creates the copresence matrix
copresence <- mat %*% t(mat)

# add character names
rownames(copresence) <- c$character
```

```r
colnames(copresence) <- c$character

# since it's a square matrix, we don't need the bottom right triangle
# and diagonales.
copresence[lower.tri(copresence,diag=TRUE)] <- NA

par(mar=c(10,10,1,1)) # plot margins
image(copresence,
      col = rgb(256,111,184, alpha=(seq(0,255)),
                maxColorValue = 256),
      xaxt= "n",  # no x axis
      yaxt= "n",  # no y axis
      frame=TRUE  # print a frame around the heatmap
      )

# include values as labels
text(y=(rep(1:ncol(copresence), each=nrow(copresence))-1)/(nrow(copresence)-1),
     x=(1:nrow(copresence)-1)/(nrow(copresence)-1),
     labels=as.vector(copresence))

# add the x axis
axis(1, at = seq(0,1,length.out = length(c$character)), labels = c$character, las=3)
# add the y axis
axis(2, at = seq(0,1,length.out = length(c$character)), labels = c$character, las=1)
```



Apparently, Marinelli and the prince have the most shared scenes. Marinelli also shares a scene with most other figures (sum of the vertical bar: 16).

5.2.2 As Network

The same information can also be visualized as a copresence network. For this, we employ the R-package `igraph`. A nice introduction to `igraph` can be found in (Arnold and Tilton, 2015), particularly for literary networks.

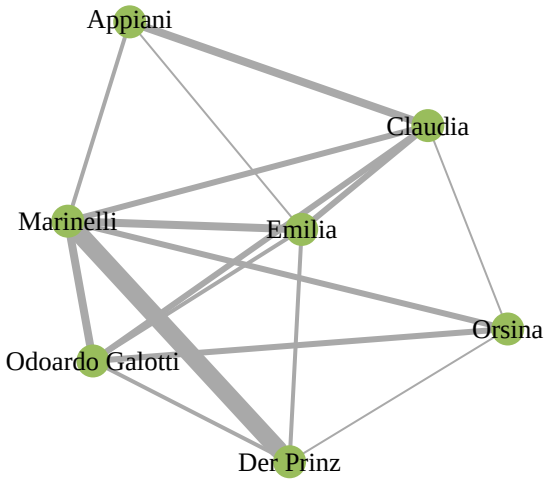
Technically, the matrix we created before is an adjacency matrix. It is therefore simple to convert it to a graph, and igraph offers the function `graph_from_adjacency_matrix()` for this.

```
c <- configuration(rksp.0, onlyPresence = TRUE, segment="Scene") %>%
  filter(rksp.0, threshold = 7) %>%
  format(rksp.0)

# extract a matrix
mat <- as.matrix(c)

# multiply the matrix with its inverse
# this creates the copresence matrix
copresence <- mat %*% t(mat)

# add character names
rownames(copresence) <- c$character
colnames(copresence) <- c$character

# convert the adjacency matrix to a graph object
g <- igraph::graph_from_adjacency_matrix(copresence,
                                 weighted=TRUE,     # weighted graph
                                 mode="undirected", # no direction
                                 diag=FALSE         # no looping edges
                                )
```

Now the variable `g` holds the graph object. There are different things we can do with the graph. First, we can visualise it. `igraph` uses the same mechanism of R that we have used before for plotting, specifying a `plot()` function that can plot graph objects.

```
# Now we plot
plot(g,
     layout=layout_with_gem,       # how to lay out the graph
     main="Copresence Network: Emilia Galotti",  # title
     vertex.label.cex=0.8,         # label size
     vertex.label.color="black",   # font color
     vertex.color=qd.colors[6],    # vertex color
     vertex.frame.color=NA,        # no vertex border
     edge.width=E(g)$weight        # scale edges according to their weight
)
```

Copresence Network: Emilia Galotti



5.2.3 Network properties

`igraph` offers a number of function to extract graph properties. We will show them below without explanation. The documentation of each function can be found here.

5.2.3.1 Density

Function: `graph.density()`

```
graph.density(g)
```

```
## [1] 0.7619048
```

5.2.3.2 Average nearest neighbor degree

Function: `knn()`

```
knn(g)
```

```
## $knn
##       Der Prinz       Marinelli         Claudia Odoardo Galotti
##        4.214286        2.760000        4.714286        5.214286
##          Emilia         Appiani          Orsina
##        6.166667        7.285714        8.375000
##
## $knnk
## [1]      NaN      NaN 7.285714 6.294643 5.365079 2.760000
```

5.2.3.3 Edge connectivity

Function: `edge_connectivity()`

```
edge_connectivity(g)
```

```
## [1] 3
```

5.2.4 Graph Export

As a final step, one might want to further work on the graph using Gephi, or other tools. In order to do so, one can export the graph into an appropriate file:

```
igraph::write_graph(g,
                    "rksp.0.graphml",    # specify the file name
                    format="graphml"     # specify the file format
)
```

This results in a file called `rksp.0.graphml`, that starts similarly as this:

```
<?xml version="1.0" encoding="UTF-8"?>
<graphml xmlns="http://graphml.graphdrawing.org/xmlns"
         xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
         xsi:schemaLocation="http://graphml.graphdrawing.org/xmlns
         http://graphml.graphdrawing.org/xmlns/1.0/graphml.xsd">
<!-- Created by igraph -->
  <key id="v_name" for="node" attr.name="name" attr.type="string"/>
  <key id="e_weight" for="edge" attr.name="weight" attr.type="double"/>
  <graph id="G" edgedefault="undirected">
    <node id="n0">
      <data key="v_name">DER KAMMERDIENER</data>
    </node>
    <node id="n1">
      <data key="v_name">DER PRINZ</data>
    </node>
    ...
```

This file can be opened with Gephi.

Chapter 6

Character Exchange

Character exchange – entering and leaving the stage – often takes place at scene boundaries. We offer multiple ways to measure and visualize this exchange over the course of the play.

6.1 Hamming Distance

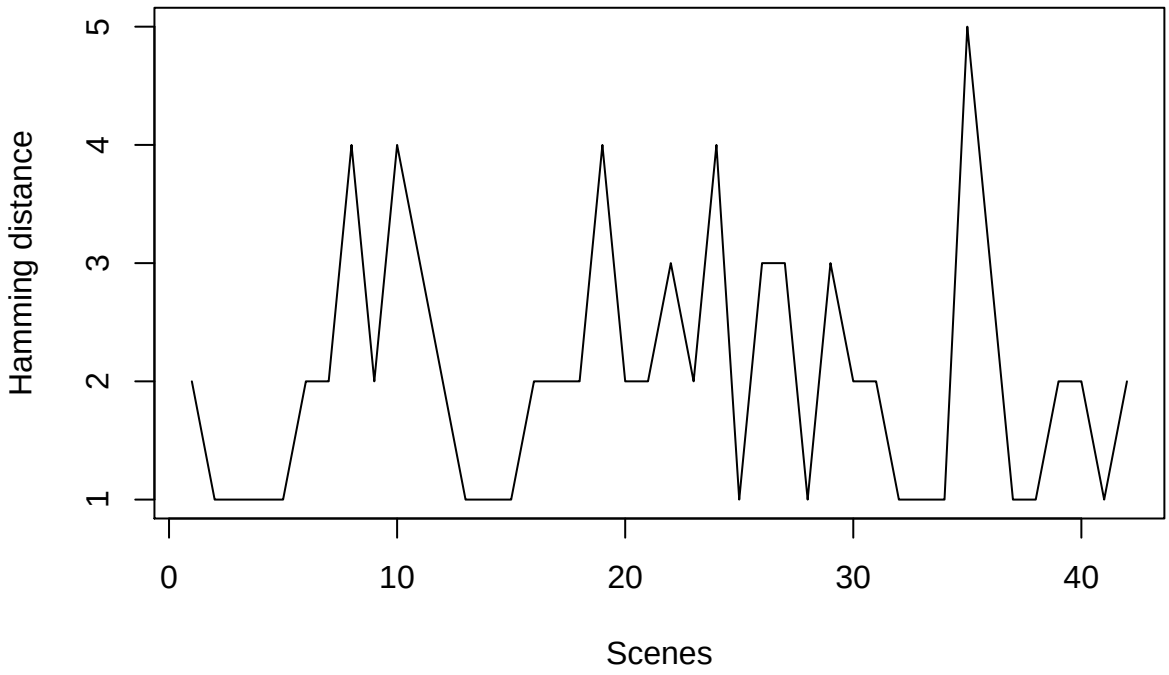
The hamming distance has been introduced in information theory by Richard Hamming (Hamming, 1950). Intuitively, the hamming distance expresses the number of characters that either enter or leave the stage. Its application to a dramatic text is straightforward, using the function `hamming()`. By default, the hamming distance is calculated over scene boundaries.

