

Tutorial 1: Processing of textual data

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2020-07-03

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In this tutorial, we demonstrate how to read text data in R, tokenize texts and create a document-term matrix.

1. Reading various file formats with the `readtext` package,
2. From text to a corpus,
3. Create a document-term matrix and investigate Zipf's law

First, let's create a new R Project (File -> New Project -> Existing directory) in the provided tutorial folder. Then we create a new R File (File -> New File -> R script) and save it as "Tutorial_1.R".

Reading txt, pdf, html, docx, ...

In case you have already a collection of document files on your disk, you can import them into R in a very convenient way provided by the `readtext` package. The package depends on some other programs or libraries in your system, e.g. to provide extraction of text from Word- and PDF-documents.

Hence, some persons encountered hurdles to install the package due to missing libraries. In this case, carefully read error messages and install the missing libraries.

For demonstration purpose, we provide in `data/documents` a random selection of documents in various file formats. First, we request a list of files in the directory to extract text from.

```
data_files <- list.files(path = "data/documents", full.names = T, recursive = T)

# View first file paths
head(data_files, 3)
```

```
## [1] "data/documents/bundestag/17_16_580-F_neu.pdf"
## [2] "data/documents/bundestag/prot_17_95.pdf"
## [3] "data/documents/bundestag/stellungnahme---buendnis-buergerenergie-e--v--data.pdf"
```

The `readtext` function from the package with the same name, detects automatically file formats of the given files list and extracts the content into a `data.frame`. The parameter `docvarsfrom` allows you to set metadata variables by splitting path names. In our example, `docvar3` contains a source type variable derived from the sub folder name.

```
require(readtext)

extracted_texts <- readtext(data_files, docvarsfrom = "filepaths", dvsep = "/")

# View first rows of the extracted texts
head(extracted_texts)

# View beginning of the second extracted text
cat(substr(extracted_texts$text[2], 0, 300))
```

Again, the `extracted_texts` can be written by `write.csv2` to disk for later use.

```
write.csv2(extracted_texts, file = "data/text_extracts.csv", fileEncoding = "UTF-8")
```

We choose CSV as a convenient text column based format for easy import and export in R and other programs. Also our example data for the rest of the tutorials is provided as CSV file. Windows users: Take care of setting UTF-8 file encoding explicitly when writing text data to the hard drive.

From text to a corpus object

Set global options at the beginning of each script! When working with textual data strings, it is recommended to turn R's automatic conversion of strings into factors off.

```
# Global options
options(stringsAsFactors = FALSE)
```

The `read.csv` command reads a **CSV** (Comma Separated Value) file from disk. Such files represent a table whose rows are represented by single lines in the files and columns are marked by a *separator* character within lines. Arguments of the command can be set to specify whether the CSV file contains a line with column names (`header = TRUE` or `FALSE`) and the character set.

We read a CSV containing 233 “State of the Union” addresses of the presidents of the United States. The texts are freely available from <http://stateoftheunion.onetwothree.net>.

Our CSV file has the format: `"doc_id";"speech_type";"president";"date";"text"`. Text is encapsulated into quotes (`"`). Since separation is marked by `;` instead of `,`, we need to specify the separator char.

```
# read csv into a data.frame
textdata <- read.csv("data/sotu.csv", header = TRUE, sep = ";", encoding = "UTF-8")
```

The texts are now available in a `data.frame` together with some metadata (document id, speech type, president). Let us first see how many documents and metadata we have read.

```
# dimensions of the data frame
dim(textdata)
```

```
## [1] 233 5
```

```
# column names of text and metadata
colnames(textdata)
```

```
## [1] "doc_id" "speech_type" "president" "date" "text"
```

How many speeches do we have per president? This can easily be counted with the command `table`, which can be used to create a cross table of different values. If we apply it to a column, e.g. *president* of our data frame, we get the counts of the unique *president* values.

```
table(textdata[, "president"])
```

```
##
##      Abraham Lincoln      Andrew Jackson      Andrew Johnson
##              4              8              4
##      Barack Obama      Benjamin Harrison      Calvin Coolidge
##              8              4              6
##      Chester A. Arthur      Donald J. Trump      Dwight D. Eisenhower
##              4              3              9
##      Franklin D. Roosevelt      Franklin Pierce      George H.W. Bush
##              12              4              4
```

