# Package 'tmT'

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#### 1 Introduction

(TODO) how to install and so on, main target group... textmeta objekte sind grundlage des pakets; was bei eigenen texten; was wenn restriktionen nicht erfüllt von textmeta objekt, standard-Datum erklären; readTextmeta erklären; hilfeseiten anpreisen...

```
install.packages("tmT")
library("tmT")
```

### 2 Data Preprocessing

A basic functionality of the package is data preprocessing. Therefore several functions are given for reading text data, creating text objects, manipulating these objects and especially handling duplicates of different forms in the text data.

#### 2.1 Read the Corpus - textmeta, readWikinews

Read the corpus data through one of your self implemented read-functions and create a textmeta object with the function of the same name and the arguments text, meta and metamult. The text component should be a list of character vectors or a list of lists of character vectors, whereas meta is a data.frame and metamult is intended for mainly unstructured meta-information as a list. Furthermore meta must contain columns id, date and title. You can test whether your object meets the requirements of a textmeta object with the function is.textmeta.

A read-function which is part of the package tmT is the function readWikinews. readWikinews reads XML-files created by the wikinews export page: https://en.wikinews.org/wiki/Special:Export. By default readWikinews reads all XML-files in the working directory. The function creates a textmeta object. For this Vignette we used two categories: Politics\_and\_conflicts and Economy\_and\_business. The pages were downloaded on 2018-03-05 in a file for each category. We can use readWikinews for reading both files, if they are in the same folder.

```
corpus <- readWikinews()
```

```
## Wikinews-20180305075910_politics.xml
## Wikinews-20180305080159_economy.xml
## number of observations in text: 7327
##
## NAs in text:
##
    NA.abs NA.rel
##
         0
##
##
## meta: 7327 observations of 3 variables
##
## NAs in meta:
##
         abs rel
## id
           0 0.00
## date 195 0.03
## title
## range of date: 2004-11-13 till 2018-03-04
## NAs in date: 195 (0.03)
```

Another method to read both files is to read both files seperately and merge them with the function mergeTextmeta. This function should be used if we want to merge data from different sources using different read-functions.

```
politics <- readWikinews(file="Wikinews-20180305075910_politics.xml")</pre>
## Wikinews-20180305075910_politics.xml
## number of observations in text: 5000
##
## NAs in text:
##
  NA.abs NA.rel
##
        0
##
## meta: 5000 observations of 3 variables
##
## NAs in meta:
##
        abs rel
          0 0.00
## id
         67 0.01
## date
## title
         0 0.00
## -----
## range of date: 2004-11-13 till 2009-11-19
## NAs in date: 67 (0.01)
economy <- readWikinews(file="Wikinews-20180305080159_economy.xml")</pre>
## Wikinews-20180305080159_economy.xml
## number of observations in text: 2327
##
## NAs in text:
##
  NA.abs NA.rel
##
        0
##
## meta: 2327 observations of 3 variables
##
## NAs in meta:
##
        abs rel
## id
          0 0.00
## date 128 0.06
## title
          0 0.00
## -----
##
## range of date: 2004-11-17 till 2018-03-04
## NAs in date: 128 (0.06)
corpus2 <- mergeTextmeta(list(politics, economy))</pre>
```

## NOTE: There are duplicates in the names of texts, could result in problems with unambiguity.

We get a note about duplicated texts (texts that appear in both categories). We have to handle this issue later. If we merge corpora with different meta-variables we can decide if all variables will be used for the merged corpora (all = TRUE, default) or only variables that appear in all corpora (all = FALSE).

After reading the raw data the texts need to be preprocessed.

#### 2.2 Remove Umlauts and XML/HTML Tags - removeXML removeHTML removeUmlauts

You can use removeXML to delete XML-tags (<...>) in character strings or a list of character vectors. The value you receive back will be a character vector or a list, if the input was a list.

If you retexts contain html entities use removeHTML. If you want to transform the entities in UTF-8 characters you can choose between the entity-type (dec=TRUE: ø, hex=TRUE: ø or entity=TRUE: ø). To choose which character should be replaced you can choose from all 16 ISO-8859 lists, e.g. symbolList=c(1,15) for ISO-8859-1 (latin1) and ISO-8859-15 (latin9). If delete=TRUE all remaining entities will be deleted. To replace german umlauts (ä ö ü ß -> ae oe ue ss) use removeUmlauts.

We remove XML-tags and HTML-entities from our Wikinews corpus. Since we have only punctuation as HTML-entities in the Corpus we remove it completely.

```
corpus$text <- removeXML(corpus$text)
corpus$text <- removeHTML(corpus$text, dec=FALSE, hex=FALSE, entity=FALSE)</pre>
```

#### ## delete remaining entities

It is possible to apply the function to the meta component of a textmeta object as well, for example to remove XML tags or umlauts from the title of the Wikipedia pages.

```
corpus$meta$title <- removeXML(corpus$meta$title)
corpus$meta$title <- removeHTML(corpus$meta$title, dec=FALSE, hex=FALSE, entity=FALSE)</pre>
```

#### ## delete remaining entities

After applying the function to the text component, we have removed all database relicts like XML-tags. At this point you should deal with identifying different types of duplicates in your text data.

#### 2.3 Identifying Duplicates - deleteAndRenameDuplicates, duplist

You should ensure unique IDs in all three components of your textmeta object. If you cannot ensure that, it is recommended to use the function deleteAndRenameDuplicates, which deletes "complete duplicates" - which means, that there are at least two entries with same ID and same information in text and in meta - and renames so called "real duplicates" - at least two entries with same ID and text, but different information in meta - and renames also "fake duplicates" - at least two entries with same ID but different text components.

TODO:[Genauer was umbenannt wird, tabelle mit beschreibung] It is important to know that for technical reasons - expecting duplicates in the names of the lists - this is the only function, which works with classic indexing, so that it assumes the same order of articles in all three components.

Additionally you can identify text component duplicates in your corpus with the function duplist, which creates a list of different types of duplicates. Non-unique IDs are not supported by the function, which implies that deleteAndRenameDuplicates should be executed before.