```
hamming(rksp.0, variant="Hamming")
```

```
##  [1] 2 1 1 1 1 2 2 4 2 4 3 2 1 1 1 2 2 2 4 2 2 3 2 4 1 3 3 1 3 2 2 1 1 1 5
## [36] 3 1 1 2 2 1 2
```

The return value of the function is just a vector of numbers, one less than the number of scenes in the play. Of course, this can be plotted:

```
plot(hamming(rksp.0, variant="Hamming"),
     type="l",
     ylab="Hamming distance",
     xlab="Scenes")
```



Hamming distance, in its original definition as well as the `Hamming` variant in our function, returns absolute values. As usual, these values should be normalized, and the function `hamming()` provides two ways of doing that.

6.1.1 NormalizedHamming

Straightforwardly, one can normalise by the overall number of characters in the play. This value is only maximal, if at once scene boundary, one subset of characters enters the stage, and the entire other subset leaves the stage. As we can see below, this is not the case in our demo text.

```
hamming(rksp.0, variant="NormalizedHamming")
```

```
##  [1] 0.15384615 0.07692308 0.07692308 0.07692308 0.07692308 0.15384615
##  [7] 0.15384615 0.30769231 0.15384615 0.30769231 0.23076923 0.15384615
## [13] 0.07692308 0.07692308 0.07692308 0.15384615 0.15384615 0.15384615
## [19] 0.30769231 0.15384615 0.15384615 0.23076923 0.15384615 0.30769231
## [25] 0.07692308 0.23076923 0.23076923 0.07692308 0.23076923 0.15384615
## [31] 0.15384615 0.07692308 0.07692308 0.07692308 0.38461538 0.23076923
## [37] 0.07692308 0.07692308 0.15384615 0.15384615 0.07692308 0.15384615
```

6.1.2 Trilcke

One more possible normalization has been proposed by Trilcke et al. (2017). Instead of normalizing with all characters in the play, the Trilcke variant only normalizes with the characters in the two adjacent scenes. Thus, if all characters leave the stage, and a new set of characters enter it, the distance is maximal. This however does not have to include *all* characters in the play.

```
hamming(rksp.0, variant="Trilcke")
```

```
##  [1] 0.6666667 0.5000000 0.5000000 0.5000000 0.5000000 0.6666667 0.6666667
##  [8] 1.0000000 0.6666667 1.0000000 0.7500000 0.6666667 0.5000000 0.3333333
## [15] 0.3333333 0.5000000 0.5000000 0.6666667 1.0000000 0.6666667 0.6666667
## [22] 0.7500000 0.5000000 0.8000000 0.3333333 0.7500000 0.7500000 0.3333333
## [29] 0.7500000 0.6666667 0.6666667 0.3333333 0.3333333 0.3333333 1.0000000
```

```
## [36] 1.0000000 0.5000000 0.5000000 0.6666667 0.6666667 0.5000000 0.5000000
```

The Trilcke variant is the default setting of the `hamming()`-function.

Chapter 7

Text Collections

So far, we have mostly processed single texts. In this chapter, we will cover the important things to know when processing collections of texts – be it genres or the works of an author.

The following code can only be reproduced if the GerDraCor data set and the collection repository have been installed. This can be done with the following code (assuming an internet connection).

```
installData("gdc")
```

```
## Pulling new data from https://github.com/quadrama/data_gdc.git.
```

```
## Already up-to-date
```

```
installCollectionData()
```

```
## Warning in utils::unzip(tf, exdir = dataDirectory): Fehler 1 während des
## Extrahierens aus Zipfile
```

```
## Warning in file.rename(from = file.path(repoDirectory, "collections"), to =
## getOption("qd.collectionDirectory")): kann Datei '/Users/reiterns/QuaDramA/
## Data2/metadata-master/collections' nicht in '/Users/reiterns/QuaDramA/
## Data2/collections' umbenennen. Grund 'No such file or directory'
```

7.1 Defining collections

Before processing, it's necessary to define a collection of texts, by assembling their ids in a list. These are considered to be sets of plays without internal structure (e.g., no play is marked as prototypical).

```
# load ids that are comedies and tragedies
ids.tragedy <- unique(loadSet(c("trauerspiel", "tragödie", "tragoedie")))
ids.comedy <- unique(loadSet(c("lustspiel", "komödie", "komoedie")))

# make sure we're using the GerDraCor corpus
ids.tragedy <- sub("tg:", "gdc:", ids.tragedy, fixed = TRUE)
ids.comedy <- sub("tg:", "gdc:", ids.comedy, fixed = TRUE)
```

7.2 Loading multiple plays

Loading multiple plays at once is straightforward, and not different from loading a single one.

Part II

Text Analysis

Chapter 8

Word Field Analysis

What we consider a word field here may differ from specific uses in linguistics. In this context, a word field is a list of words that belong to a common theme / topic / semantic group. Multiple word fields can be assembled to create a dictionary. On a technical level, what we describe in the following works for arbitrary lists of words. A semantic relation between the words is technically not required. Thus, the following pieces of code can be used with arbitrary word lists.

For **demo purposes** (this is really a toy example), we will define the word field of `Love` as containing the words `Liebe` (love) and `Herz` (heart). In R, we can put them in a character vector, and we use lemmas:

```
wf_love <- c("liebe", "herz")
```

We will test this word field on Emilia Galotti, which should be about love.

