

Package ‘qdap’

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Type Package

Title Bridging the gap between qualitative data and quantitative analysis

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Depends R (>= 2.15), ggplot2 (>= 0.9.2)

Imports gridExtra, chron, scales, RColorBrewer, igraph, tm, wordcloud, venneuler, openNLPmodels.en, Snowball, gplots, gridExtra, gdata, openNLP, XML, RCurl, parallel

Suggests reshape2, plyr, koRpus

LazyData TRUE

ByteCompile yes

Description This package automates many of the tasks associated with quantitative discourse analysis of transcripts containing discourse including frequency counts of sentence types, words, sentence, turns of talk, syllable counts and other assorted analysis tasks. The package provides parsing tools for preparing transcript data. Many functions enable the user to aggregate data by any number of grouping variables providing analysis and seamless integration with other R packages that undertake higher level analysis and visualization of text. This provides the user with a more efficient and targeted analysis.

License GPL-2

URL <https://github.com/trinker/qdap/wiki>

Collate

'adjacency_matrix.R' 'all_words.R' 'automated_readability_index.R' 'bag.o.words.R' 'blank2NA.R' 'bracketX.R' 'capitalize.R' 'qprep.R' 'rank_freq_plot.R' 'read.docx.R' 'read.transcript.R' 'reducer.R' 'reduceWORDS.R' 'replace_abbrev.R'

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abbreviations	<i>Small Abbreviations Data Set</i>
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Description

A dataset containing abbreviations and their qdap friendly form.

Format

A data frame with 14 rows and 2 variables

Details

- abv. Common transcript abbreviations
- rep. qdap representation of those abbreviations

action.verbs

*Action Word List***Description**

A dataset containing a vector of action words. This is a subset of the **Moby project: Moby Part-of-Speech**.

Format

A vector with 1569 elements

Details

From Grady Ward's Moby project: "This second edition is a particularly thorough revision of the original Moby Part-of-Speech. Beyond the fifteen thousand new entries, many thousand more entries have been scrutinized for correctness and modernity. This is unquestionably the largest P-O-S list in the world. Note that the many included phrases means that parsing algorithms can now tokenize in units larger than a single word, increasing both speed and accuracy."

References

<http://icon.shef.ac.uk/Moby/mpos.html>

adjacency_matrix

*Takes a Matrix and Generates an Adjacency Matrix***Description**

Takes a matrix (wfm) or termco object (.a, .c or .d) and generates an adjacency matrix for use with igraph

Usage

```
adjacency_matrix(matrix.obj)
```

```
adjmat(matrix.obj)
```

Arguments

matrix.obj A matrix object, preferably, of the class "termco_d" or "termco_c" generated from terco.a, termco.d or termco.c.

Value

Generates an adjacency matrix

See Also[dist](#)**Examples**

```
## Not run:
wordLIST <- c(" montague", " capulet", " court", " marry")
(raj.termco <- with(raj.act.1, termco.a(dialogue, person,
  wordLIST, ignore.case = T)))
(raj.adjmat <- adjmat(raj.termco))
names(raj.adjmat) #see what's available from the adjacency_matrix object
library(igraph)
g <- graph.adjacency(raj.adjmat$adjacency, weighted=TRUE, mode ='undirected')
g <- simplify(g)
V(g)$label <- V(g)$name
V(g)$degree <- degree(g)
layout1 <- layout.auto(g)
plot(g, layout=layout1)

## End(Not run)
```

adverb

*Adverb Word List***Description**

A dataset containing a vector of adverbs words. This is a subset of the **Moby project: Moby Part-of-Speech**.

Format

A vector with 13398 elements

Details

From Grady Ward's Moby project: "This second edition is a particularly thorough revision of the original Moby Part-of-Speech. Beyond the fifteen thousand new entries, many thousand more entries have been scrutinized for correctness and modernity. This is unquestionably the largest P-O-S list in the world. Note that the many included phrases means that parsing algorithms can now tokenize in units larger than a single word, increasing both speed and accuracy."

References

<http://icon.shef.ac.uk/Moby/mpos.html>

`all_words`*Searches Text Column for Words*

Description

A convenience function to find words that begin with or contain a letter chunk and returns the frequency counts of the number of occurrences of each word.

Usage

```
all_words(text.var, begins.with = NULL, contains = NULL,  
          alphabetical = TRUE)
```

Arguments

<code>text.var</code>	The text variable
<code>begins.with</code>	This argument takes a word chunk. Default is NULL. Use this if searching for a word beginning with the word chunk.
<code>contains</code>	This argument takes a word chunk. Default is NULL. Use this if searching for a word containing the word chunk.
<code>alphabetical</code>	logical. If True orders rows alphabetically, if false orders the rows by frequency.

Value

Returns a dataframe with frequency counts of words that begin with or contain the provided word chunk.

Note

Can not provide both `begins.with` and `contains` arguments at once. If both `begins.with` and `contains` are NULL `all.words` returns a frequency count for all words.

See Also

[term.match](#)

Examples

```
## Not run:  
all_words(raj$dialogue, begins.with="re")  
all_words(raj$dialogue, "q")  
all_words(raj$dialogue, contains="conc")  
all_words(raj$dialogue)  
  
## End(Not run)
```

automated_readability_index

Readability Measures

Description

automated_readability_index - Apply Automated Readability Index to transcript(s) by zero or more grouping variable(s).

coleman_liau - Apply Coleman Liau Index to transcript(s) by zero or more grouping variable(s).

SMOG - Apply SMOG Readability to transcript(s) by zero or more grouping variable(s).

flesch_kincaid - Flesch-Kincaid Readability to transcript(s) by zero or more grouping variable(s).

fry - Apply Fry Readability to transcript(s) by zero or more grouping variable(s).

linsear_write - Apply Linsear Write Readability to transcript(s) by zero or more grouping variable(s).