In the given example corpus complete duplicates are only expected if pages were associated to both categories. These duplicates are deleted.

```
corpus <- deleteAndRenameDuplicates(corpus)</pre>
```

#### ## success

The function deleteAndRenameDuplicates deleted 286 complete duplicates, so that duplist is applicable to the corpus.

```
dups <- duplist(corpus)

## ID-Fake-Dups... next step

## ID-Real-Dups... next step

## unique (and not-duplicated) texts... next step</pre>
```

```
## same texts... success
## duplist, list of (lists of) IDs with names:
## "uniqueTexts", "notDuplicatedTexts", "idFakeDups", "idRealDups",
## "allTextDups", "textOnlyDups", "textMetaDups", "textOthersDups".
## 5793 unique texts
## 5553 not-duplicated texts
## 0 texts with 0 different Fake-Dup IDs (ID equals, text differs)
## 0 texts with 0 different Real-Dup IDs (ID and text equals, meta differs)
## 1774 text-duplicates with 240 different texts
## 1708 textOnly-duplicates (text equals, meta differs) with 226 different texts
## 52 (text and meta)-duplicates with 21 different (text and meta) combinations, excluding ID
## 14 other text-duplicates
```

There is a possibility to visualize duplicates over time by the function plotScot which is explained in section 3.2.

For further analysis, especially performing the Latent Dirichlet Allocation, it is important that for each duplicate only one page is considered. Therefore it is the aim to reduce the corpus, so that it contains all pages which appear only once and a representative page for all pages which appear twice or more frequent. In our example we have only duplicated texts containing the empty string "" or small relicts like "\_\_NOTOC\_\_" or "\* \* \*"

#### 2.4 Clear Corpus - makeClear

For further preprocessing of text corpora tosca offers the function makeClear. It removes punctuation, numbers and stopwords. By default it removes english stopwords. It uses the stopword list of the function stopwords from the tm package. For the german stopword list are some additional words implemented (e.g. "dass" and "fuer"). You can control which stopwords should be removed with the argument sw. In addition the function changes all words to lowercase and tokenizes the documents. The result is a list of character vectors, or if paragraph is set TRUE (default) a list of lists of character vectors. The sublists should represent paragraphs of a document. If you hand over a textmeta object instead of a list of texts you will receive one back. In this case you have to hand it over to the parameter object instead of text.

The examples corpus language is english, so that sw should be set to stopwords() from the tm package, which includes english stopwords by default (kind = "en").

```
corpusClear <- makeClear(object = corpus)

## check articles on UTF8: next step

## change to lowercase: next step

## remove punctuation: next step

## remove numbers: next step

## remove stopwords: next step

## remove redundant whitespace: next step

## tokenize: next step

## remove empty articles: 1234 success</pre>
```

The function makeClear deletes all meta entries which did not belong to one of the texts (e.g. deleted empty texts). To create a textmeta object including this data the corresponding function is used.

```
textClear2 = makeClear(text = corpus$text)

## check articles on UTF8: next step

## change to lowercase: next step

## remove punctuation: next step

## remove numbers: next step
```

```
## remove stopwords: next step
## remove redundant whitespace: next step
## tokenize: next step
## remove empty articles: 1234 success
corpusClear2 = textmeta(text = textClear2, meta = corpus$meta)
```

#### 2.5 Generate Wordlist - makeWordlist

After clearing the corpus with the function makeClear we are able to call the function makeWordlist, which creates a table of occurring words in a given corpus. THe function table needs much RAM. That's a problem for bis Corpora. In makeWordlist we use the parameter k (default: 100000L) to reduce the number of texts which are processed at once. Large ks lead to faster calculation but more RAM usage.

For calculating wordlists a tokenized corpus must be used. In the given example corpusClear\$text is handed over to the function accordingly.

```
wordtable = makeWordlist(corpusClear$text)

## number of articles: 5807

## find out vocabularies...

## done:

## 0

## 5807 next step

## calculate counts...

## done:

## 0
```

## 3 Descriptive Analysis

After preprocessing the text data there is a typical workflow we recommend for looking at the corpus. This workflow contains the generic functions print and summary as well as the highly adaptable functions plotScot and plotFreq. These graphical functions should be part of any initial analysis of text data.

#### 3.1 Generic Functions - print, summary

Some information about the (one to) three components of the textmeta object is obtained by calling the generic function print.

```
print(corpus)

## Object of class "textmeta":

## number of observations in text: 7041

## meta: 7327 observations of 3 variables

## range of date: 2004-11-13 till 2018-03-04

## NAs in date: 195 (0.03)
```

The function provides the count of pages in the corpus (7041) and there are two additional columns in meta to the mandatory ones id, date and title. The pages are dated from 2004-11-13 till 2018-03-04.

You will get more information, especially about counts of NAs and tables of some candidates (default: resource and downloadDate) with the generic function summary. In addition to candidates you can handover the argument list.names (default: names(object)) for specifying the components out of text, meta and metamult which should be analysed by the function.

#### summary(corpus)

```
## number of observations in text: 7041
##
## NAs in text:
##
   NA.abs NA.rel
##
         0
##
## meta: 7327 observations of 3 variables
##
## NAs in meta:
##
         abs rel
           0 0.00
## id
## date 195 0.03
## title
           0 0.00
## ----
## range of date: 2004-11-13 till 2018-03-04
## NAs in date: 195 (0.03)
```

3.2 Visualisation of Corpus over Time - plotScot

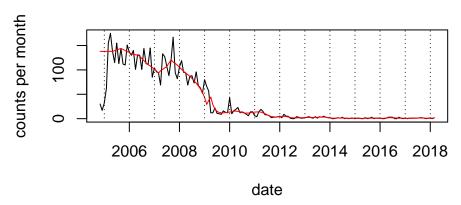
Apparently there are 191 NAs in the variable date.

One of the descriptive plotting functions in the package is plotScot (SubCorpusOverTime) which creates a plot of counts or proportion of either documents or words in a (sub)corpus over time. The subcorpus is specified by id and it is possible to set the unit to which the dates should be floored (default: "month"). The argument curves = c("exact", "smooth", "both") determine which curve(s) should be plotted. If you select type = "words", the object which you handover should be a tokenized textmeta object. If type = "docs" (default) you can hand over untokenized textmeta object as well.

First of all the number of texts per month in the complete example corpus is plotted, as exact and smoothed curve.