```
data("rksp.0")
```

8.1 Single word field

The core of the word field analysis is collecting statistics about a dictionary. Therefore, we use the function called `dictionaryStatisticsSingle()` (single, because we only want to analyse a single word field):

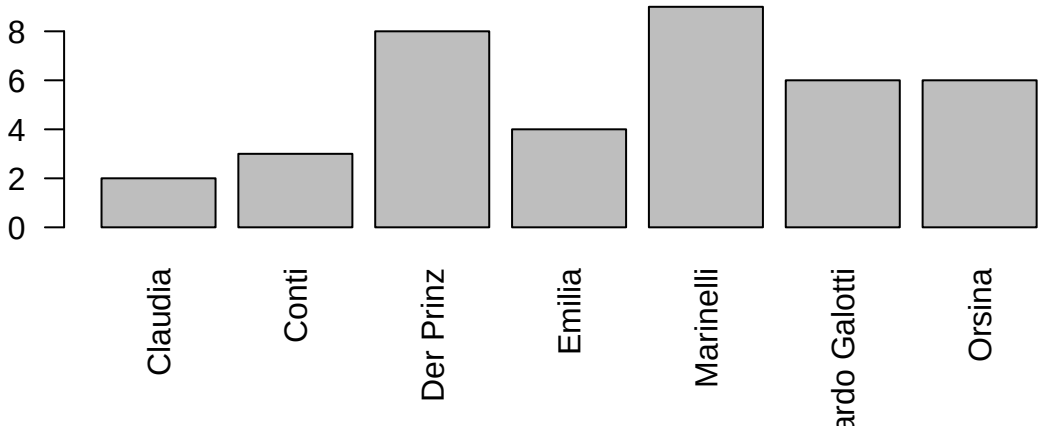
```
dictionaryStatisticsSingle(
  rksp.0,              # the text we want to process
  wordfield=wf_love    # the word field
)
```

```
##    corpus  drama        character x
## 1    test rksp.0           angelo 0
## 2    test rksp.0          appiani 0
## 3    test rksp.0         battista 0
## 4    test rksp.0     camillo_rota 0
## 5    test rksp.0  claudia_galotti 2
## 6    test rksp.0            conti 3
## 7    test rksp.0 der_kammerdiener 0
## 8    test rksp.0        der_prinz 8
## 9    test rksp.0           emilia 4
## 10   test rksp.0        marinelli 9
## 11   test rksp.0          odoardo 6
## 12   test rksp.0           orsina 6
## 13   test rksp.0            pirro 0
```

What this table shows us the number of times the characters in the play use words that appear in this list. By default, these are absolute numbers.

We can visualise these counts in a simple bar chart:

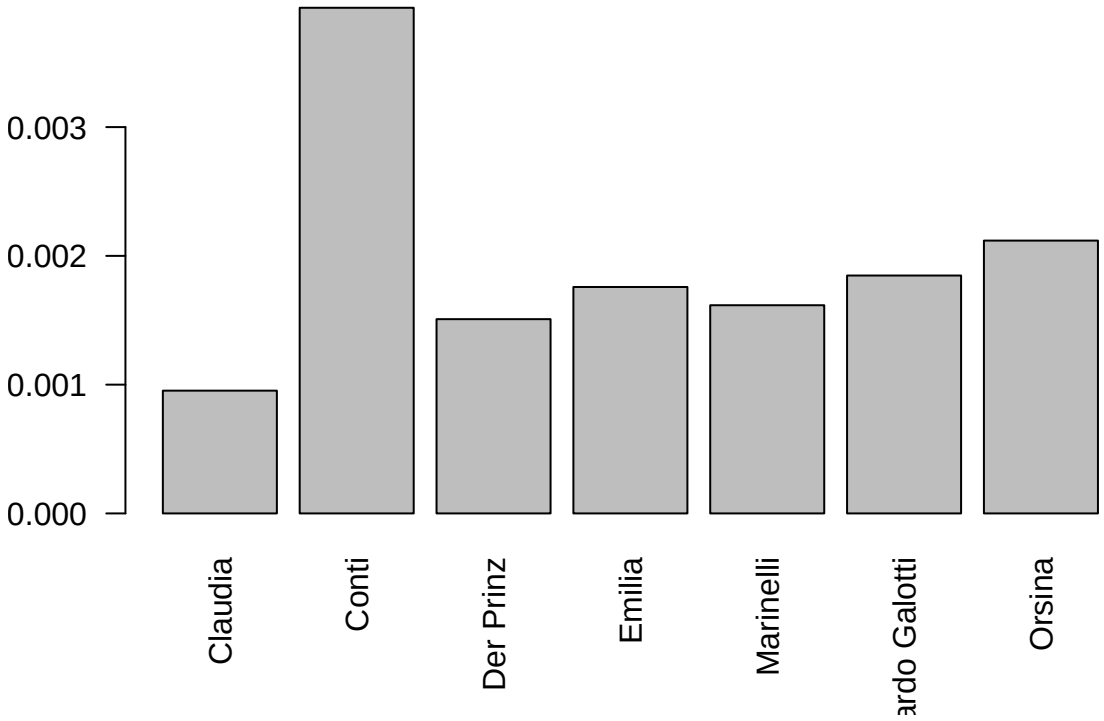
```
# retrieve counts and replace character ids by names
dstat <- dictionaryStatisticsSingle(rksp.0, wordfield = wf_love) %>%
  format(rksp.0)

# remove figures not using these words at all
dstat <- dstat[dstat$x>0,]

# create a bar plot
barplot(dstat$x,                        # what to plot
        names.arg = dstat$character,    # x axis labels
        las=2                           # turn axis labels
)
```



Apparently, the prince and Marinelli are mentioning these words a lot more than the other characters.

When comparing characters, it often (but not always) makes sense to normalize the counts according to the total number of spoken words by a character. This can be enabled by setting the argument `normalizeByFigure=TRUE`. This will divide the number of words in this field by the total number of words a character speaks.

```
dstat <- dictionaryStatisticsSingle(rksp.0, wordfield=wf_love,
  normalizeByFigure = TRUE   # apply normalization
) %>% format(rksp.0)         # reformat character names

# remove figures not using these words at all
dstat <- dstat[dstat$x>0,]

barplot(dstat$x,
        names.arg = dstat$character,
        las=2
)
```



8.2 Multiple Word Fields

The function `dictionaryStatistics()` can be used to analyse multiple dictionaries at once. To this end, dictionaries are represented as lists of character vectors. The (named) outer list contains the keywords, the vectors are just words associated with the keyword.

New dictionaries can be easily created like this:

```
wf <- list(Liebe=c("liebe","herz","schmerz"), Hass=c("hass","hassen"))
```

This dictionary contains the two entries `Liebe` (love) and `Hass` (hate), with 3 respective 2 entries. Dictionaries can be created in code, like shown above. In addition, the function `loadFields()` can be used to download dictionary content from a URL or a directory. By default, the function loads this dictionary from GitHub (that we used in publications), for the keywords `Liebe` and `Familie` (family). Since version 2.3.0, this dictionary is included in the package as `base_dictionary` and can be used right away (without internet connection). It is also the default dictionary used by `dictionaryStatistics()`.

The function `loadFields()` offers parameters to load from different URLs via http or to load from plain text files that are stored locally. The latter can be achieved by specifying the directory as `baseurl`. Entries for each keyword should then be stored in a file named like the keyword, and ending with `txt` (by default, can be overwritten). See `?loadFields` for details. Some of the options can also be specified through `dictionaryStatistics()`, as exemplified below.

The following examples use the `base_dictionary`, i.e., a specific version of the fields we have been using in QuaDramA.