Usage

```
automated_readability_index(text.var,
  grouping.var = NULL, rm.incomplete = FALSE, ...)

coleman_liau(text.var, grouping.var = NULL,
  rm.incomplete = FALSE, ...)

SMOG(text.var, grouping.var = NULL, output = "valid",
  rm.incomplete = FALSE, ...)

flesch_kincaid(text.var, grouping.var = NULL,
  rm.incomplete = FALSE, ...)

fry(text.var, grouping.var = NULL, labels = "automatic",
  rm.incomplete = FALSE, ...)

linsear_write(text.var, grouping.var = NULL,
  rm.incomplete = FALSE, ...)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
rm.incomplete	logical. If TRUE removes incomplete sentences from the analysis.
...	Other arguments passed to endf.
output	A character vector character string indicating output type. One of "valid" (default and congruent with McLaughlin's intent) or "all".
labels	A character vector character string indicating output type. One of "automatic" (default; adds labels automatically) or "click" (interactive).

Value

Returns a dataframe with selected readability statistic by grouping variable(s). The `fry` function returns a graphic representation of the readability.

Note

Many of the indices (e.g. Automated Readability Index) are derived from word difficulty (letters per word) and sentence difficulty (words per sentence). If you have not run the `sentSplit` function on your data the results may not be accurate.

References

- Coleman, M., & Liau, T. L. (1975). A computer readability formula designed for machine scoring. *Journal of Applied Psychology*, Vol. 60, pp. 283-284.
- Flesch R. (1948). A new readability yardstick. *Journal of Applied Psychology*. Vol. 32(3), pp. 221-233. doi: 10.1037/h0057532.
- Gunning, T. G. (2003). *Building Literacy in the Content Areas*. Boston: Allyn & Bacon.
- McLaughlin, G. H. (1969). SMOG Grading: A New Readability Formula. *Journal of Reading*, Vol. 12(8), pp. 639-646.
- Senter, R. J., & Smith, E. A.. (1967) Automated readability index. Technical Report AMRLTR-66-220, University of Cincinnati, Cincinnati, Ohio.

Examples

```
## Not run:
with(rajSPLIT, automated_readability_index(dialogue, list(person, act)))
with(rajSPLIT, automated_readability_index(dialogue, list(sex, fam.aff)))

with(rajSPLIT, coleman_liau(dialogue, list(person, act)))
with(rajSPLIT, coleman_liau(dialogue, list(sex, fam.aff)))

with(rajSPLIT, SMOG(dialogue, list(person, act)))
with(rajSPLIT, SMOG(dialogue, list(sex, fam.aff)))

with(rajSPLIT, flesch_kincaid(dialogue, list(person, act)))
with(rajSPLIT, flesch_kincaid(dialogue, list(sex, fam.aff)))

(x <- with(rajSPLIT, fry(dialogue, list(sex, fam.aff))))
with(rajSPLIT, fry(dialogue, list(sex, fam.aff), labels = "click"))

with(rajSPLIT, linsear_write(dialogue, list(person, act)))
with(rajSPLIT, linsear_write(dialogue, list(sex, fam.aff)))

## End(Not run)
```

Description

`bag.o.words` - Reduces a text column to a bag of words.

`breaker` - Reduces a text column to a bag of words and qdap recognized end marks.

`word.split` - Reduces a text column to a list of vectors of bag of words and qda recognized end-marks (i.e. ".", "!", "?", "*", "-").

Usage

```
bag.o.words(text.var, apostrophe.remove = FALSE, ...)
```

```
breaker(text.var)
```

```
word.split(text.var)
```

Arguments

`text.var` The text variable.

`apostrophe.remove`

logical. If TRUE removes apostrophe's from the output.

`...` further arguments passed to strip function.

Value

Returns a vector of striped words.

`breaker` - returns a vector of striped words and qdap recognized endmarks (i.e. ".", "!", "?", "*", "-").

Examples

```
## Not run:
bag.o.words(DATA$state)
by(DATA$state, DATA$person, bag.o.words)
lapply(DATA$state, bag.o.words)
bag.o.words("I'm going home!", apostrophe.remove = FALSE)

DATA
breaker(DATA$state)
by(DATA$state, DATA$person, breaker)
lapply(DATA$state, breaker)

word.split(c(NA, DATA$state))

## End(Not run)
```

blank2NA

Replace Blanks in Data Frame

Description

Replaces blank (empty) cells in a dataframe. generally, for internal use.

Usage

```
blank2NA(dataframe, missing = NA)
```

Arguments

dataframe	A dataframe with blank (empty) cells.
missing	Value to replace empty cells with.

Value

Returns a dataframe with blank spaces replaced.

See Also

[rm_row](#)

Examples

```
## Not run:
dat <- data.frame(matrix(sample(c(1:4, ""), 50, TRUE),
  10, byrow = TRUE), stringsAsFactors = FALSE)
dat
blank2NA(dat)

## End(Not run)
```

bracketX

*Bracket Parsing***Description**

bracketX - Apply bracket removal to character vectors.

bracketXtract - Apply bracket extraction to character vectors.

Usage

```
bracketX(text.var, bracket = "all", missing = NULL,
  names = FALSE)
```

```
bracketXtract(text.var, bracket = "all", with = FALSE)
```

Arguments

text.var	The text variable
bracket	The type of bracket (and encased text) to remove. This is one of the strings "curly", "square", "round", "angle" and "all". These strings correspond to: {, [, (, < or all four types.
missing	Value to assign to empty cells.
names	logical. If TRUE the sentences are given as the names of the counts.
with	logical. If TRUE returns the brackets and the bracketted text.

Value

bracketX - returns a vector of text with brackets removed.

bracketXtract - returns a list of vectors of bracketed text.

Author(s)

Martin Morgan and Tyler Rinker <tyler.rinker@gmail.com>.

References

<http://stackoverflow.com/questions/8621066/remove-text-inside-brackets-parens-and-or-braces>

Examples

```
## Not run:
examp2 <- structure(list(person = structure(c(1L, 2L, 1L, 3L),
  .Label = c("bob", "greg", "sue"), class = "factor"), text =
  c("I love chicken [unintelligible]!",
  "Me too! (laughter) It's so good.[interrupting]",
  "Yep it's awesome {reading}."), .Names =
  c("person", "text"), row.names = c(NA, -4L), class = "data.frame")

examp1
bracketX(examp2$text, 'square')
bracketX(examp2$text, 'curly')
bracketX(examp2$text)

examp2
bracketXtract(examp2$text, 'square')
bracketXtract(examp2$text, 'curly')
bracketXtract(examp2$text)
bracketXtract(examp2$text, with = TRUE)

paste2(bracketXtract(examp2$text, 'curly'), " ")

## End(Not run)
```

BuckleySaltonSWL

Buckley & Salton Stopword List

Description

A stopword list containing a character vector of stopwords.

Format

A character vector with 546 elements

Details

From Onix Text Retrieval Toolkit API Reference: "This stopword list was built by Gerard Salton and Chris Buckley for the experimental SMART information retrieval system at Cornell University. This stopword list is generally considered to be on the larger side and so when it is used, some implementations edit it so that it is better suited for a given domain and audience while others use this stopword list as it stands."

Note

Reduced from the original 571 words to 546.

References

<http://www.lextek.com/manuals/onix/stopwords2.html>

capitalizer	<i>Capitalize Select Words</i>
-------------	--------------------------------

Description

A helper function for word_list that allows the user to supply vectors of words to be capitalized.

Usage