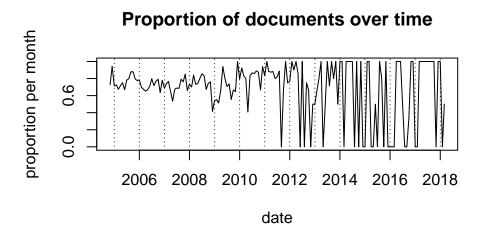
```
plotScot(corpusClear, curves = "both")
```

#### Count of documents over time



The black curve is the exact one and the red curve represents the smoothed values. The grafic gives a first impression about the distribution of the texts over time. Most of the news articles where written between 2005 and 2009. If we want to identify the distribution of duplicates over the time we can use plotScot to plot the IDs of the not duplicated texts in the corpus.

plotScot(corpus, id = dups\$notDuplicatedTexts, rel = TRUE)



Most zeros in the plot result from no articles in the whole corpus during these time periods. It is possible to get these values as NAs by setting natozero = FALSE in plotScot. This option works if rel = TRUE and is offered by many other functions in the package. Usually all plot functions in the package return the data belonging to the plot as invisible output. These plot functions offer a lot more functionality, which is described in the corresponding help functions.

#### 3.3 Frequency Analysis - plotFreq

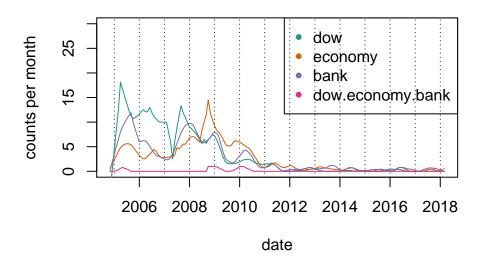
The other descriptive plotting function is plotFreq which performs a frequency analysis. Most of the arguments does not differ from plotScot. But the options wordlist and link = c("and", "or") are added for specifying the words of the frequency analysis and their link within one vector. In detail wordlist could either be a list of character vectors or a single character vector, which will be coerced to a list of

the vector's length. Each list entry represents a set of words which all (default link = "and") or one of them (link = "or") should appear in an article to be counted. The function uses filterWord with out = "count", which is explained later on, for counting.

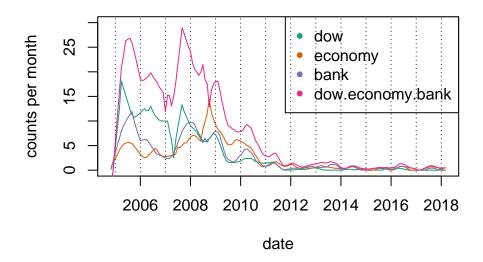
The example corpus contains Wikinews articles concerning the categories *Politics\_and\_conflicts* and *Economy\_and\_business*. Therefore some typical words out of these categories were taken to perform a frequency analysis. First of all the words *dow*, *economy* and *bank* were taken.

```
wordsEconomy <- list("dow", "economy", "bank", c("dow", "economy", "bank"))
plotFreq(corpusClear, wordlist = wordsEconomy, curves = "smooth",
   ylim = c(0, 30), legend = "topright")
plotFreq(corpusClear, wordlist = wordsEconomy, link = "or", curves = "smooth",
   ylim = c(0, 30), legend = "topright")</pre>
```

#### Count of wordlist-filtered documents over time - link:



#### Count of wordlist-filtered documents over time – link

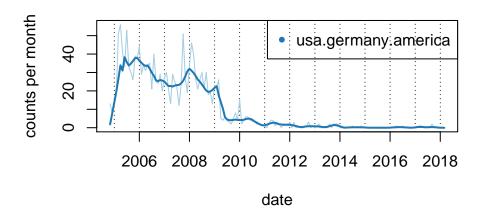


In the figures above you can see the difference between the and link and or link. In the first figure the three curves indicate the single words. The fourth curve shows the number of texts in which all three words appear. For most dates no texts meets this requirements. In the second figure the same single curves are shown. The fourth curve represents all three words again, but setting link = "or". The curve lies above the three others in every point. Due to smoothing it is possible that the line falls under one of the single word lines. This can be avoided by choosing curves = "exact".

In another figure the counts of pages in which the words usa, germany and america appear, are analysed. You can see that it is often useful to compare smoothed and exact curves to visualize the variance and a trend in the data.

```
plotFreq(corpusClear, wordlist = list(c("usa", "germany", "america")), link = "or",
    curves = "both", both.lwd = 2, legend = "topright")
```

#### Count of wordlist-filtered documents over time - link



#### 3.4 Write CSV Files - showTexts, showMeta

There are two functions for writing csv files implemented in the package. Both needs an textmeta object in showTexts, respectively the meta component of any-formated textmeta object in showMeta. The default of the parameter id in showTexts are all document IDs of the corpus as a character vector, but it is possible to handover a character matrix as well, so that each column will be represented in seperated csv file. In the first column of the csv file there will be the ID of each document, in the second and third the title and the date, whereas in the fourth column there will be the text itself.

Six IDs are sampled from the whole corpus with a given seed. These pages are saved as corpusllesen.csv and are returned as invisible to temp.

```
set.seed(123)
id <- sample(corpus$meta$id, 6)</pre>
temp <- showTexts(corpus, id = id)</pre>
temp[, c("id", "date", "title")]
##
           id
                    date
## 1
      ID46998 2006-08-11
## 2
      ID56896 2006-12-19
## 3 ID65115 2007-04-12
## 4 ID117206 2008-11-27
## 5 ID186836 2010-05-25
       ID7674 2005-04-09
## 6
##
                                                                               title
## 1
                                        Green Party refines 'Buy Kiwi Made' scheme
## 2
                                Pakistan: New policy on renewable energy launched
## 3
                                     Two Iraqi MPs killed in bombing at parliament
## 4
                                                 Wikinews Shorts: November 27, 2008
## 5
                                      Oil spilled after ships collide in Singapore
## 6 Bush EPA nominee abandons insecticide-on-children study after Senate hearing
```

We now have a look at the meta data. The default of the parameter id in showMeta are the IDs which are in the column meta\$id. You can also handover a matrix of IDs like in showTexts and you can specify which columns of the meta component you want to be written in the csv by setting the argument cols (default: colnames(meta)).