```
dstat <- dictionaryStatistics(
  rksp.0,  # the text
  fieldnames = # fields we want to measure (from base_dictionary)
    c("Liebe", "Familie", "Ratio", "Krieg", "Religion"),
  normalizeByFigure = TRUE,   # normalization by figure
  normalizeByField = TRUE     # normalization by field
```

```
)
dstat
```

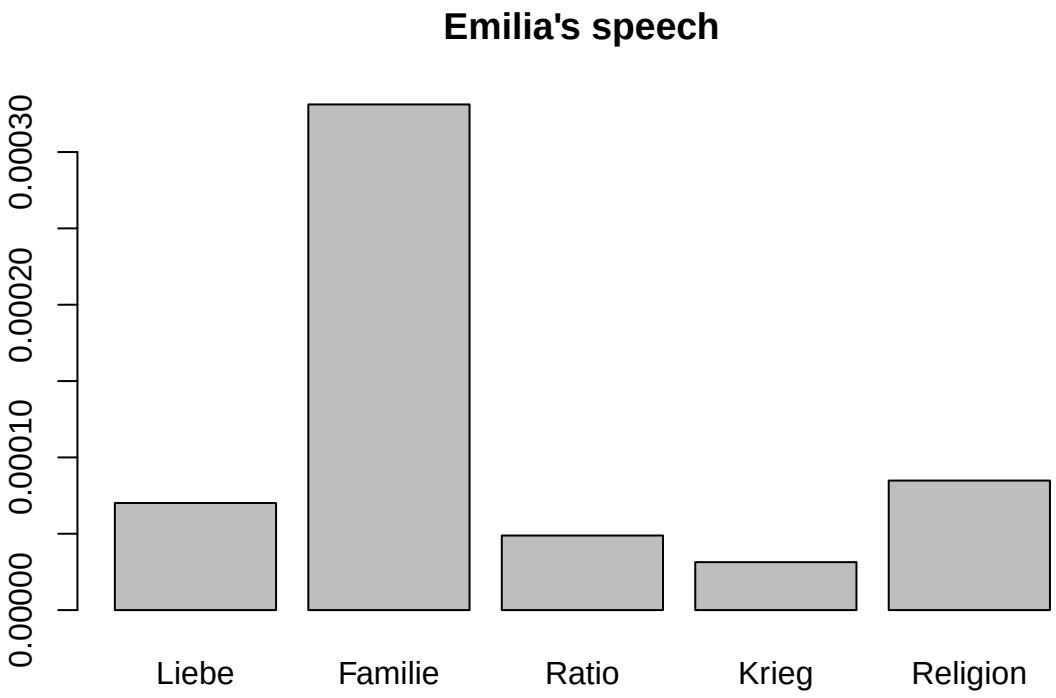
```
##    corpus  drama        character        Liebe      Familie        Ratio
## 1    test rksp.0           angelo 5.025967e-05 4.314529e-05 7.290755e-05
## 2    test rksp.0          appiani 8.610133e-05 1.355080e-04 6.661338e-05
## 3    test rksp.0         battista 0.000000e+00 1.404988e-04 0.000000e+00
## 4    test rksp.0     camillo_rota 1.003613e-04 1.292324e-04 4.367575e-04
## 5    test rksp.0  claudia_galotti 6.591891e-05 2.350575e-04 3.972037e-05
## 6    test rksp.0            conti 1.531692e-04 5.379043e-05 3.635835e-05
## 7    test rksp.0 der_kammerdiener 0.000000e+00 0.000000e+00 0.000000e+00
## 8    test rksp.0        der_prinz 7.021317e-05 6.974600e-05 4.714313e-05
## 9    test rksp.0           emilia 7.014262e-05 3.311757e-04 4.884005e-05
## 10   test rksp.0        marinelli 8.026020e-05 1.451804e-04 3.991777e-05
## 11   test rksp.0          odoardo 5.895608e-05 2.319657e-04 6.271666e-05
## 12   test rksp.0           orsina 5.259046e-05 8.223048e-05 4.250366e-05
## 13   test rksp.0            pirro 0.000000e+00 1.597508e-04 2.699492e-05
##           Krieg     Religion
## 1  5.624297e-05 0.000000e+00
## 2  4.817575e-05 0.000000e+00
## 3  0.000000e+00 0.000000e+00
## 4  8.423181e-05 0.000000e+00
## 5  4.681329e-05 5.853528e-05
## 6  2.337322e-05 0.000000e+00
## 7  0.000000e+00 0.000000e+00
## 8  2.862261e-05 3.308290e-05
## 9  3.139717e-05 8.482745e-05
## 10 3.528446e-05 1.890842e-05
## 11 4.123417e-05 4.861291e-05
## 12 2.522195e-05 1.238973e-05
## 13 0.000000e+00 1.022966e-04
```

The variable `dstat` now contains multiple columns, one for each word field. We have been using the option `normalizeByFigure` before. When comparing multiple fields, it often happens that they have a different size (i.e., different number of words). In this case, it makes sense to *also* normalize with the number of words in the word field. This is achieved by `normalizeByField=TRUE`. This makes the numbers very small, but they should be used in comparison anyway.

8.2.1 Bar plot by character

It is now straightforward to show the distribution of fields for a single character:

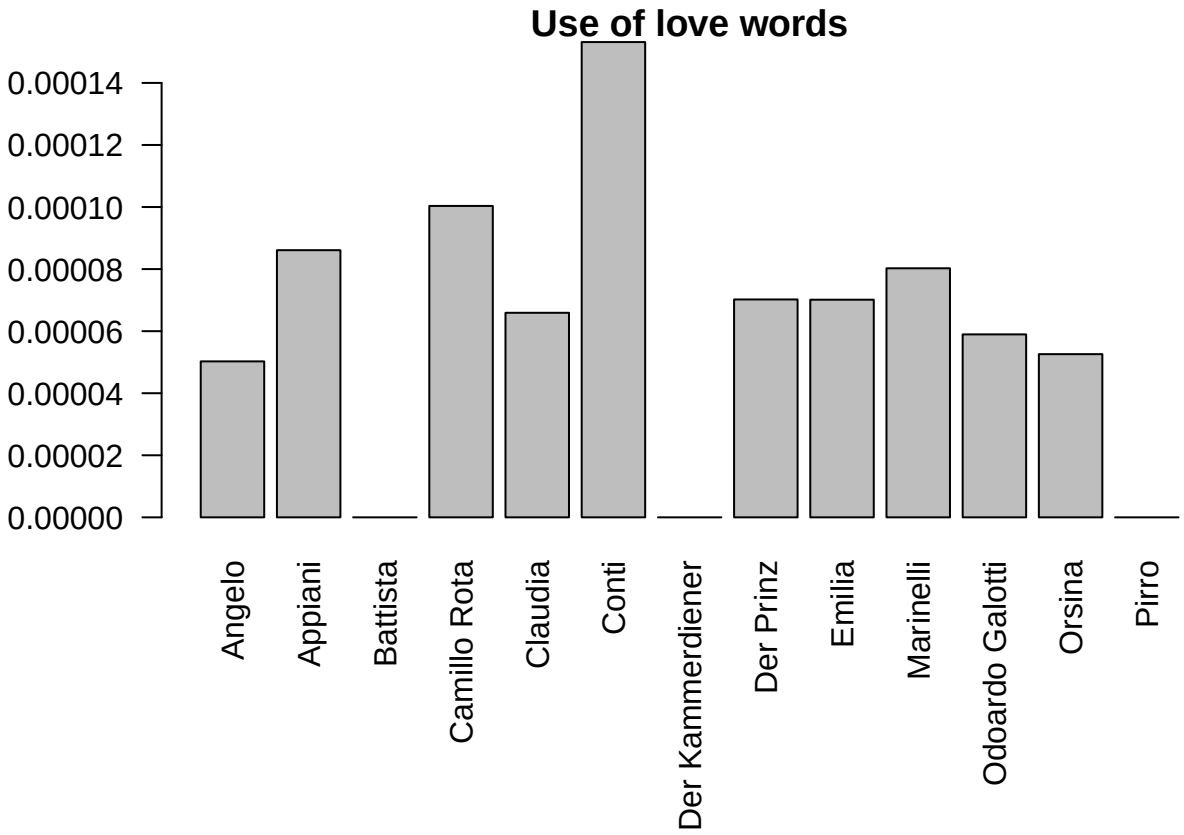
```
mat <- as.matrix(dstat)