```
capitalizer(text, caps.list = NULL, I.list = TRUE,
  apostrophe.remove = FALSE)
```

Arguments

text	A vector of words (generally from bag.o.words or breaker).
caps.list	A list of words to capitalize.
I.list	logical. If TRUE capitalizes I words and contractions.
apostrophe.remove	logical, asking if apostrophes have been removed. If TRUE will try to insert apostrophe's back into words appropriately.

Value

Returns a vector of capitalized words based on supplied capitalization arguments.

Note

Not intended for general use. Acts as a helper function to several qdap functions.

Examples

```
## Not run:
capitalizer(bag.o.words("i like it but i'm not certain"), "like")
capitalizer(bag.o.words("i like it but i'm not certain"), "like", FALSE)

## End(Not run)
```

clean	<i>Remove Escaped Characters</i>
-------	----------------------------------

Description

Pre process data to remove escaped characters

Usage

```
clean(text.var)
```

Arguments

text.var	The text variable
----------	-------------------

Value

Returns a vector of character strings with escaped characters removed.

Examples

```
## Not run:
x <- "I go \r
      to the \tnext line"
x
clean(x)

## End(Not run)
```

cm_code.blank	<i>Blank Code Transformation</i>
---------------	----------------------------------

Description

Transform codes with any binary operator combination.

Usage

```
cm_code.blank(x2long.obj, combine.code.list,
             rm.var = NULL, overlap = TRUE)
```

Arguments

x2long.obj	An object from cm_range2long, cm_time2long or cm_df2long
combine.code.list	A list of named character vectors of at least two code column names to combine
rm.var	Name of the repeated measures column.
overlap	logical, integer or character of binary operator + integer. If TRUE finds the overlap. If FALSE finds anywhere any of the codes occur. If integer finds that exact combination of overlaps. If character must be a logical vector c(>, <, ==, ==>, ==, !=) followed by an integer and wrapped with quotes.

Value

Returns a dataframe with transformed occurrences of supplied overlapping codes added.

Note

For most jobs `cm_code.transform` will work. This adds a bit of flexibility in exclusion and partial matching. The code column must be named `code` and your start and end columns must be named `"start"` and `"end"`.

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#), [cm_code.overlap](#), [cm_code.combine](#), [cm_code.exclude](#), [cm_code.transform](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)
foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='1:4, 10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
z <- cm_range2long(foo, foo2, v.name="time")
nots <- list(notAABB=qcv(AA, BB), notAACC=qcv(AA, CC), notBBCC=qcv(BB, CC))
z <- cm_code.blank(z, nots, "time", overlap=0)
z <- cm_code.blank(z, list(atleastAABBCC=qcv(AA, BB, CC)), "time", overlap=1)
z <- cm_code.blank(z, list(AACC=qcv(AA, CC)), "time", overlap=FALSE) #combined
cm_code.blank(z, list(AACCnoAA=qcv(AACC, AA)), "time", overlap=1) #remove the AA part

#WITH cm_time2long
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

y <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

dat <- cm_time2long(x, y)
cm_code.blank(dat, list(P=qcv(A, B), Q=qcv(B, C), R=qcv(A, B, C)),
  "variable", overlap=TRUE)
```

```
## End(Not run)
```

cm_code.combine	<i>Combine Codes</i>
-----------------	----------------------

Description

Combine all occurrences of codes into a new code.

Usage

```
cm_code.combine(x2long.obj, combine.code.list,
  rm.var = NULL)
```

Arguments

x2long.obj	An object from cm_range2long, cm_time2long or cm_df2long
combine.code.list	A list of named character vectors of at least two code column names to combine
rm.var	Name of the repeated measures column.

Value

Returns a dataframe with combined occurrences of supplied overlapping codes added.

Note

The code column must be named code and your start and end columns must be named "start" and "end".

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#), [cm_code.blank](#), [cm_code.exclude](#), [cm_code.overlap](#), [cm_code.transform](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='1:4, 10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
z <- cm_range2long(foo, foo2, v.name="time")
```



```

combines <- list(AB=qcv(AA, BB), ABC=qcv(AA, BB, CC))
cm_code.combine(x, list(AB=qcv(AA, BB)))
cm_code.combine(x, list(ALL=qcv(AA, BB, CC)))
cm_code.combine(z, combines, "time")

#WITH cm_time2long
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

y <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

dat <- cm_time2long(x, y)
cm_code.combine(dat, list(P=qcv(A, B), Q=qcv(B, C), R=qcv(A, B, C)), "variable")

## End(Not run)

```

cm_code.exclude	<i>Exclude Codes</i>
-----------------	----------------------

Description

Find the occurrences of n codes excluding the nth code. e.g. You have times/words coded for a teacher and you also have times/words coded for happiness. You can find all the happiness times excluding the teacher times or vice versa.

Usage

```
cm_code.exclude(x2long.obj, exclude.code.list,
  rm.var = NULL)
```

Arguments

x2long.obj	An object from cm_range2long, cm_time2long or cm_df2long
exclude.code.list	A list of named character vectors of at least two code column names to compare and exclude. The last column name is the one that will be excluded.
rm.var	Name of the repeated measures column.

Value

Returns a dataframe with n codes excluding the nth code.

Note

The code column must be named code and your start and end columns must be named "start" and "end".

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#), [cm_code.blank](#), [cm_code.combine](#), [cm_code.overlap](#), [cm_code.transform](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='1:4, 10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
z <- cm_range2long(foo, foo2, v.name="time")
cm_code.exclude(x, list(ABnoC=qcv(AA, BB, CC)))
cm_code.exclude(z, list(ABnoC=qcv(AA, BB, CC)), rm.var="time")
excludes <- list(ABnoB=qcv(AA, BB), ABnoC=qcv(AA, BB, CC))
cm_code.exclude(z, excludes, rm.var="time")
#WITH cm_time2long
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

y <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

dat <- cm_time2long(x, y)
cm_code.exclude(dat, list(P=qcv(A, B), Q=qcv(B, C), R=qcv(A, B, C)), "variable")

## End(Not run)
```

cm_code.overlap	<i>Find Co-occurrence Between Codes</i>
-----------------	---

Description

Combine co-occurrences of codes into a new code.