Now we want to transfer the loaded text source into a **corpus object** of the **quanteda**-package. Quanteda provides a large number of highly efficient convenience functions to process text in R [1]. First we load the package.

```
require(quanteda)
```

A corpus object is created with the `corpus` command. As parameter, the command gets the fulltext of the documents. In our case, this is the `text`-column of the `textdata`-data.frame. The `docnames`-parameter of the `corpus` function defines which unique identifier is given to each text example in the input (values from other columns of the data frame could be imported as metadata to each document but we will not use them in this tutorial).

```
sotu_corpus <- corpus(textdata$text, docnames = textdata$doc_id)
# have a look on the new corpus object
summary(sotu_corpus)
```

```
## Corpus consisting of 233 documents, showing 100 documents:
```

```
##
## Text Types Tokens Sentences
## 1 460 1167 24
## 2 593 1504 40
## 3 816 2476 60
## 4 772 2287 61
## 5 803 2121 56
```

##	6	1137	3197	79
##	7	821	2154	53
##	8	1005	3095	78
##	9	732	2236	58
##	10	833	2368	54
##	11	597	1624	35
##	12	562	1489	40
##	13	1098	3493	90
##	14	823	2397	62
##	15	837	2463	48
##	16	746	2286	50
##	17	984	3171	78
##	18	969	3113	76
##	19	858	2606	62
##	20	956	2921	72
##	21	699	1950	40
##	22	893	2615	61
##	23	829	2439	46
##	24	1126	3495	87
##	25	1074	3521	67
##	26	831	2282	50
##	27	1050	3389	59
##	28	1086	3614	72
##	29	1222	4814	122
##	30	1243	4728	114
##	31	1196	5091	131
##	32	1070	3762	84
##	33	1319	6323	149
##	34	1265	5145	117
##	35	1593	6926	184
##	36	1750	9175	247
##	37	2147	9798	212
##	38	1817	8406	172
##	39	1819	7578	165
##	40	1832	7937	195
##	41	2356	11408	299
##	42	2767	16271	391
##	43	1792	7716	165
##	44	1916	8476	199
##	45	1741	8494	181
##	46	2581	14454	318
##	47	2240	11567	239
##	48	2421	13212	294
##	49	2368	12341	281
##	50	2388	12426	264
##	51	2638	14498	341
##	52	2153	9700	182
##	53	2002	8836	197
##	54	2037	9018	205
##	55	1958	8635	193
##	56	2115	10019	267
##	57	2689	17517	446
##	58	2782	19849	476
##	59	2687	17782	440

##	60	3289	23273	597
##	61	1863	8222	210
##	62	2061	9004	232
##	63	2696	14328	351
##	64	2379	10705	284
##	65	2292	10375	231
##	66	2427	10985	267
##	67	2458	12581	274
##	68	2325	11379	255
##	69	2575	14929	406
##	70	2912	17828	518
##	71	2453	13596	396
##	72	2638	15360	484
##	73	1872	7651	213
##	74	2061	9367	319
##	75	1719	6643	200
##	76	1755	6585	207
##	77	2153	10077	276
##	78	1871	7790	202
##	79	2668	13156	381
##	80	2352	10846	281
##	81	1924	8481	247
##	82	2173	9577	266
##	83	1758	7016	211
##	84	1162	4336	112
##	85	2377	11007	292
##	86	2143	10074	273
##	87	2574	13397	347
##	88	1787	7429	192
##	89	1965	8706	218
##	90	2040	8641	237
##	91	2443	12643	313
##	92	1768	7209	191
##	93	1351	4095	118
##	94	1062	3295	92
##	95	1315	4079	108
##	96	2355	9667	288
##	97	3813	21377	559
##	98	3110	16350	416
##	99	1394	5722	123
##	100	2402	9794	239

A corpus is an extension of R list objects. With the `[[]]` brackets, we can access single list elements, here documents, within a corpus. We print the text of the first element of the corpus using the `texts` command.

```
# getting a single text documents content
cat(texts(sotu_corpus[1]))
```

```
## Fellow-Citizens of the Senate and House of Representatives:
##
## I embrace with great satisfaction the opportunity which now presents itself
## of congratulating ...
```

The command `cat` prints a given character vector with correct line breaks (compare the difference of the output with the `print` method instead).