barplot(mat[9,],     # we select Emilia's line
        main="Emilia's speech",
        names.arg=colnames(mat)
)
```



8.2.2 Bar plot by field

Conversely, we can also show who uses words of a certain field how often:

```
dstat <- dictionaryStatistics(
  rksp.0,  # the text
  fieldnames = # fields we want to measure (from base_dictionary)
    c("Liebe", "Familie", "Ratio", "Krieg", "Religion"),
  normalizeByFigure = TRUE,    # normalization by figure
  normalizeByField = TRUE      # normalization by field
) %>%
  format(rksp.0)

mat <- as.matrix(dstat)

par(mar=c(9,4,1,1))

barplot(mat[,1],                  # Select the row for 'love'
        main="Use of love words", # title for plot
        beside = TRUE,            # not stacked
        names.arg = dstat$character,   # x axis labels
        las=2                     # rotation for labels
)
```



8.3 Dictionary Based Distance

Technically, the output of `dictionaryStatistics()` is a `data.frame`. This is suitable for most uses. In some cases, however, it's more suited to work with a matrix that only contains the raw numbers (i.e., number of family words). Calculating character distance based on dictionaries, for instance. For these cases, the package provides an S3 method that extracts the numeric part of the data.frame and creates a matrix. We have used this function `as.matrix()` already above.

The matrix doesn't have row names by default. The snippet below can be used to add row names.

```
ds <- dictionaryStatistics(rksp.0,
                           fieldnames=c("Liebe", "Familie", "Ratio", "Krieg", "Religion"),
                           normalizeByFigure=TRUE)
m <- as.matrix(ds)
rownames(m) <- ds$character
m
```

```
##                       Liebe     Familie       Ratio       Krieg
## angelo           0.004724409 0.003149606 0.007874016 0.006299213
## appiani          0.008093525 0.009892086 0.007194245 0.005395683
## battista         0.000000000 0.010256410 0.000000000 0.000000000
## camillo_rota     0.009433962 0.009433962 0.047169811 0.009433962
## claudia_galotti  0.006196378 0.017159199 0.004289800 0.005243089
## conti            0.014397906 0.003926702 0.003926702 0.002617801
## der_kammerdiener 0.000000000 0.000000000 0.000000000 0.000000000
## der_prinz        0.006600038 0.005091458 0.005091458 0.003205733
## emilia           0.006593407 0.024175824 0.005274725 0.003516484
## marinelli        0.007544458 0.010598168 0.004311119 0.003951859
```

```
## odoardo          0.005541872 0.016933498 0.006773399 0.004618227
## orsina           0.004943503 0.006002825 0.004590395 0.002824859
## pirro            0.000000000 0.011661808 0.002915452 0.000000000
##                      Religion
## angelo           0.0000000000
## appiani          0.0000000000
## battista         0.0000000000
## camillo_rota     0.0000000000
## claudia_galotti  0.0033365110
## conti            0.0000000000
## der_kammerdiener 0.0000000000
## der_prinz        0.0018857251
## emilia           0.0048351648
## marinelli        0.0010777798
## odoardo          0.0027709360
## orsina           0.0007062147
## pirro            0.0058309038
```

Every character is now represented with five numbers, which can be interpreted as a vector in five-dimensional space. This means, we can easily apply distance metrics supplied by the function `dist()` (from the default package `stats`). By default, `dist()` calculates Euclidean distance.

```
cdist <- dist(m)
# output not shown
```

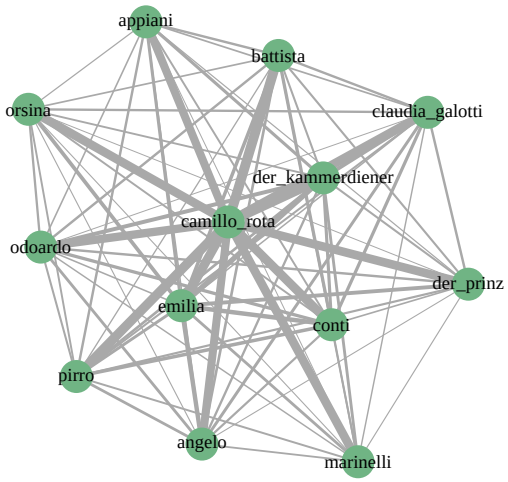
The resulting data structure is similar to the one in the weighted configuration matrix, which means everything from Section 5.2.2 can be applied here. In particular, we can convert these distances into a network:

```
require(igraph)
g <- graph_from_adjacency_matrix(as.matrix(cdist),
                                 weighted=TRUE,     # weighted graph
                                 mode="undirected", # no direction
                                 diag=FALSE         # no looping edges
                                )
```

This network can of course be visualised again.