Usage

```
cm_code.overlap(x2long.obj, overlap.code.list,
  rm.var = NULL)
```

Arguments

x2long.obj	An object from cm_range2long, cm_time2long or cm_df2long
overlap.code.list	A list of named character vectors of at least two code column names to aggregate co-occurrences.
rm.var	Name of the repeated measures column.

Value

Returns a dataframe with co-occurrences of supplied overlapping codes added.

Note

The code column must be named code and your start and end columns must be named "start" and "end".

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#), [cm_code.combine](#), [cm_code.transform](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='1:4, 10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
z <- cm_range2long(foo, foo2, v.name="time")
combines <- list(AB=qcv(AA, BB), ABC=qcv(AA, BB, CC))
cm_code.overlap(x, list(AB=qcv(AA, BB)))
```

```

cm_code.overlap(x, list(ALL=qcv(AA, BB, CC)))
cm_code.overlap(z, combines, "time")

#WITH cm_time2long
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

y <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

dat <- cm_time2long(x, y)
cm_code.overlap(dat, list(P=qcv(A, B), Q=qcv(B, C), R=qcv(A, B, C)), "variable")

## End(Not run)

```

cm_code.transform	<i>Transform Codes</i>
-------------------	------------------------

Description

Transform co-occurences and/or combinations of codes into a new code(s).

Usage

```

cm_code.transform(x2long.obj, overlap.code.list = NULL,
  combine.code.list = NULL, exclude.code.list = NULL,
  rm.var = NULL)

```

Arguments

x2long.obj	An object from cm_range2long, cm_time2long or cm_df2long
overlap.code.list	A list of named character vektors of at least two code column names to aggregate co-occurences.
combine.code.list	A list of named character vektors of at least two code column names to combine
exclude.code.list	A list of named character vektors of at least two code column names to compare and exclude. The last column name is the one that will be excluded.
rm.var	Name of the repeated measures column.

Value

Returns a dataframe with overlapping, combined occurrences, and/or exclusion of supplied overlapping codes added.

Note

The code column must be named code and your start and end columns must be named "start" and "end".

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#), [cm_code.blank](#), [cm_code.combine](#), [cm_code.exclude](#), [cm_code.overlap](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='1:4, 10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
z <- cm_range2long(foo, foo2, v.name="time")
overlaps <- list(AB=qcv(AA, BB), ABC=qcv(AA, BB, CC))
cm_code.transform(x, overlap.code.list=list(AB=qcv(AA, BB)))
cm_code.transform(x, combine.code.list = list(ALL=qcv(AA, BB, CC)))
cm_code.transform(x, overlap.code.list=list(AB=qcv(AA, BB)),
  combine.code.list = list(ALL=qcv(AA, BB, CC)))
cm_code.transform(z, overlaps, rm.var="time")
cm_code.transform(z, overlaps,
  exclude.code.list=list(AABB_no_CC = qcv(AA, BB, CC)), rm.var="time")
#WITH cm_time2long
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

y <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)

dat <- cm_time2long(x, y)
cm_code.transform(dat, list(P=qcv(A, B), Q=qcv(B, C), R=qcv(A, B, C)),
  list(S=qcv(A, B), T=qcv(B, C), U=qcv(A, B, C)),
  list(ABnoC = qcv(A, B, C)), rm.var="variable")

## End(Not run)
```

cm_combine.dummy	<i>Find Co-occurrence Between Codes</i>
------------------	---

Description

Combine code columns where they co-occur.

Usage

```
cm_combine.dummy(cm.l2d.obj, combine.code,
  rm.var = "time", overlap = TRUE)
```

Arguments

cm.l2d.obj	An object from cm_long2dummy
combine.code	A list of named character vectors of at least two code column names to combine
rm.var	Name of the repeated measures column. Default is "time".
overlap	logical, integer or character of binary operator + integer. If TRUE finds the overlap. If FALSE finds anywhere any of the codes occur. If integer finds that exact combination of overlaps. If character must be a logical vector c(>, <, <=, >=, ==, !=) followed by an integer.

Value

Returns a dataframe with co-occurrences of provided code columns.

See Also

[cm_long2dummy](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
D1 <- cm_long2dummy(x)

z <- cm_range2long(foo, foo2, v.name="time")
D2 <- cm_long2dummy(z, "time")
cm_combine.dummy(D1, combine.code = list(AA=qcv(AA, BB)))
```

```

cm_combine.dummy(D1, combine.code = list(AB=qcv(AA, BB)), overlap=="=1")
cm_combine.dummy(D1, combine.code = list(AB=qcv(AA, BB)), overlap!="=1")
D1 <- cm_combine.dummy(D1, combine.code = list(AB=qcv(AA, BB)), overlap=0)
D1 <- cm_combine.dummy(D1, combine.code = list(CAB=qcv(AB, CC)), overlap=FALSE)

combines <- list(AB=qcv(AA, BB), ABC=qcv(AA, BB, CC))
cm_combine.dummy(D1, combine.code = combines)
cm_combine.dummy(D2, combine.code = combines)

## End(Not run)

```

cm_df.fill

*Range Coding of a Code Matrix***Description**

Allows range coding of words for efficient coding.

Usage

```

cm_df.fill(dataframe, ranges, value = 1, text.var = NULL,
           code.vars = NULL, transform = FALSE)

```

Arguments

dataframe	A dataframe containing a text variable.
ranges	A named list of ranges to recode. Names correspond to code names in dataframe.
value	The recode value. Takes a vector of length one or a vector of length equal to the number of code columns.
text.var	The name of the text variable.
code.vars	Optional vector of codes.
transform	logical. If TRUE the words are located across the top of dataframe.

Value

Generates a dummy coded dataframe.

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_df.temp](#), [cm_df2long](#)

Examples

```
## Not run:
codes <- qcv(dc, sf, wes, pol, rejk, lk, azx, mmm)
X <- cm_df.temp(DATA, "state", codes)
cm_df.fill(X, list(dc=c(1:3, 5), sf=c(4, 6:9, 11), wes=0, pol=0, rejk=0,
lk=0, azx=1:30, mmm=5))
cm_df.fill(X, list(sf=c(4, 6:9, 11), dc=c(1:3, 5), azx=1:30, mmm=5))

## End(Not run)
```

cm_df.temp	<i>Break Transcript Dialogue into Blank Code Matrix</i>
------------	---

Description

Breaks transcript dialogue into words while retaining the demographic factors associate with each word. The codes argument provides a matrix of zeros that can serve as a dummy coded matrix of codes per word.

Usage