Success!!! We now have 233 speeches for further analysis available in a convenient `tm` corpus object!

Text statistics

A further aim of this exercise is to learn about statistical characteristics of text data. At the moment, our texts are represented as long character strings wrapped in document objects of a corpus. To analyze which word forms the texts contain, they must be **tokenized**. This means that all the words in the texts need to be identified and separated. Only in this way it is possible to count the frequency of individual word forms. A word form is also called “**type**”. The occurrence of a type in a text is a “**token**”.

For text mining, texts are further transformed into a numeric representation. The basic idea is that the texts can be represented as statistics about the contained words (or other content fragments such as sequences of two words). The list of every distinct word form in the entire corpus forms the **vocabulary** of a corpus.

For each document, we can count how often each word of the vocabulary occurs in it. By this, we get a term **frequency vector** for each document. The dimensionality of this term vector corresponds to the size of the vocabulary. Hence, the word vectors have the same form for each document in a corpus. Consequently, multiple term vectors representing different documents can be combined into a matrix. This data structure is called **document-term matrix** (DTM).

The function `dfm` (Document-Feature-Matrix; Quanteda treats words as features of a text-based dataset) of the `quanteda` package creates such a DTM. If this command is called without further parameters, the individual word forms are identified by using the tokenizer of `quanteda` as the word separator (see `help(tokens)` for details). `Quanteda` has 3 different word separation methods. The standard and smartest way uses word boundaries and punctuations to separate the text sources. The other methods rely on whitespace information and work significantly faster but not as accurate.

```
# Create a DTM (may take a while)
DTM <- dfm(sotu_corpus)
# Show some information
DTM

## Document-feature matrix of: 233 documents, 30,358 features (94.3% sparse).
##      features
## docs fellow-citizens  of the senate and house representatives : i embrace
## 1           1 69 97      2 41      3           3 3 11      1
## 2           1 89 122     2 45      3           3 3 8       0
## 3           1 159 242     3 73      3           3 4 6       0
## 4           1 139 195     2 56      3           3 3 21      0
## 5           1 132 180     2 49      3           3 3 12      0
## 6           1 187 273     2 86      4           3 3 18      0
## [ reached max_ndoc ... 227 more documents, reached max_nfeat ... 30,348 more features ]

# Dimensionality of the DTM
dim(DTM)

## [1] 233 30358
```

The dimensions of the DTM, 233 rows and 30358 columns, match the number of documents in the corpus and the number of different word forms (types) of the vocabulary.

A first impression of text statistics we can get from a word list. Such a word list represents the frequency counts of all words in all documents. We can get that information easily from the DTM by summing all of its column vectors.