Analogously to showTexts the following code example will create three files named corpus<i>meta.csv, where i = 1, 2, 3 stands for the i-th column of the matrix of IDs.

```
temp <- showMeta(corpus$meta, id = matrix(id, nrow = 2),</pre>
  cols = c("title", "date"))
temp
## $`1`
##
                                                       title
                                                                    date
                Green Party refines 'Buy Kiwi Made' scheme 2006-08-11
## 5776 Pakistan : New policy on renewable energy launched 2006-12-19
##
## $`2`
##
                                                  title
                                                               date
## 2996 Two Iraqi MPs killed in bombing at parliament 2007-04-12
## 4751
                   Wikinews Shorts: November 27, 2008 2008-11-27
## 6468
                   Wikinews Shorts: November 27, 2008 2008-11-27
##
## $`3`
##
                                                                                   title
```

```
## 334 Bush EPA nominee abandons insecticide-on-children study after Senate hearing
## 6888 Oil spilled after ships collide in Singapore
## date
## 334 2005-04-09
## 6888 2010-05-25
```

### 4 Generating Subcorpora

The preprocessing which was done before is mandatory. For further preparation the package offers functions for filtering the corpus by dates, wordcount or search terms to generate subcorpora.

#### 4.1 Filter Corpus by Dates - filterDate

There are three implemented ways to filter your text corpora: One of these is the function filterDate, which filters a given textmeta object by a time period. The function works on any formated objects of class textmeta and extracts documents out of the text component, from which the date column in the meta component is in between s.date and e.date - including texts from both dates. The return value is the filtered textmeta object or a list which could be the text component of a textmeta object respectively, if you hand over the text and meta component not as a textmeta object.

The example corpus is filtered to articles dated between 2006 and 2009.

```
corpusDate <- filterDate(corpusClear, s.date = "2006-01-01", e.date = "2009-12-31")
print(corpusClear)

## Object of class "textmeta":
## number of observations in text: 5807
## meta: 6036 observations of 3 variables
## range of date: 2004-11-13 till 2018-03-04
## NAs in date: 42 (0.01)
print(corpusDate)

## Object of class "textmeta":
## number of observations in text: 3909
## meta: 3909 observations of 3 variables
## range of date: 2006-01-01 till 2009-12-31</pre>
```

The filtered corpus contains only the 3909 texts from the period 2006 till 2009.

#### 4.2 Filter Corpus by Wordcount - filterCount

#### TODO:(folgt...)

#### 4.3 Filter Corpus by Words - filterWord

The use of filterWord works analogously. It filters the text component of a textmeta object by appearances of specific words. The function uses regular expressions. It filters the given documents in the text component by words handed over by search, which could be a simple character vector or a list of data.frames. In the case of a character vector handed over the entries of the vector are linked by an *or*, so if *any* of the words appears in one specific document, it is returned.

Maybe you are not interested in the texts of the documents itself. Therefore you can set out to control the output: By default (out = text) you receive the filtered documents or if you hand over the argument object the corresponding textmeta respectively. If you choose out = bin you get the corresponding logical vector of indices and if you choose out = count you get a matrix - with the number of documents rows and the search-length respectively vector-length columns - which indicates in row i and column j how often the j-th word of the wordlist appears in the according i-th document.

First of all small examples are given for understanding the functionality of the function filterWord. An examples for the *or*-link is given by the next code example.

```
toyCorpus <- list(text1 = "dataset", text2 = "anything")
searchterm <- c("data", "as", "set", "anything")
filterWord(text = toyCorpus, search = searchterm, out = "bin")</pre>
```

```
## text1 text2
## TRUE TRUE
```

The returned values are TRUE twice. There is at least one pattern in the searchterm vector which appears at least once in each of the strings dataset and anything.

In the case of a list of data.frames handed over each data.frame is linked by an or and should contain columns pattern, word and count. The parameter pattern includes the search terms, the column word is a logical variable which controls whether words (TRUE) or patterns are searched. Alternatively word can be a character string containing the keyword left or right for left- or right-truncated search. You must set the argument count to an integer. This argument controls how often a word or pattern must appear in a document for it to be returned. Rows in each data.frame are linked by an and. An example is given by the following code.

```
searchframe <- data.frame(pattern = searchterm, word = FALSE, count = 1)
filterWord(text = toyCorpus, search = searchframe, out = "bin")</pre>
```

```
## text1 text2
## FALSE FALSE
```

In the case that words is handed over as data.frame, the *and* link is active. The function checks whether all of the patterns appear as part of words in the two entries of texts. Therefore the function returns FALSE twice.

For another sample case, we will delete the word anything from the search terms.

```
filterWord(text = toyCorpus, search = searchframe[1:3,], out = "bin")
## text1 text2
## TRUE FALSE
```

If you omit the word *anything* from words you receive a TRUE for text1 (*dataset*) - all three patterns appear in it - and a FALSE for text2 (*anything*), because not all patterns appear in it, not even one of them.

An example of out = count to receive a count for each text and search term combination is given by the following.

```
filterWord(text = list(text1 = c("i", "was", "here", "text"),
  text2 = c("some", "text", "about", "some", "text", "and", "something", "else")),
  search = c("some", "text"), out = "count")
```

```
## some text
## text1 0 1
## text2 3 2
```

In the case of out = count it is useful, that search is a simple character vector.

Another application of filterWord is to apply the function with word = TRUE, so that the function searchs only for single words, not for strings containing these words. This is displayed by the following example.

```
searchterm <- list(text1 = "land and and", text2 = c("and", "land", "and", "and"))
searchframe <- list(
  data.frame(pattern = "and", word = FALSE, count = 1),
  data.frame(pattern = "and", word = TRUE, count = 1))
filterWord(text = searchterm, search = searchframe, out = "count")

## and and_w
## text1 3 2
## text2 4 3</pre>
```

The function returns counts c(3, 4) for the simple pattern search and c(2, 3) for the word search, because the word and appears once in every document of searchterm only as pattern and not as single word.

After understanding the functionality of the function, finally it is used for filtering the Wikipedia corpus. The example corpus is filtered to those pages which fit to the chosen categories sofar, that the name of one of the catagories should appear on the page at least once, at least as pattern. It is not necessary to set ignore.case because the Wikipedia corpus was cleared before. This step includes that all words are lowercase now. TODO:[AND OR, OR Logik]

```
searchterm <- list(
  data.frame(pattern = "economy", word = FALSE, count = 1),
  data.frame(pattern = c("big", "data"), word = FALSE, count = 1),
  data.frame(pattern = "politics", word = FALSE, count = 1))
corpusFiltered = filterWord(corpusDate, search = searchterm)
print(corpusDate)

## Object of class "textmeta":
## number of observations in text: 3909
## meta: 3909 observations of 3 variables
## range of date: 2006-01-01 till 2009-12-31

print(corpusFiltered)

## Object of class "textmeta":
## number of observations in text: 451
## meta: 451 observations of 3 variables
## range of date: 2006-01-02 till 2009-12-17</pre>
```

The date and word filtered corpus consists of 451 texts compared to 3909 texts in the original corpusDate corpus.