```
cm_df.temp(dataframe, text.var, codes = NULL, csv = TRUE,
file.name = NULL, transpose = FALSE, strip = FALSE)
```

Arguments

dataframe	A dataframe containing a text variable.
text.var	The name of the text variable.
codes	Optional list of codes.
csv	logical. If TRUE creates a csv in the working directory.
file.name	The name of the csv file. If NULL defaults to the dataframe name.
transpose	logical. If TRUE transposes the dataframe so that the text is across the top.
strip	logical. If TRUE all punctuation is removed.

Value

Generates a dataframe, and optional csv file, of individual words while maintaing demographic information. If a vector of codes is provided the outcome is a matrix of words used by codes filled with zeros. This dataframe is useful for dummy coded (1=yes code exists; 2-no it does not) representation of data and can be used for visualizations and statistical analysis.

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

```
cm\_range2long, #`cm\_df.fill`
```


Examples

```
## Not run:
codes <- qcv(dc, sf, wes, pol, rejk, lk, azx, mmm)
cm_df.temp(DATA, "state", codes)
cm_df.temp(DATA, "state", codes, transpose = TRUE)
head(cm_df.temp(raj.act.1, "dialogue", codes))
cm_df.temp(raj.act.1, "dialogue", codes, transpose = TRUE)[, 1:9]

## End(Not run)
```

cm_df.transcript	<i>Transcript With Word Number</i>
------------------	------------------------------------

Description

Out put a transcript with word number/index above for easy input back into qdap after coding.

Usage

```
cm_df.transcript(text.var, grouping.var, file = NULL,
  indent = 4, width = 70)
```

Arguments

text.var	text.var The text variable
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
file	A connection, or a character string naming the file to print to (e.g. .doc, .txt).
indent	Number of spaces to indent.
width	Width to output the file (defaults to 70; this is generally a good width and indent for a .docx file).

Value

Returns a transcript by grouping variable with word number above each word. This makes use with cm_df2long transfer/usage easier because the researcher has coded on a transcript with the numeric word index already.

Note

It is recommended that the researcher actually codes on the out put from this file. If a file already exists cm_df.transcript will append to that file.

Author(s)

DWin, Gavin Simpson and Tyler Rinker <tyler.rinker@gmail.com>.

See Also

See Also as [cm_df2long](#) See Also as [cm_df.temp](#)

Examples

```
## Not run:
with(mraja1spl, cm_df.transcript(dialogue, list(person)))
with(mraja1spl, cm_df.transcript(dialogue, list(sex, fam.aff, died)))
with(mraja1spl, cm_df.transcript(dialogue, list(person), file="foo.doc"))
# delete("foo.doc") #delete the file just created

## End(Not run)
```

cm_df2long

Transform Codes to Start-End Durations

Description

Transforms the range coding structure(s) from `cm_df$temp` (in list format) into a data frame of start and end durations in long format.

Usage

```
cm_df2long(df.temp.obj, v.name = "variable",
  list.var = TRUE, code.vars = NULL, no.code = NA,
  add.start.end = TRUE, repeat.vars = NULL,
  rev.code = FALSE)
```

Arguments

<code>df.temp.obj</code>	a character vector of names of object(s) created by <code>cm_df\$temp</code> , a list of <code>cm_df\$temp</code> created objects or a data frame created by <code>cm_df\$temp</code> .
<code>v.name</code>	an optional name for the column created for the <code>list.var</code> argument
<code>list.var</code>	logical. If TRUE creates a column for the data frame created by each time.list passed to <code>cm_t2l</code>
<code>code.vars</code>	a character vector of code variables. If NULL uses all variables from the first column after the column named <code>word.num</code> .
<code>no.code</code>	the value to assign to no code; default is NA
<code>add.start.end</code>	logical. If TRUE adds a column for start and end times
<code>repeat.vars</code>	a character vector of repeated/stacked variables. If NULL uses all non <code>code.vars</code> variables.
<code>rev.code</code>	logical. If TRUE reverses the order of <code>code.vars</code> and <code>no.code</code> variables.

Value

Generates a data frame of start and end times for each code.

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_time2long](#), [cm_range2long](#), [cm_df.temp](#)

Examples

```
## Not run:
#' codes <- qcv(dc, sf, wes, pol, rejk, lk, azx, mmm)
x1 <- cm_df.temp(DATA, "state", codes)
cm_df2long(x1, code.vars = codes)
x1[, 7:14] <- lapply(7:14, function(i) sample(0:1, nrow(x1), TRUE))
cm_df2long(x1, code.vars = codes)

## End(Not run)
```

cm_distance	<i>Distance Matrix Between Codes</i>
-------------	--------------------------------------

Description

Generate distance measures to ascertain a mean distance measure between codes.

Usage

```
cm_distance(dataframe, time.var = NULL, parallel = FALSE,
  code.var = "code", causal = FALSE, start.var = "start",
  end.var = "end", mean.digits = 2, sd.digits = 2,
  stan.digits = 2)
```

Arguments

dataframe	a data frame from the cm_x2long family (cm_range2long; cm_df2long; cm_time2long)
time.var	an optional variable to split the dataframe by (if you have data that is by various times this must be supplied).
parallel	logical. If TRUE runs the cm_distance on multiple cores. This is effective with larger data sets but may actually be slower with smaller data sets.
code.var	the name of the code variable column. Defaults to "codes" as outputted by x2long family
causal	logical. If TRUE measures the distance between x and y given that x must precede y
start.var	the name of the start variable column. Defaults to "start" as outputted by x2long family
end.var	the name of the end variable column. Defaults to "end" as outputted by x2long family
mean.digits	the number of digits to be displayed in the mean matrix
sd.digits	the number of digits to be displayed in the sd matrix

Value

An object of the class `cm.dist`. This is a list of `n` lists with the following components per each list (`time.var`):

<code>mean</code>	A distance matrix of average distances between codes
<code>sd</code>	A matrix of standard deviations of distances between codes
<code>n</code>	A matrix of counts of distances between codes
<code>combined</code>	A matrix of combined mean, sd and n of distances between codes
<code>standardized</code>	A matrix of standardized values of distances between codes. The closer a value is to zero the closer two codes relate.

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='02:03, 05'),
  BB = qcv(terms='1:2, 3:10'),
  CC = qcv(terms='1:9, 100:150')
)
foo2 <- list(
  AA = qcv(terms='40'),
  BB = qcv(terms='50:90'),
  CC = qcv(terms='60:90, 100:120, 150'),
  DD = qcv(terms='')
)
(dat <- cm_range2long(foo, foo2, v.name = "time"))
(out <- cm_distance(dat, time.var = "time", causal=T))
names(out)
names(out$foo2)
out$foo2
#=====
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 6.32:7.00, 9.00, 10.00:11.00, 59.56"),
  B = qcv(terms = "3.01:3.02, 5.01, 19.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.32:7.00, 9.00, 17.01")
)
dat <- cm_time2long(x)
gantt_wrap(dat, "code", border.color = "black", border.size = 5, sig.dig.line.freq = -2)
(a <- cm_distance(dat))
names(a)
names(a$dat)
a$dat

## End(Not run)
```

cm_dummy2long

Convert cm_combine.dummy Back to Long

Description

`cm_combine.dummy` back to long.