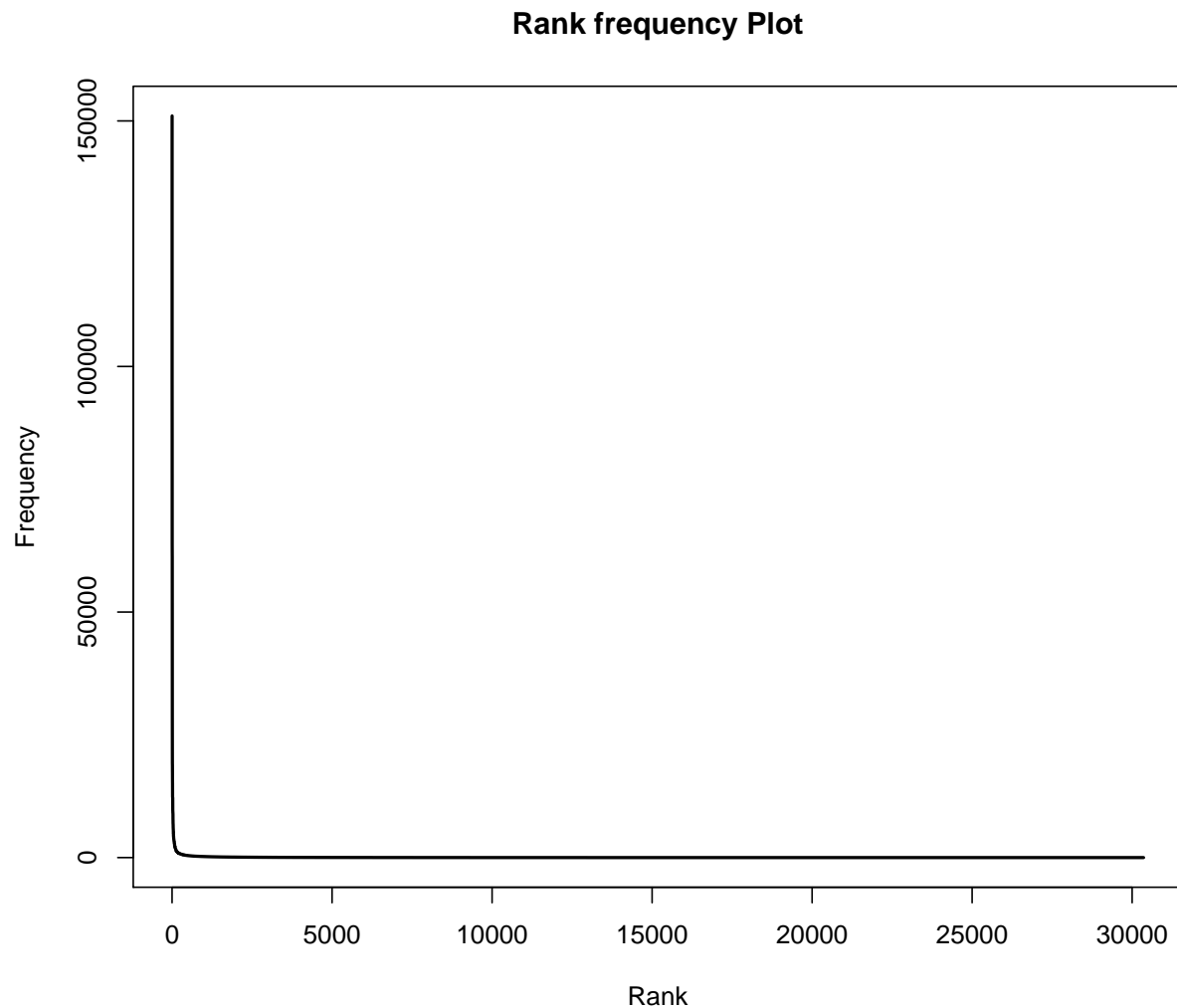
A so-called **sparse matrix** data structure is used for the document term matrix in the *quanteda* package (*quanteda* inherits the *Matrix* package for sparse matrices). Since most entries in a document term vector are 0, it would be very inefficient to actually store all these values. A sparse data structure instead stores only those values of a vector/matrix different from zero. The *Matrix* package provides arithmetic operations on sparse DTMs.

```
# sum columns for word counts
freqs <- colSums(DTM)
# get vocabulary vector
words <- colnames(DTM)
# combine words and their frequencies in a data frame
wordlist <- data.frame(words, freqs)
# re-order the wordlist by decreasing frequency
wordIndexes <- order(wordlist[, "freqs"], decreasing = TRUE)
wordlist <- wordlist[wordIndexes, ]
# show the most frequent words
head(wordlist, 25)
```

```
##      words  freqs
## the      the 151018
## of       of  97395
## ,        ,  85384
## .        .  63784
## and      and  61565
## to       to  61304
## in       in  38980
## a        a  28351
## that     that 21960
## for      for  19183
## be       be  18787
## our      our  17678
## is       is  17197
## it       it  15321
## by       by  15074
## we       we  12377
## which    which 12354
## as       as  12277
## this     this  12215
## have     have  12172
## with     with  12119
## will     will   9615
## on       on   9520
## i        i   9372
## has      has   9104
```