#### 5 Latent Dirichlet Allocation

The main analytical functionality requested by text mining tools is to perform and analyse a Latent Dirichlet Allocation. In the package tmT this is ensured by the function LDAgen for performing the LDA, functions for validating the LDA results and various functions to visualize the results in different ways, especially over time. It is possible to analyse individual articles and its topic allocations as well. In addition a function for preparing your corpus for performing a Latent Dirichlet Allocation is given. This function creates a object which can be handed over to the function you could use for a LDA.

#### 5.1 Transform Corpus - LDAprep

The last step before performing a Latent Dirichlet Allocation is to create corpus data, which could be handed over to the function lda.collapsed.gibbs.sampler from the lda package or the function LDAgen from this package respectively. This is done by using the function LDAprep with its arguments text (text component of a textmeta object) and vocab (character vector of vocabularies). These vocabularies are the words which are taken into account for LDA.

You can have a look at the documentation of the lda.collapsed.gibbs.sampler for further information about lda. The function LDAprep offers options ldacorrect, excludeNA and reduce set all TRUE by default. The returned value is a list in which every entry symbolizes an article and contains a matrix with two rows. In the first row there is the index of each word in vocab minus one, in the second row there is the number of appearances of each word in the article. The option ldacorrect = TRUE ensures the second row is always one and the number of the appearances of the word will be shown by the number of columns belonging to this word. This structure is needed by lda.collapsed.gibbs.sampler.

Looking at the example corpus at first a new wordlist must be generated based on the filtered corpus.

```
wordtableFiltered <- makeWordlist(corpusFiltered$text)
## number of articles: 451
## find out vocabularies...
    done:
##
     0
##
     451 next step
##
  calculate counts...
##
    done:
##
##
     451 success
head(sort(wordtableFiltered$wordtable, decreasing = TRUE))
##
         said
                    will
                              people
                                                         us government
                                          party
##
         1230
                    1203
                                 852
                                            832
words5 <- wordtableFiltered$words[wordtableFiltered$wordtable > 5]
pagesLDA = LDAprep(text = corpusFiltered$text, vocab = words5)
```

After receiving the words which appear at least six times in the whole filtered corpus, the function LDAprep is applied to the example corpus with vocab = words5. The object pagesLDA will be handed over to the function which performs a Latent Dirichlet Allocation.

#### 5.2 Performing LDA - LDAgen

The function which has to be applied first to the corpus manipulated by LDAprep is LDAgen. Therefore the function offers the options K (integer, default: K = 100L) to set the number of topics, vocab (character vector) for specifying the words which are considered in the manipulation of the corpus and several more e.g. number of iterations for the burnin (default: burnin = 70) and the number of iterations for the gibbs sampler (default: num.iterations = 200). The result will be saved in a R workspace, the first part of the results name can be specified by setting the option folder (default: folder = "lda-result").

In the concrete example corpus the manipulated corpus pagesLDA is used for documents, the topic number is set to K = 10 and for reproducibility a seed is set to seed = 123. The filename consists of the folder argument followed by the options of K, num.iterations, burnin and the seed of the LDA.

```
LDAgen(documents = pagesLDA, K = 10L, vocab = words5, seed = 123)
load("lda-result-k10i200b70s123alpha0.1eta0.1.RData")
```

For validation of the LDA result and further analysis, the result is loaded back to workspace.

#### 5.3 Validation of LDA Results - intruderWords, intruderTopics

For validation of LDA results there are two functions in the package. These functions expect user input, the user works like a text labeller. The LDA result is handed over by setting beta = result\$topics. During the function intruderWords the labeler gets a set of words. The number of words can be set by numOutwords (default: 5). This set represents one topic. It includes a number of intruders (default: numIntruder = 1), which can also be zero. In general, if the user identifies the intruder(s) correctly this is an identifier for a good topic allocation. You can set options numTopwords (default: 30) to control which top words of each topic are considered for this validation. In addition it is possible to enable or disable the possibility for the user to mark nonsense topics. By default this option is enabled (noTopic = TRUE). The true intruder can be printed to the console after each step with printSolution = TRUE (default: FALSE).

#### TODO:[non-preprocessed corpus nach removexml, .

The LDA result of the example corpus is checked by intruderWords with a number of intruders of zero or one.

```
set.seed(101)
intWords <- intruderWords(beta = result$topics, numIntruder = 0:1)

## counter 1
## 1 countries
## 2 will
## 3 billion
## 4 budget
## 5 yeah</pre>
```

As an illustration the first set is shown. Obviously the word newspaper does not fit into the set with the words computational, cell, cells and engineering. Therefore the user would type 5 and press enter. If the user wants to mark nonsense topics he would type an x (in the summary the number of meaningful topics is shown) and 0 if he thinks there is no intruder word. Actually newspaper is the true intruder in the set above. As an example user input c(5, 0, 0, 5, 1, 2, 0, 5, 3, 5) is considered.

```
print(intWords)
```

```
## Parameters:
##
    byScore numTopwords numIntruder numOutwords noTopic
##
                       30
                                   0 1
                                                         TRUE
##
## Results:
          numIntruder missIntruder falseIntruder
##
##
    [1,]
                     0
                                   0
                                                   0
##
    [2,]
                     0
                                   0
                                                   1
##
    [3,]
                     0
                                   0
                                                   0
##
   [4,]
                     1
                                   0
                                                   0
##
   [5,]
                     0
                                   0
                                                   1
##
    [6,]
                     1
                                   1
                                                   0
##
   [7,]
                     1
                                   0
                                                   0
##
   [8,]
                     1
                                   0
                                                   0
## [9,]
                                   0
                                                   0
                     1
## [10,]
                                   0
```

By printing the object of <code>intruderWords</code> to the console, you get information about options for the validation strategy and a results matrix with ten rows an three columns. The rows indicate the different sets of potential intruders. For each set the matrix contains information how many intruder are in the specific set, how many intruders were missed by the user and how many false intruders were named. Of course the columns <code>missIntr</code> und <code>falseIntr</code> matchs if <code>numIntruder</code> is a scalar and the user names exactly this number of potential intruders for each set.