Usage

```
cm_dummy2long(cm.comb.obj, rm.var = "time")
```

Arguments

cm.comb.obj An object from cm_combine.dummy

rm.var Name of the repeated measures column. Default is "time".

Value

Returns a dataframe with co-occurrences of provided code columns.

See Also

[cm_long2dummy](#), [cm_combine.dummy](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1:10'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4:8'),
  BB = qcv(terms='10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
D1 <- cm_long2dummy(x)

z <- cm_range2long(foo, foo2, v.name="time")
D2 <- cm_long2dummy(z, "time")
cm_combine.dummy(D1, combine.code = list(AB=qcv(AA, BB)))

combines <- list(AB=qcv(AA, BB), ABC=qcv(AA, BB, CC))

A <- cm_combine.dummy(D2, combine.code = combines)
B <- cm_combine.dummy(D1, combine.code = combines)

cm_dummy2long(A)
cm_dummy2long(B, "time")

## End(Not run)
```

cm_long2dummy

*Stretch and Dummy Code cm_xxx2long***Description**

Stretches and dummy codes a cm_xxx2long dataframe to allow for combining columns.

Usage

```
cm_long2dummy(dataframe, rm.var = NULL, code = "code",
  start = "start", end = "end")
```

Arguments

dataframe	A dataframe that contains the person variable.
rm.var	An optional character argument of the name of a repeated measures column.
code	A character argument of the name of a repeated measures column. Default is "code".
start	A character argument of the name of a repeated measures column. Default is "start".
end	A character argument of the name of a repeated measures column. Default is "end".

Value

Returns a dataframe or a list of stretched and dummy coded dataframe(s).

See Also

[cm_range2long](#), [cm_time2long](#), [cm_df2long](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:3, 5:6')
)

foo2 <- list(
  AA = qcv(terms='4'),
  BB = qcv(terms='10:12'),
  CC = qcv(terms='1, 11, 15:20'),
  DD = qcv(terms='')
)

x <- cm_range2long(foo)
cm_long2dummy(x)

z <- cm_range2long(foo, foo2, v.name="time")
cm_long2dummy(z, "time")
```

```
## End(Not run)
```

cm_range.temp	<i>Range Code Sheet</i>
---------------	-------------------------

Description

Generates a range coding sheet for coding words.

Usage

```
cm_range.temp(codes, file = NULL)
```

Arguments

codes	List of codes.
file	A connection, or a character string naming the file to print to (.txt is recommended).

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_time.temp](#)

Examples

```
## Not run:
cm_range.temp(qcv(AA, BB, CC), file = "foo.txt")
# delete("foo.txt")

## End(Not run)
```

cm_range2long	<i>Transform Codes to Start-End Durations</i>
---------------	---

Description

Transforms the range coding structure(s) from cm_range.temp (in list format) into a data frame of start and end durations in long format.

Usage

```
cm_range2long(..., v.name = "variable", list.var = TRUE,
  debug = TRUE)
```

Arguments

...	list object(s) in the form generated by cm_time.temp.
v.name	sn optional name for the column created for the list.var argument.
list.var	logical. If TRUE creates a column for the data frame created by each time.list passed to cm_t2l.
star.end	logical. If TRUE outputs stop and end times for each cm_time.temp list object.
debug	logical. If TRUE debugging mode is on. cm_time2long will return possible errors in time span inputs.

Value

Generates a data frame of start and end times for each code.

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_df2long](#) [cm_time.temp](#)

Examples

```
## Not run:
foo <- list(
  AA = qcv(terms='1'),
  BB = qcv(terms='1:2, 3:10, 19'),
  CC = qcv(terms='1:9, 100:150')
)

foo2 <- list(
  AA = qcv(terms='40'),
  BB = qcv(terms='50:90'),
  CC = qcv(terms='60:90, 100:120, 150'),
  DD = qcv(terms='')
)
dat <- cm_range2long(foo, foo2, v.name = "time")
gantt_wrap(dat, "code", "time")

## End(Not run)
```

cm_time.temp

Time Span Code Sheet

Description

Generates a time span coding sheet and coding format sheet.

Usage

```
cm_time.temp(codes, start = ":00", end = NULL,
             file = NULL)
```

Arguments

codes	List of codes.
start	A character string in the form of "00:00" indicating start time (default is ":00").
end	A character string in the form of "00:00" indicating end time.
file	A connection, or a character string naming the file to print to (.txt is recommended).

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_range.temp](#),

Examples

```
## Not run:
cm_time.temp(qcv(AA, BB, CC), ":30", "7:40", file = "foo.txt")
# delete("foo.txt")
x <- list(
  transcript_time_span = qcv(terms='00:00 - 1:12:00'),
  A = qcv(terms='2.40:3.00, 5.01, 6.62:7.00, 9.00'),
  B = qcv(terms='2.40, 3.01:3.02, 5.01, 6.62:7.00, 9.00, 1.12.00:1.19.01'),
  C = qcv(terms='2.40:3.00, 5.01, 6.62:7.00, 9.00, 17.01')
)
cm_time2long(x)
cm_time.temp(qcv(AA, BB, CC))

## End(Not run)
```

cm_time2long

Transform Codes to Start-End Times

Description

Transforms the range coding structure(s) from `cm_time.temp` (in list format) into a data frame of start and end times in long format.

Usage

```
cm_time2long(..., v.name = "variable", list.var = TRUE,
             start.end = FALSE, debug = TRUE)
```

Arguments

...	List object(s) in the form generated by <code>cm_time.temp</code> .
<code>v.name</code>	An optional name for the column created for the <code>list.var</code> argument
<code>list.var</code>	logical. If TRUE creates a column for the data frame created by each <code>time.list</code> passed to <code>cm_t2l</code> .
<code>start.end</code>	logical. If TRUE outputs stop and end times for each <code>cm_time.temp</code> list object.
<code>debug</code>	logical. If TRUE debugging mode is on. <code>cm_time2long</code> will return possible errors in time span inputs.