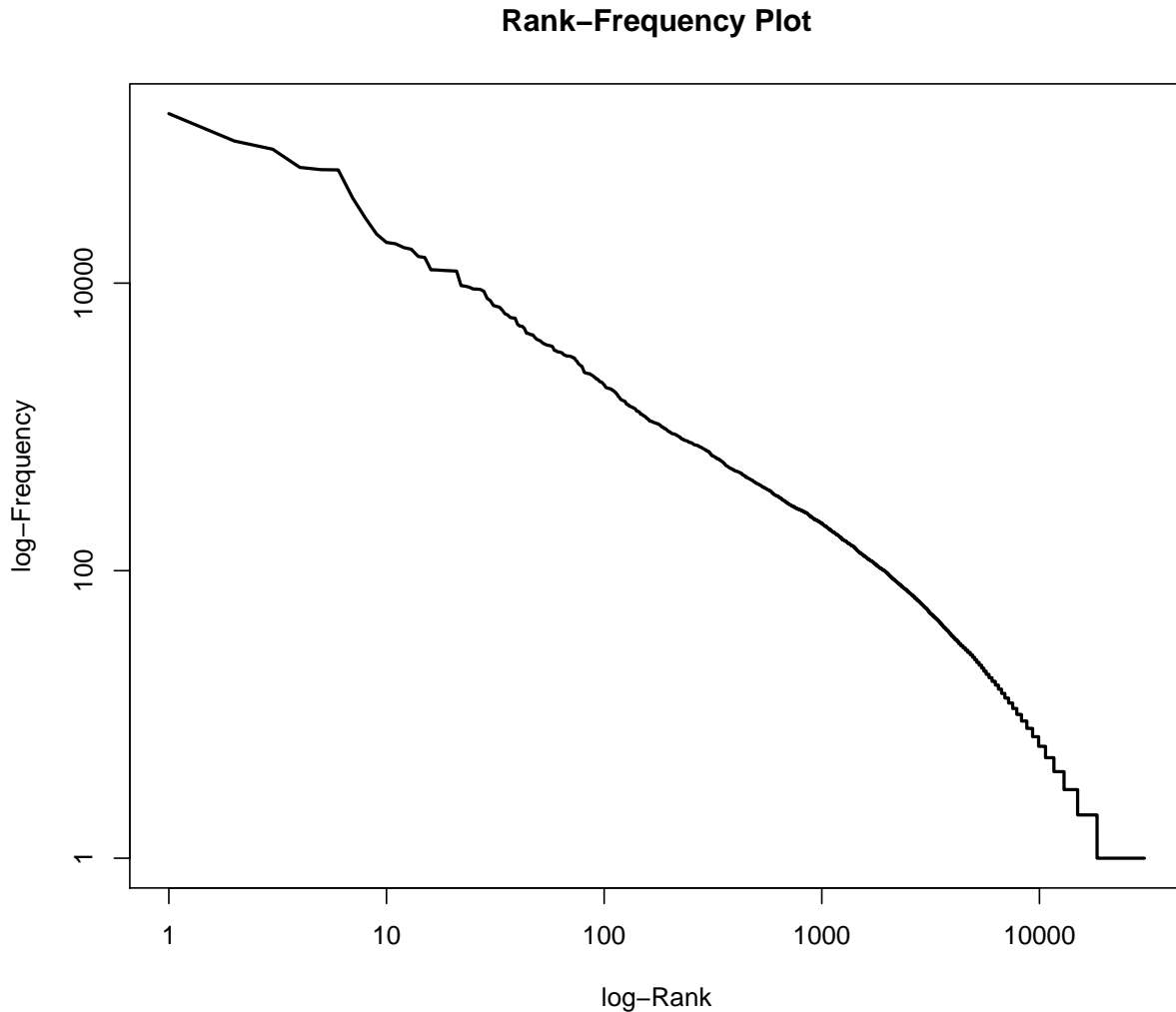
The words in this sorted list have a ranking depending on the position in this list. If the word ranks are plotted on the x axis and all frequencies on the y axis, then the Zipf distribution is obtained. This is a typical property of language data and its distribution is similar for all languages.

```
plot(wordlist$freqs , type = "l", lwd=2, main = "Rank frequency Plot", xlab="Rank", ylab = "Frequency")
```