#### summary(intWords)

```
## 0 not interpretable topics
##
    0 not evaluated topics
##
## Parameters:
##
    byScore numTopwords numIntruder numOutwords noTopic
##
                     30
                                 0 1
                                                     TRUE
## 10 out of 10 meaningful topics (100 %)
   7 correct Topics (70 %)
##
## Table of Intruders:
## 0 1
## 4 6
##
## Mean number of missed Intruders: 0.1
## Table of missed Intruders:
## 0 1
## 9 1
## Mean number of false Intruders: 0.2
## Table of false Intruders:
## 0 1
## 8 2
```

Applying summary to an object of type intruderWords will result in an ouput of some measures concerning the validation. Each function call contains ten sets. You are able to continue labelling by calling intruderWords with oldResult = intWords if your set was not finished.

```
intWords <- intruderWords(oldResult = intWords)</pre>
```

Analogously to intruderWords you can use intruderTopics for validation the other way around. This function is used for validation of topics associated to a specific document instead of validation of words associated to one topic. Therefore the document is displayed in another window and a sample of topics represented by the ten top.topic.words - is shown in the console. You should hand over in text the text component of the original untokenized corpus before manipulation by makeClear, so that the document is kind of readable. The user then names the intruder(s). There are options for different numbers of topics and intruders as in intruderWords as well. The parameter theta should be set to result\$document\_expects given result is the LDA result. An example call is given below.

```
intruderTopics(text = corpus$text, id = ldaID,
  beta = result$topics, theta = result$document_expects)
```

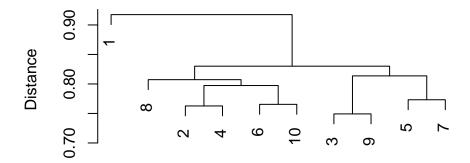
#### 5.4 Clustering of Topics - clusterTopics, mergeLDA

For analysing topic similarities it is useful to cluster the topics. The function clusterTopics implements this. The main argument is topics and should be set to the topics element of the result object. You could specify file, width and height (both integers) to write the resulting plot to a pdf. Other options

are topicnames for labelling the topics in the plot and method (default: "average"), which influences the way the topics are clustered. The method statement is used for applying the distance matrix to the function hclust. The distance matrix is computed based on the hellinger distance and is returned in a list together with the value of the hclust call as invisible by clusterTopics.

```
clustRes <- clusterTopics(ldaresult = result, xlab = "Topic", ylab = "Distance")</pre>
```

### **Cluster Dendrogram**



Topic hclust (\*, "average")

names(clustRes)

## [1] "dist" "cluster"

The same plot as above can be recreated by calling plot(clustRes\$cluster). In the plot you can see the similarities concerning the hellinger distance of the topics.

It is possible to merge different results of LDAs by calling mergeLDA(list(result1, result2, ..., resultN)). The function mergeLDA binds the topics elements of the results by row and only consider words which appears in all results. As result you receive the topics matrix including all topics from the given results.

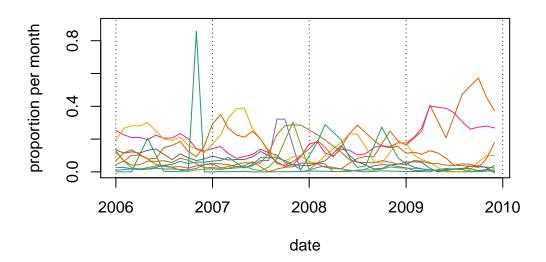
#### 5.5 Visualisation of Topics over Time - plotTopic

As extension of the highly flexible functions plotScot and plotFreq the package tmT offers another plotting function of the same type. The function plotTopic does something very similar to these two functions. It pictures the counts or proportion of words allocated to different topics of a LDA result over time. The result object is handed over in ldaresult, the belonging IDs of documents as a character vector in ldaid. In object the function expect a strictly tokenized textmeta object. You could set select for selecting topics by an integer vector. By default all topics are selected. Analoguesly to wnames in plotFreq it is possible to set topic names with tnames. By default the index and the most representative word (top.topic.words) per topic are chosen as names. For further individualisation the function offers mostly the same options as plotScot and plotFreq.

Often it is useful to choose curves = "smooth" if you do not select topics, because there is a massive fluctuation of exact curves. However, it is important to have a look at the exact curves, because the smoothed curves are someway manipulated by the statement smooth, so the user is tempted to optimise the smoothing parameter for getting the curves he or she wants.

```
plotTopic(object = corpusFiltered, ldaresult = result, ldaID = ldaID,
    rel = TRUE, curves = "smooth", smooth = 0.1, legend = "none", ylim = c(0, 0.9))
```

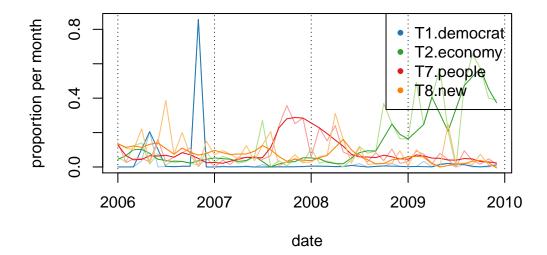
## Proportion of topics over time



There is no difference of handing over an inflated corpus with documents which were not used for LDA. But the corpus must contain all documents of the LDA.

```
plotTopic(object = corpusClear, ldaresult = result, ldaID = ldaID,
    select = c(1:2, 7:8), rel = TRUE, curves = "both", smooth = 0.1)
```

## Proportion of topics over time

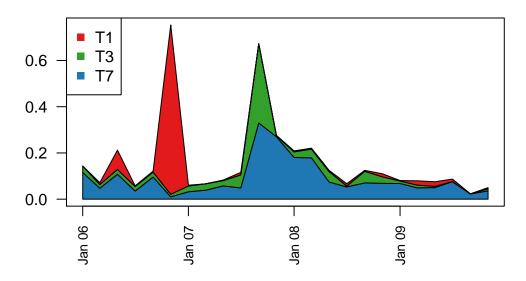


#### 5.6 Visualisation of Topic Share over Time - plotArea

The function plotArea offers possibilities to create so called area visualisations of topics over time. It requires arguments ldaresult, ldaid and meta as introduced before. There are options select, tnames, unit and others. Additionally you can set threshold to a numeric between 0 and 1, as a limit, which a topics proportion have to surpass at least once to be plotted.