```
# Now we plot
plot.igraph(g,
            layout=layout_with_fr,       # how to lay out the graph
            main="Dictionary distance network",  # title
            vertex.label.cex=0.6,         # label size
            vertex.label.color="black",   # font color
            vertex.color=qd.colors[5],    # vertex color
            vertex.frame.color=NA,        # no vertex border
            edge.width=E(g)$weight*100    # scale edges according to their weight
                                          # (since the distances are so small, we multiply)
            )
```

Dictionary distance network



Although this network is similar to the one shown in Section 5.2.2 (both undirected and weighted), it displays a totally different kind of information. The networks based on copresence connect characters that appear together on stage, while this network connects characters with similar thematic profile in their speech (within the limits of being able to detect thematic profiles via word fields).

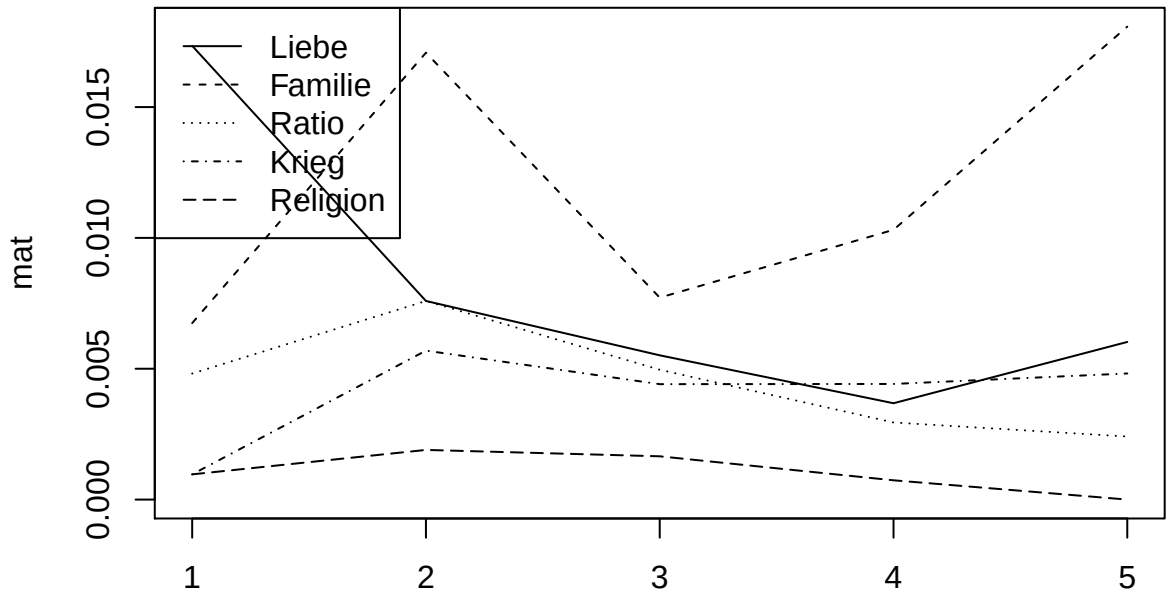
8.4 Development over the course of the play

Finally, the function `dictionaryStatistics()` can be used to track word field for segments of the play. To do that, we use the parameter `segment`, and set it to either "Act" or "Scene".

8.4.1 Individual characters

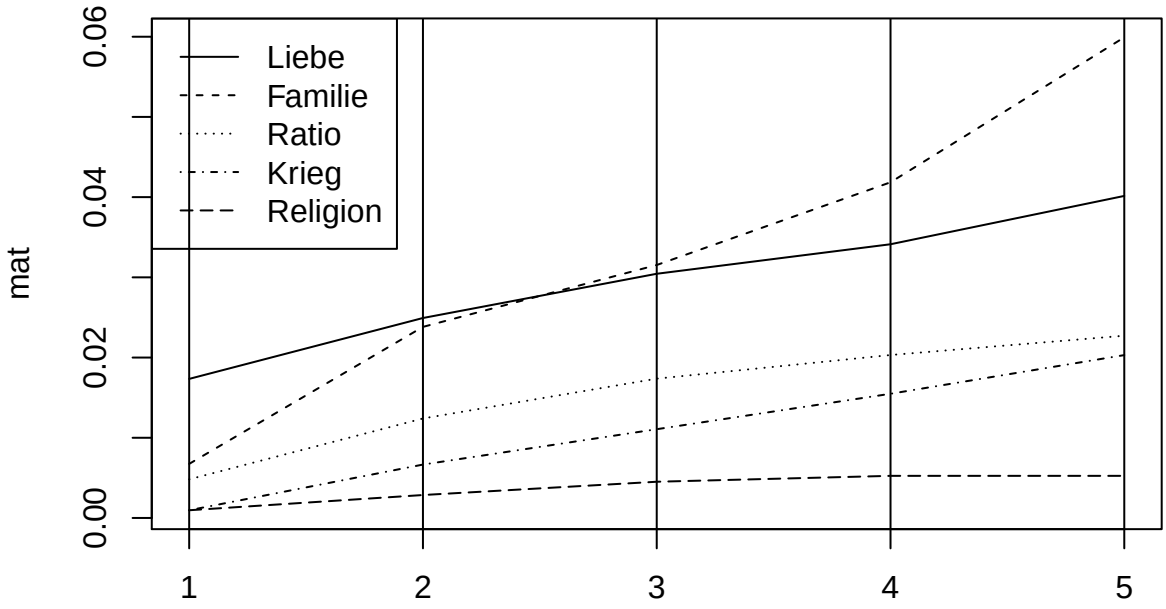
We can now plot the theme progress over the course of the play. This can be done for specific characters, as shown below.

```
dsl <- dictionaryStatistics(rksp.0,
                            fieldnames=c("Liebe", "Familie", "Ratio", "Krieg", "Religion"),
                            normalizeByFigure=TRUE,
                            segment="Act")

mat <- as.matrix(dsl[dsl$character=="marinelli",])

matplot(mat, type="l", col="black")
legend(x="topleft",legend=colnames(mat), lty=1:ncol(dsl))
```



Depending on the use case, it might be easier to interpret if the numbers are cumulatively added up. The snippet below shows how this works.

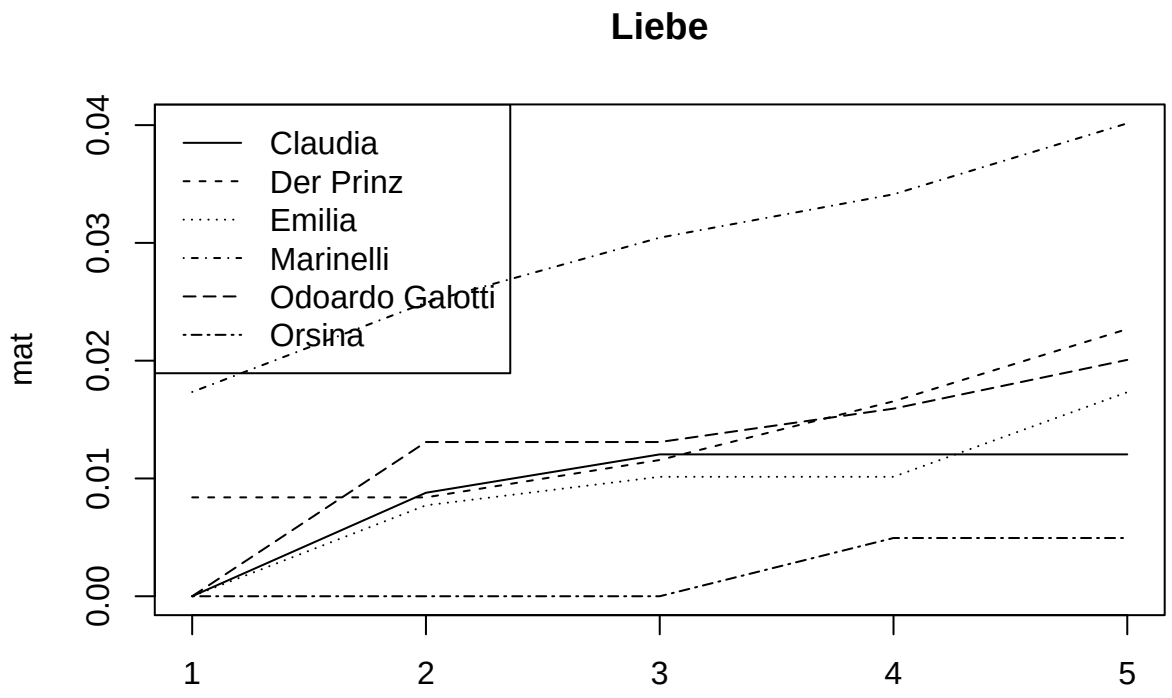
```
dsl <- dictionaryStatistics(rksp.0,
                            fieldnames=c("Liebe", "Familie", "Ratio", "Krieg", "Religion"),
                            normalizeByFigure=TRUE,
                            segment="Act")