Value

Generates a data frame of start and end times for each code.

References

Miles, M. B. & Huberman, A. M. (1994). An expanded sourcebook: Qualitative data analysis. 2nd ed. Thousand Oaks, CA: SAGE Publications.

See Also

[cm_df2long](#) [cm_time.temp](#)

Examples

```
## Not run:
x <- list(
  transcript_time_span = qcv(00:00 - 1:12:00),
  A = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00"),
  B = qcv(terms = "2.40, 3.01:3.02, 5.01, 6.02:7.00, 9.00, 1.12.00:1.19.01"),
  C = qcv(terms = "2.40:3.00, 5.01, 6.02:7.00, 9.00, 17.01")
)
dat <- cm_time2long(x)
gantt_wrap(dat, "code", border.color = "black", border.size = 5)

## End(Not run)
```

colSplit	<i>Separate a Column Pasted by paste2</i>
----------	---

Description

Separates a `paste2` column into separate columns.

Usage

```
colSplit(column, col.sep = ".", name.sep = "&")
```

Arguments

<code>column</code>	The pasted vector.
<code>col.sep</code>	The column separator used in <code>paste2</code> .
<code>name.sep</code>	Name separator used in the column (internal use within <code>colsplit2df</code>).

Value

Returns a dataframe of split columns.

See Also

[colsplit2df](#), [paste2](#)

Examples

```
## Not run:
(foo <- paste2(C02[, 1:3]))
colSplit(foo)
(bar <- paste2(mtcars[, 1:3], sep="|"))
colSplit(bar, col.sep = "|")

## End(Not run)
```

colsplit2df

Wrapper for colSplit that Returns a Dataframe

Description

Wrapper for colSplit that returns a dataframe.

Usage

```
colsplit2df(dataframe, splitcol = 1, new.names = NULL,
  sep = ".", keep.orig = FALSE)
```

Arguments

dataframe	A dataframe with a column that has been pasted together.
splitcol	The name of the column that has been pasted together.
new.names	A character vector of new names to assign to the columns. Default attempts to extract the original names before the paste.
sep	The character that used in paste2 to paste the columns.
keep.orig	logical. If TRUE the original pasted column will be retained as well.

Value

Returns a dataframe with the pasted column cplit into new columns.

See Also

[colSplit](#), [paste2](#)

Examples

```
## Not run:
C02$'Plant&Type&Treatment' <- paste2(C02[, 1:3])
C02 <- C02[, -c(1:3)]
head(colsplit2df(C02, 3))
head(colsplit2df(C02, 3, qcv(A, B, C)))
head(colsplit2df(C02, 3, qcv(A, B, C), keep.orig=TRUE))
head(colsplit2df(C02, "Plant&Type&Treatment"))
C02 <- datasets::C02

## End(Not run)
```

common

Find Common Words Between Groups

Description

Find common words between grouping variables (e.g. people).

Usage

```
common(x, ...)

## Default S3 method:
common(..., overlap = "all",
  equal.or = "equal")

## S3 method for class 'list'
common(word.list, overlap = "all",
  equal.or = "more")
```

Arguments

<code>word.list</code>	A list of names character vectors.
<code>overlap</code>	Minimum/exact amount of overlap.
<code>equal.or</code>	A character vector of <code>c("equal", "greater", "more", "less")</code> .
<code>...</code>	In lieu of <code>word.list</code> the user may input <code>n</code> number of character vectors.

Value

Returns a dataframe of all words that match the criteria set by `overlap` and `equal.or`.

NULL

NULL

Examples

```
## Not run:
a <- c("a", "cat", "dog", "the", "the")
b <- c("corn", "a", "chicken", "the")
d <- c("house", "feed", "a", "the", "chicken")
common(a, b, d, overlap=2)
common(a, b, d, overlap=3)

r <- list(a, b, d)
common(r)
common(r, overlap=2)

common(word_list(DATA$state, DATA$person)$cwl, overlap = 2)

## End(Not run)
```

convert*Convert Seconds to h:m:s*

Description

Converts a vector of seconds to h:m:s

Usage

```
convert(x)
```

Arguments

x A vector of times in seconds.

Value

Returns a vector of times in h:m:s format. Generally, this function is for internal use.

Examples

```
## Not run:
convert(c(256, 3456, 56565))

## End(Not run)
```

DATA

Fictitious Classroom Dialogue

Description

A fictitious dataset useful for small demonstrations.

Format

A data frame with 11 rows and 5 variables

Details

- person. Speaker
- sex. Gender
- adult. Dummy coded adult (0-no; 1-yes)
- state. Statement (dialogue)
- code. Dialogue coding scheme

DATA2

Fictitious Repeated Measures Classroom Dialogue

Description

A repeated measures version of the [DATA](#) dataset.

Format

A data frame with 74 rows and 7 variables

Details

- day. Day of observation
- class. Class period/subject of observation
- person. Speaker
- sex. Gender
- adult. Dummy coded adult (0-no; 1-yes)
- state. Statement (dialogue)
- code. Dialogue coding scheme

delete	<i>Easy File Handling</i>
--------	---------------------------

Description

delete - Deletes files and directories.

folder - Create a folder/directory.

Usage

```
delete(file = NULL)
```

```
folder(folder.name = NULL)
```

Arguments

file	The name of the file in the working directory or the path to the file to be deleted. If NULL provides a menu of files from the working directory.
folder.name	The name of the folder to be created. Default NULL creates a file in the working directory with the creation date and time stamp.

Value

delete permanently removes a file/directory.

folder creates a folder/directory.

See Also

[unlink](#), [file.remove](#), [dir.create](#)

Examples

```
## Not run:  
(x <- folder("DELETE.ME"))  
which(dir() == "DELETE.ME")  
delete("DELETE.ME")  
which(dir() == "DELETE.ME")  
  
## End(Not run)
```

 DICTIONARY

Nettalk Corpus Syllable Data Set

Description

A dataset containing syllable counts.

Format

A data frame with 20137 rows and 2 variables

Details

- word. The word
- syllables. Number of syllables

Note

This data set is based on the Nettalk Corpus but has some researcher word deletions and additions based on the needs of the `syllable.sum` algorithm.

References

Sejnowski, T.J., and Rosenberg, C.R. (1987). "Parallel networks that learn to pronounce English text" in *Complex Systems*, 1, 145-168. Retrieved from: [http://archive.ics.uci.edu/ml/datasets/Connectionist+Bench+\(Nettalk+Corpus\)](http://archive.ics.uci.edu/ml/datasets/Connectionist+Bench+(Nettalk+Corpus))

[UCI Machine Learning Repository website](#)

 dissimilarity

Dissimilarity Statistics

Description

Uses the distance function to calculate dissimilarity statistics by grouping variables.