The distribution follows an extreme power law distribution (very few words occur very often, very many words occur very rare). The Zipf law says that the frequency of a word is reciprocal to its rank ($1 / r$). To make the plot more readable, the axes can be logarithmized.

```
plot(wordlist$freqs , type = "l", log="xy", lwd=2, main = "Rank-Frequency Plot", xlab="log-Rank", ylab = "log-Frequency")
```

In the plot, two extreme ranges can be determined. Words in ranks between ca. 10,000 and 30358 can be observed only 10 times or less. Words below rank 100 can be observed more than 1000 times in the documents. The goal of text mining is to automatically find structures in documents. Both mentioned extreme ranges of the vocabulary often are not suitable for this. Words which occur rarely, on very few documents, and words which occur extremely often, in almost every document, do not contribute much to the meaning of a text.

Hence, ignoring very rare / frequent words has many advantages:

- reducing the dimensionality of the vocabulary (saves memory)
- processing speed up
- better identification of meaningful structures.

To illustrate the range of ranks best to be used for analysis, we augment information in the rank frequency plot. First, we mark so-called **stop words**. These are words of a language that normally do not contribute to semantic information about a text. In addition, all words in the word list are identified which occur less than 10 times.

The `%in%` operator can be used to compare which elements of the first vector are contained in the second vector. At this point, we compare the words in the word list with a loaded stopwords list (retrieved by the

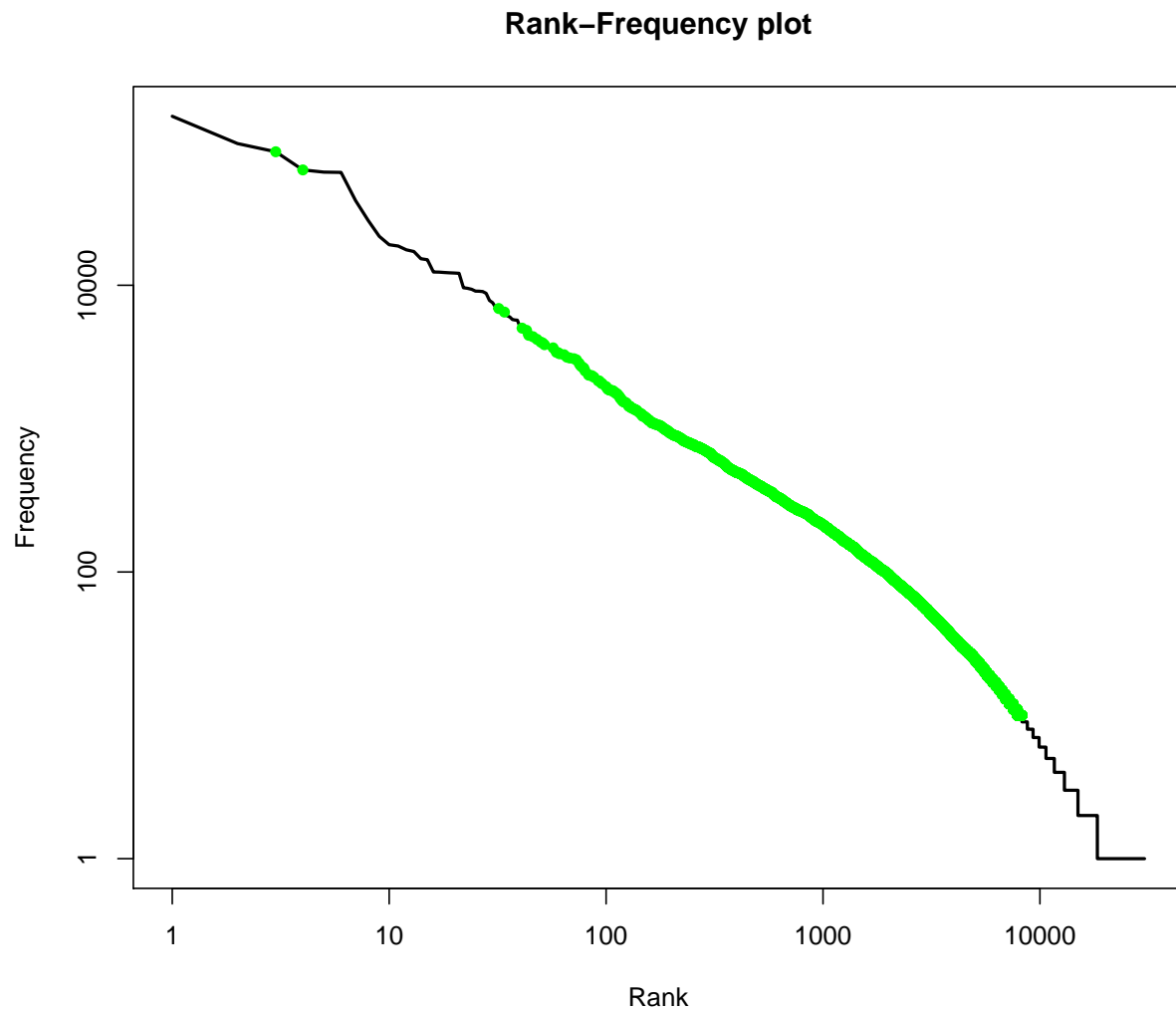
function `stopwords` of the `tm` package) . The result of the `%in%` operator is a boolean vector which contains TRUE or FALSE values.