Because this seems to be interesting topics *T1.syndrome* (red curve), *T3.health* (green) and *T7.medicine* (blue) are plotted in a sediment plot. The chosen unit is "bimonth" (default is "quarter").

```
plotArea(ldaresult = result, ldaID = ldaID, meta = corpusFiltered$meta,
    select = c(7, 3, 1), unit = "bimonth", sort = FALSE)
```



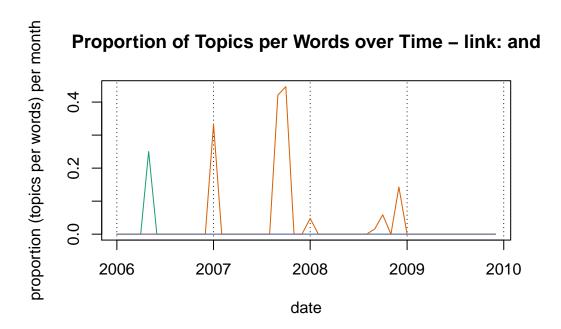
(TODO) Interpretation.

#### 5.7 Visualisation of Words in Topic over Time - plotTopicWord, plotWordpt

Another visualise topics over time is given by plotTopicword. It displays the counts or proportion of given topic-word combinations. If rel = TRUE the baseline for normalisation are the words counts, not the counts of topics. Arguments which have to specified are object (corpus, textmeta object), docs (corpus manipulated by LDAprep, the input for LDAgen) and the ldaresult with its ldaid (IDs of documents in docs or ldaresult respectively). The function asks for docs for complexity reasons. This object should be created while preparation for LDA anyway. The options wordlist and select are known from other plot functions and offer a lot of different topic word combinations which should be plotted by plotTopicword.

In the example corpus the proportion of the word *economy* in the topics one, three and seven is explored. The top.topic.words of the three chosen topics are *syndrome* (lightgreen curve), *health* (orange) and *medicine* (purple)

```
plotTopicWord(object = corpusFiltered, docs = pagesLDA, ldaresult = result, ldaID = ldaID,
   wordlist = "economy", select = c(1, 3, 7), rel = TRUE, legend = "none")
```



The graphic shows that the word *economy* is associated to the topic *health* most often.

For interpretating it is important to keep in mind the baseline, the word counts of *economy*. To display this the sums of all topic-word proportions are calculated and are expected to be one for all dates which appear at least once, otherwise zero.

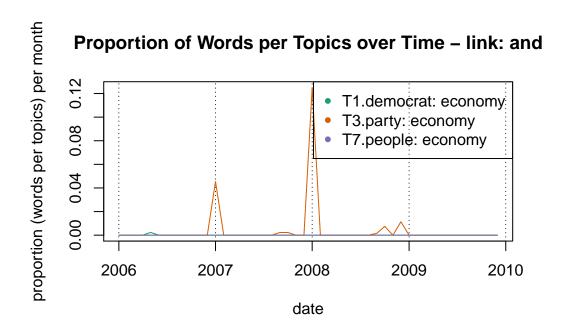
```
tab <- plotTopicWord(corpusFiltered, pagesLDA, result, ldaID, "economy", rel = TRUE)
all(round(rowSums(tab[, -1]), 10) %in% c(1, 0))</pre>
```

#### ## [1] TRUE

This is confirmed by the call above. For some analysis maybe it could be interesting to take the other possible baseline, the topic counts, into account. Therefore there is an additional function called plotWordpt.

The function plotWordpt works analogously like its pendant plotTopicWord, but with baseline topic sums instead of word sums. The difference between both functions plotWordpt and plotTopicWord is given by the fact that plotWordpt considers topic peaks. You will get the relative counts of the selected word(s) in the selected topic(s). All curves sum up to one if you choose any topic and the whole vocabulary list as wordlist.

```
plotWordpt(object = corpusFiltered, docs = pagesLDA, ldaresult = result, ldaID = ldaID,
   wordlist = "economy", select = c(1, 3, 7), rel = TRUE)
```

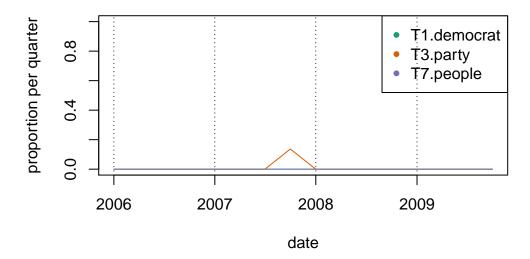


#### 5.8 Visualisation of Words in Articles allocated to Topics - plotWordSub

Imagine you want to identify words which are used frequently in articles allocated to a topic. The function which realizes a plot to the question is called plotWordSub. The first problem is allocation of topics. Therefore you set a absolute or relative limit how often words of a given article are allocated to one topic. Additionally you have to specify whether one article is allocated exactly once, maximum once or multiple times depending on the limit argument. The default is limit = 10 and alloc = "multi", so an article is allocated to a topic if it contains at least 11 words which are allocated to the given topic. Multiple or no allocations are possible. After allocating the articles to the topics the function creates subcorpora using filterWord. To control the filter you have to set the search argument. The counts of the subcorpora (normalized to their whole corpora) are plotted. There are many options to personalize your plot like in the other plot functions.

```
searcheco <- data.frame(pattern = "economy", word = TRUE, count = 3)
plotWordSub(object = corpusFiltered, ldaresult = result, ldaID = ldaID, limit = 1/3,
    select = c(1, 3, 7), search = searcheco, unit = "quarter", legend = "topright")</pre>
```

### **Proportion of Documents in given Subcorpus over Time**



The plot shows subcorpora generated by the **search** argument above, which means articles must contain the word *economy* at least three times. The corpora from which these subcorpora are generated have to contain one third of words which are allocated to the corresponding topic (limit = 1/3).

#### 5.9 Heatmap of Topics over Time including Clustering - plotHeat

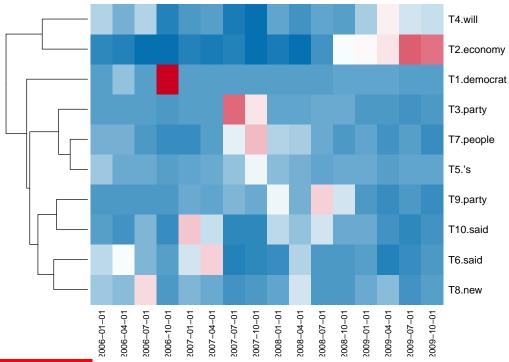
The use case for plotHeat is given by searching for explicit peaks of coverage of some topics. Therefore the resulting heatmap shows the deviation of the proportion of a given topic at this current time from its mean proportion. In addition a dendrogramm is plotted on the left side of the heatmap showing similarities of topics. The clustering is performed with hclust on the dissimilarities computed by dist.