mat <- as.matrix(dsl[dsl$character=="marinelli",])

mat <- apply(mat,2,cumsum)

matplot(mat, type="l", col="black")

# Add act lines
abline(v=1:nrow(mat))

legend(x="topleft", legend=colnames(mat), lty=1:5)
```



8.4.2 Comparing characters

Simultaneously to the setting above, we now want to compare the development of several characters for a single word field. This unfortunately requires some reshuffling of the data, using the function `reshape` (from the `stats` package).

```
dsl <- dictionaryStatistics(rksp.0,
                            fieldnames=c("Liebe"),
                            normalizeByFigure=TRUE,
                            segment="Act") %>%
  filter(rksp.0,
         threshold = 6) %>%
  format(rksp.0)

dsl <- reshape(dsl, direction = "wide",  # the table becomes wider
               timevar = "Number.Act",   # the column that specifies multiple readings
               times = "Liebe",          # the number to distribute
               idvar=c("corpus","drama","character")   # what identifies a character
)

mat <- as.matrix.data.frame(dsl[,4:ncol(dsl)])
rownames(mat) <- dsl$character
mat <- apply(mat,1,cumsum)

matplot(mat, type="l", lty = 1:ncol(mat), col="black", main="Liebe")
legend(x="topleft", legend=colnames(mat), lty=1:ncol(mat))
```


Liebe
mat
0.00
0.01
0.02
0.03
0.04
1
2
3
4
5
Claudia
Der Prinz
Emilia
Marinelli
Odoardo Galotti
Orsina

Chapter 9

Advanced Text Analysis

9.1 When are characters mentioned?

When characters are speaking on stage, they are actively present. But they can also be passively present, if other characters refer to them. Both levels of presence can be extracted with the `presence()` function:

```
# Load Emilia Galotti
data(rksp.0)

pres <- presence(rksp.0)
pres
```

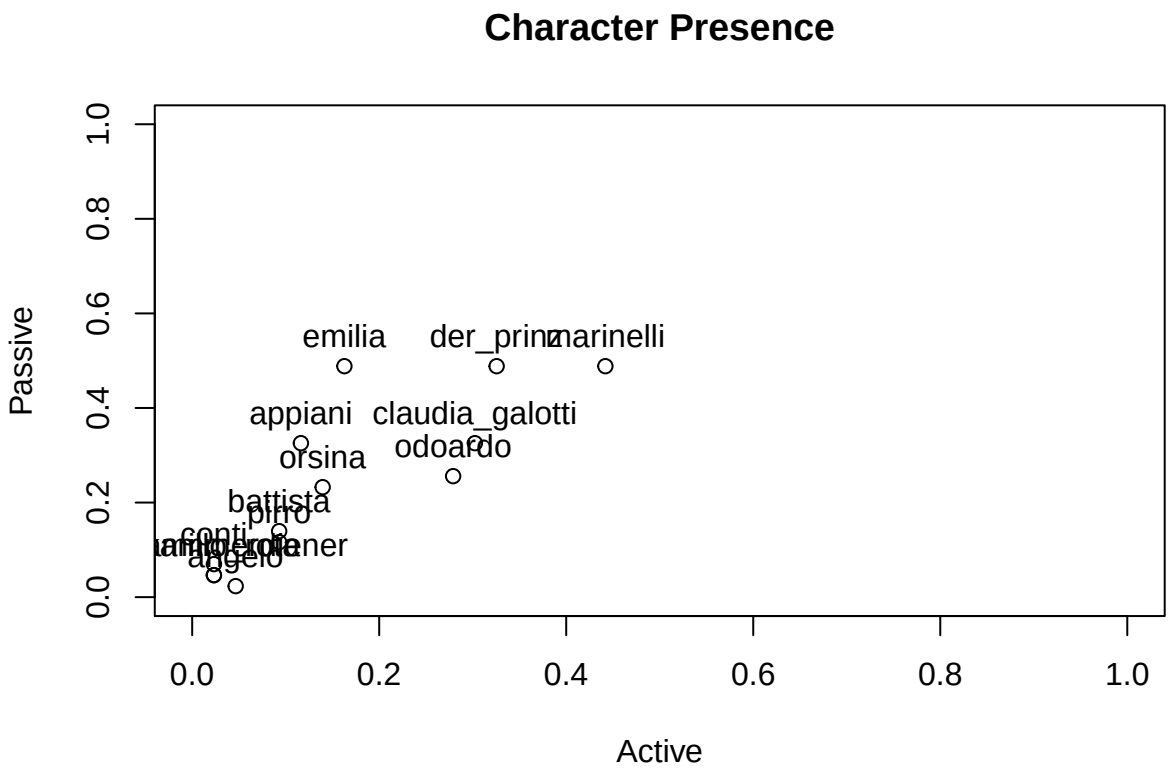
```
##    corpus  drama        character scenes actives passives    presence
## 1    test rksp.0           angelo     43       2        1  0.02325581
## 2    test rksp.0          appiani     43       5       14 -0.20930233
## 3    test rksp.0         battista     43       4        6 -0.04651163
## 4    test rksp.0     camillo_rota     43       1        2 -0.02325581
## 5    test rksp.0  claudia_galotti     43      13       14 -0.02325581
## 6    test rksp.0            conti     43       1        3 -0.04651163
## 7    test rksp.0 der_kammerdiener     43       1        2 -0.02325581
## 8    test rksp.0        der_prinz     43      14       21 -0.16279070
## 9    test rksp.0           emilia     43       7       21 -0.32558140
## 10   test rksp.0        marinelli     43      19       21 -0.04651163
## 11   test rksp.0          odoardo     43      12       11  0.02325581
## 12   test rksp.0           orsina     43       6       10 -0.09302326
## 13   test rksp.0            pirro     43       4        5 -0.02325581
```

As we can see, each character has a few numbers associated: The column `actives` shows the number of scenes in which the character is actively present. This is equivalent to the information in the configuration matrix. The column `passives` shows the number of scenes in which a character is mentioned. By default, this excludes the scenes in which they are present themselves (this behaviour can be changed by adding the parameter `passiveOnlyWhenNotActive = TRUE` to the call of the `presence` function).

A simple visualisation that shows the characters active and passive presence in one plot can be generated like this: The first line (`plot()`) is responsible for the plotting of the symbols, the second line (`text()`) adds the character names or ids numbers.