A boolean value (or a vector of boolean values) can be inverted with the `!` operator (TRUE gets FALSE and vice versa). The `which` command returns the indices of entries in a boolean vector which contain the value TRUE.

We also compute indices of words, which occur less than 10 times. With a union set operation, we combine both index lists. With a `setdiff` operation, we reduce a vector of all indices (the sequence `1:nrow(wordlist)`) by removing the stopwords indices and the low frequent word indices.

With the command “`lines`” the range of the remaining indices can be drawn into the plot.

```
plot(wordlist$freqs, type = "l", log="xy", lwd=2, main = "Rank-Frequency plot", xlab="Rank", ylab = "Frequency")
englishStopwords <- stopwords("en")
stopwords_idx <- which(wordlist$words %in% englishStopwords)
low_frequent_idx <- which(wordlist$freqs < 10)
insignificant_idx <- union(stopwords_idx, low_frequent_idx)
meaningful_range_idx <- setdiff(1:nrow(wordlist), insignificant_idx)
lines(meaningful_range_idx, wordlist$freqs[meaningful_range_idx], col = "green", lwd=2, type="p", pch=20)
```



The green range marks the range of meaningful terms for the collection.

Optional exercises

1. Print out the word list without stop words and low frequent words.

```
##              words freqs
## ,              , 85384
## .              . 63784
## government government 6884
## states          states 6502
## congress        congress 5023
## united          united 4847
## ;              ; 4478
## can             can 4378
## -              - 4196
## people          people 4014
## upon           upon 3958
## year           year 3850
## $              $ 3659
## may            may 3408
## country        country 3390
## must           must 3329
## great          great 3275
## made           made 3151
## now            now 3110
## public         public 3074
## new            new 3020
## time           time 2865
## war            war 2767
## one            one 2713
## american       american 2668
```

2. If you look at the result, are there any corpus specific terms that should also be considered as stop word?
3. What is the share of terms regarding the entire vocabulary which occur only once in the corpus?

```
## [1] 0.39413
```

4. Compute the type-token ratio (TTR) of the corpus. the TTR is the fraction of the number of tokens divided by the number of types.

```
## [1] 0.01551821
```

References

1. Welbers, K., van Atteveldt, W., Benoit, K.: Text analysis in r. Communication Methods and Measures. 11, 245–265 (2017).