By default the proportions are calculated on the article lengths, but it is possible to force calculation on only the LDA vocabulary by setting object to a textmeta object only including meta information. Otherwise a strictly tokenized textmeta object is required. The parameters ldaresult and ldaID expect a LDA result and according IDs like in functions mentioned before. Options tnames (topic label), file (if you want to save the plot in a pdf) and unit (default: round dates to "year") are given as well. Additionally it is possible to set whether the deviations should be normalised to take different topic sizes into account (default: norm = FALSE). You can change the intervals of labeling on the x-axis by setting date\_breaks. By default (date\_breaks = 1) every label is drawn. If you choose date\_breaks = 5 every fifth label will be drawn.

The peak of the topic T1.syndrome in January 2014 was mentioned several times before. This should be visible in the following heatmap as well. As compromise between clarity and interpretability unit = "quarter" is chosen.

```
plotHeat(object = corpusFiltered, ldaresult = result, ldaID = ldaID, unit = "quarter")
```

#### **Absolute Deviation of Topic Shares from Mean Topic Share**



As expected the *T1.syndrome* topics peak is clearly identifiable. The according rectangle at the first quarter of 2014 is colored by the deepest red of this figure. On the other hand mostly all other quarters of years concerning this topic are colored lightblue. Other remarkable quarters are for example the fourth quarter of 2015 or 2016, where the topic *T2.data* or *T4.cells* respectively has noticeable peaks. The dendrogramm shows that none of the topics are similar to another concerning the absolute deviations of topic proportion from the mean topic proportion per quarter. This supports the findings of clustering the topics with clusterTopics.

#### 5.10 Individual Cases Contemplation - topTexts, topicsInText

Sometimes it is useful to look at some individual cases sometimes. Especially the documents with the highest counts or proportion of words belonging to one topic are of interest. These documents can be extracted by topTexts. By default (rel = TRUE) the proportion is considered. The function requires a ldaresult and the according ldaid. It offers options select, limit and minlength, which control how much articles per topic (default: all topics) are returned (default: limit = 20) and articles of which minimum length (default: minlength = 30) are taken into account. The output value is a matrix of the according IDs.

```
In the example the top four pages from the topics T1.syndrome, T3.health and T7.medicine are requested.

topID <- topTexts(ldaresult = result, ldaID = ldaID, select = c(1, 3, 7), limit = 4)

dim(topID)
```

#### ## [1] 4 3

Obviously the corresponding matrix has four rows and three columns.

After identifying the top pages it is possible to have a deeper look at these articles. Therefore the mentioned function showTexts can be used. The returned value is a list with three entries with data.frames of four rows - the different pages - and four columns each - id, title, date and text. For displaying, the fourth column of each data.frame containing the pages content itself is removed.

```
topArt <- showTexts(corpusFiltered, id = topID)
lapply(topArt, function(x) x[, 1:3])</pre>
```

```
## $T1.democrat
##
           id
## 1
     ID54162
## 2
     ID40076
## 3 ID116056
## 4 ID38241
##
                                                                  title
## 1
                                     2006 U.S. Congressional Elections
## 2 PAP returned to power in 2006 Parliamentary Elections, Singapore
                 Third party U.S. presidential candidates hold debate
## 4
                        US Federal Reserve ceases to publish M3 index
##
           date
## 1 2006-11-08
## 2 2006-05-07
## 3 2008-11-02
## 4 2006-04-27
##
  $T3.party
##
          id
## 1 ID80831
## 2 ID80085
## 3 ID80811
## 4 ID81466
##
## 1
                         Ontario Votes 2007: Interview with Libertarian candidate Mark Scott, Toronto-D
                              Ontario Votes 2007: Interview with Green candidate Peter Ormond, Hamilton
## 3 Ontario Votes 2007: Interview with Family Coalition Party candidate Bill McMaster, Lambton-Kent-Mi
                     Ontario Votes 2007: Interview with Progressive Conservative candidate Dan McCreary
## 4
##
           date
## 1 2007-09-27
## 2 2007-09-18
## 3 2007-09-26
## 4 2007-10-02
##
## $T7.people
##
## 1 ID82665
## 2 ID83419
## 3 ID80882
## 4 ID82308
##
                                                                                  title
## 1
                 John Reed on Orwell, God, self-destruction and the future of writing
## 2
                             Gay Talese on the state of journalism, Iraq and his life
                              John Vanderslice plays New York City: Wikinews interview
## 4 Sam Brownback on running for President, gay rights, the Middle East and religion
##
           date
## 1 2007-10-18
## 2 2007-10-27
## 3 2007-09-27
## 4 2007-10-15
```

At last the function topicsInText offers the possibilty to analyse a single document's topic allocations. The function creates a HTML document with its words colored depending on the topic allocations of each word. It requires arguments ldaresult and ldaID as usual. The belonging LDAprep object should be handed over

in text, while the vocabulary set as character vector in words. You will set id to the documents ID you are interested in. It is possible to show the origing text by setting originalText to the belonging uncleared text component of your textmeta object. There are some more options - e.g. wordOrder - for modifying the output individually.

The article Family medicine with ID 1656748 from topic T3.health and category Medicine is analysed with the function topicsInText in more detail.

```
topicsInText(text = pagesLDA, ldaresult = result, ldaID = ldaID,
  id = topArt$T3.economy[4,1], vocab = words5, originaltext = corpus$text, wordOrder = "")
```

In the part of the HTML output above at first the different topics in the order of its absolute appearences in the given document are displayed. The topics are represented by its 20 top.topic.words each and are colored each in its own color. Words which were deleted by clearing the corpus are colored black. This way you are able to check plausibility of individual documents, so topicsInText can also be seen as individual cases validation.

#### 6 Conclusion

(TODO) Wichtigste Punkte des Workflows zusammenfassen, allgemeine Fallstricke (Duplikate . . . ), Diskussion des Anwendungsbeispiels (viele stopwords nicht geloescht . . . ), Ausblick? (weitere Funktionen . . . )