```
plot(x=pres$active/pres$scenes,
     y=pres$passive/pres$scenes,
     xlim=c(0,1),
     ylim=c(0,1),
```

```
     xlab="Active",
     ylab="Passive",
     main="Character Presence")
text(x=pres$actives/pres$scenes,
     y=pres$passives/pres$scenes,
     labels=substr(pres$character,0,20),
     pos=3)
```



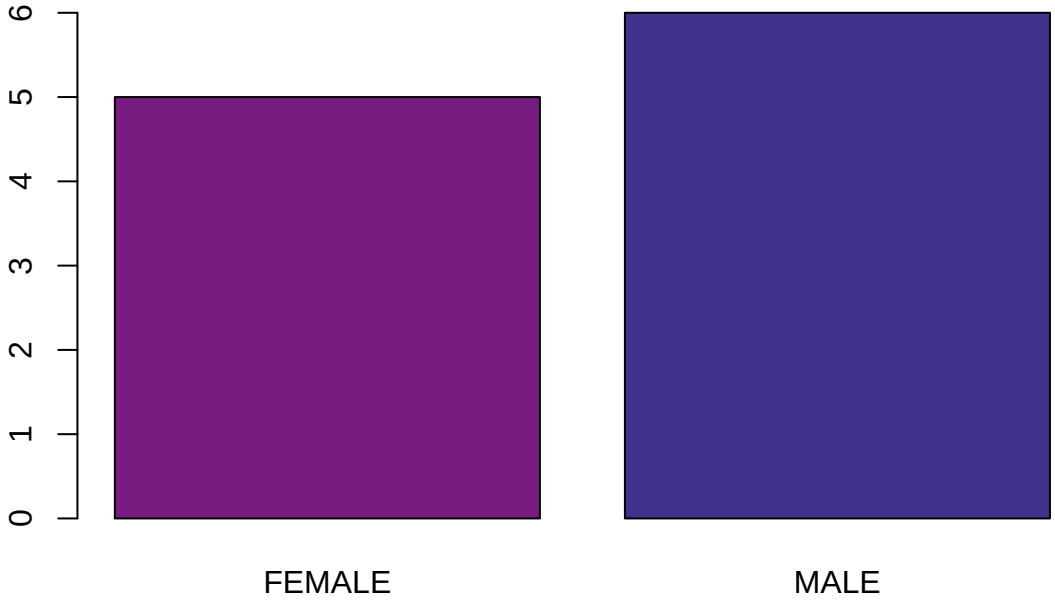
Chapter 10

Resterampe

10.0.1 Character meta data

We will now combine this information with additional meta data about characters, i.e., gender.

```
# Proportion of male / female characters
barplot(table(text$characters$Gender),col=qd.colors)
```

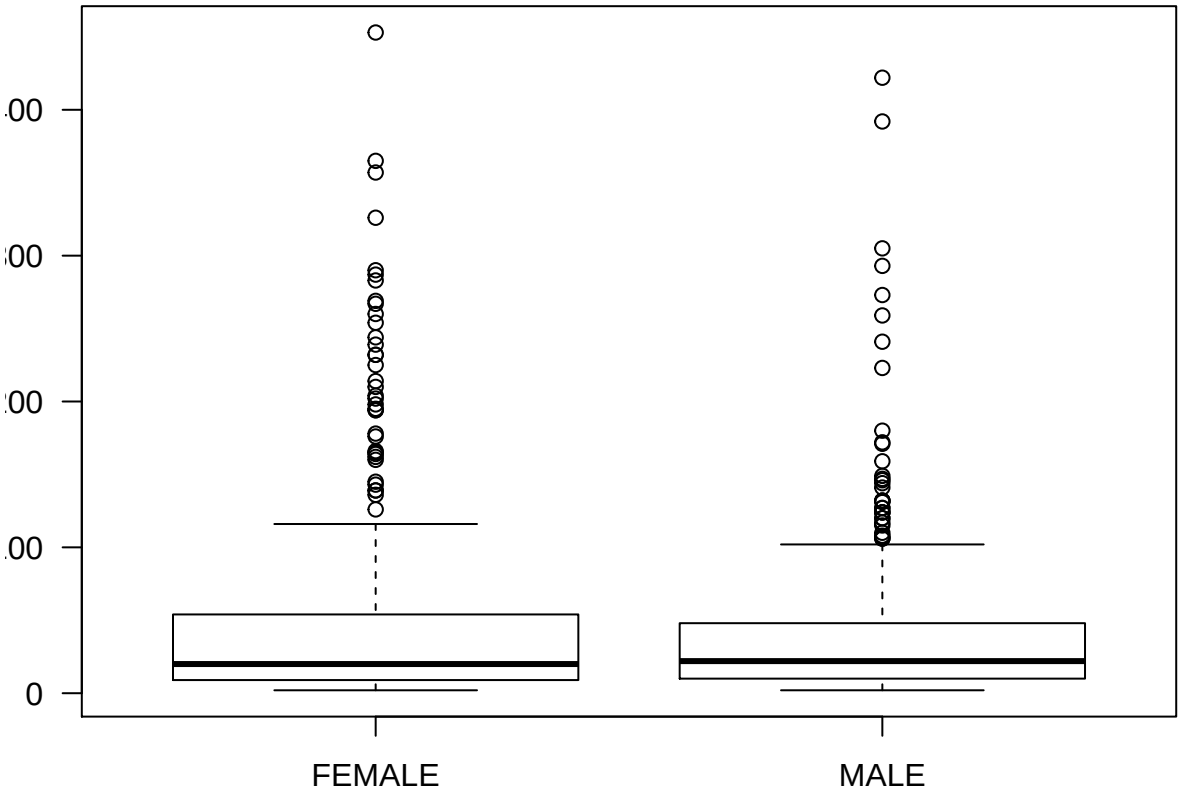


10.0.2 Character groups

Next, we want to make the same analysis not for individual characters, but for character groups, based on categories such as gender.

```
ustat <- utteranceStatistics(rjmw.0,
                             normalizeByDramaLength = FALSE # use absolute values
                            )

characterdata <- rjmw.0$characters

ustat <- merge(ustat, characterdata,
               by.x = c("corpus","drama", "character"),
```

```
              by.y = c("corpus","drama", "figure_id"))

par(mar=c(2,2,2,2))
boxplot(utteranceLength ~ Gender, # what do we want to correlate
        data=ustat,
        las = 1 # rotate axis labels
      )
```



According to this picture, female characters speak slightly longer utterances in this play.

Bibliography

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