Usage

```
dissimilarity(text.var, grouping.var = NULL,
  method = "prop", diag = FALSE, upper = FALSE, p = 2,
  digits = 3)
```


Arguments

text.var	A text variable or word frequency matrix object.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
method	Distance methods (see dist function). If "prop" (the default; the result is 1 - "binary").
diag	logical. If True returns the diagonals of the matrix
upper	logical. If True returns the upper triangle of the matrix
p	The power of the Minkowski distance
digits	integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed

Value

Returns a matrix of dissimilarity values (the agreement between text).

See Also

[dist](#)

Examples

```
## Not run:
with(DATA, dissimilarity(state, list(sex, adult)))
with(DATA, dissimilarity(state, person, diag = TRUE))

## End(Not run)
```

distTab	<i>SPSS Style Frequency Tables</i>
---------	------------------------------------

Description

Generates a dsitribution table for vectors, matrices and dataframes.

Usage

```
distTab(dataframe, breaks = NULL, digits = 2, ...)
```

Arguments

dataframe	A vector or data.frame object.
breaks	Either a numeric vector of two or more cut points or a single number (greater than or equal to 2) giving the number of intervals into which x is to be cut.
digits	Integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed
...	Other variables passed to cut.

Value

Returns a list of data frames (or singular data frame for a vector) of frequencies, cumulative frequencies, percentages and cumulative percentages for each interval.

See Also

[cut](#)

Examples

```
## Not run:
distTab(rnorm(10000), 10)
distTab(sample(c("red", "blue", "gray"), 100, T), right = FALSE)
distTab(CO2, 4)
distTab(mtcars)
distTab(mtcars, 4)

wdst <- with(mraja1spl, word_stats(dialogue, list(sex, fam.aff, died)))
distTab(wdst$gts)

## End(Not run)
```

diversity

Diversity Statistics

Description

Transcript apply diversity statistics.

Usage

```
diversity(text.var, grouping.var = NULL, digits = 3)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
digits	Number of decimal places to round.

Value

Returns a dataframe of various diversity related indices for Shannon, collision, Berger Parker and Brillouin.

Examples

```
## Not run:
colsplit2df(with(mraja1spl, diversity(dialogue, list(sex, died, fam.aff))))

## End(Not run)
```

duplicates	<i>Find Duplicated Words in a Text String</i>
------------	---

Description

Find duplicated word/word chunks in a string. Intended for internal use.

Usage

```
duplicates(string, threshold = 1)
```

Arguments

string	A character string.
threshold	An interger of the minimal number of repeats.

Value

Returns a vector of all duplicated words/chunks.

Examples

```
## Not run:  
duplicates(DATA$state)  
duplicates(DATA$state[1])  
  
## End(Not run)
```

emoticon	<i>Emoticons Data Set</i>
----------	---------------------------

Description

A dataset containing common emoticons (adapted from [Popular Emoticon List](#)).

Format

A data frame with 81 rows and 2 variables

Details

- meaning. The meaning of the emoticon
- emoticon. The graphic representation of the emoticon

References

http://www.lingo2word.com/lists/emoticon_listH.html

endf	<i>Test for Incomplete Sentences</i>
------	--------------------------------------

Description

Test for incomplete sentences and optionally remove them.

Usage

```
endf(dataframe, text.var, warning.report = TRUE,
      which.mode = FALSE)
```

Arguments

dataframe	A dataframe that contains the person and text variable.
text.var	The text variable.
warning.report	logical. If TRUE prints a warning of regarding removal of incomplete sentences.
which.mode	logical. If TRUE outputs two logical vectors: NOT (logical test of not being an incomplete sentence) and INC (logical test of being an incomplete sentence)

Value

Generates a dataframe with incomplete sentences removed.

Examples

```
## Not run:
dat <- sentSplit(DATA, "state", stem.col = FALSE)
dat$state[c(2, 5)] <- paste(strip(dat$state[c(2, 5)]), "|")
endf(dat, "state")
endf(dat, "state", warning.report = FALSE)
endf(dat, "state", which.mode = TRUE)

## End(Not run)
```

env.syl	<i>Syllable Lookup Environment</i>
---------	------------------------------------

Description

A dataset containing a syllable lookup environment (see `link[qdap]{DICTIONARY}`).

Format

A environment with

Details

For internal use.

References

[UCI Machine Learning Repository website](#)

exclude

Exclude Elements From a Vector

Description

Quickly exclude words from a word list

Usage

```
exclude(word.list, ...)
```

Arguments

<code>word.list</code>	A list of words/terms to exclude from.
<code>...</code>	A vector or sinle length objects to be excluded from the word.list.

Value

Returns a vector with the excluded terms removed.

Examples

```
## Not run:
Top25Words
exclude(Top25Words, qcv(the, of, and))
exclude(Top25Words, "the", "of", "an")
exclude(1:10, 3, 4)
exclude(1:10, 3:4)

#Using with term.match and termco.a
exclude(term.match(DATA$state, qcv(th), FALSE), "truth")
termco.a(DATA$state, DATA$person, exclude(term.match(DATA$state, qcv(th),
  FALSE), "truth"))
MTCH.LST <- exclude(term.match(DATA$state, qcv(th, i)), qcv(truth, stinks))
termco.a(DATA$state, DATA$person, MTCH.LST)

## End(Not run)
```

formality

*Formality Score***Description**

Transcript apply formality score by grouping variable(s).

Usage

```
formality(text.var, grouping.var = NULL, plot = FALSE,
          sort.by.formality = TRUE, digits = 2, point.pch = 20,
          point.cex = 0.5, point.colors = c("gray65", "red"),
          bar.colors = NULL, min.wrdcnt = NULL, ...)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates formality score for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
plot	logical. Provides a visualization for the results
sort.by.formality	logical. If TRUE orders the results by formality score.
digits	The number of digits displayed.
point.pch	The plotting symbol.
point.cex	The plotting symbol size.
point.colors	A vector of colors (length of two) to plot word count and formality score.
bar.colors	A palette of colors to supply to the bars in the visualization. If two palettes are provided to the two bar plots respectively.
min.wrdcnt	A minimum word count threshold that must be achieved to be considered in the results. Default includes all subgroups.

Details

Heylighen & Dewaele(2002)'s formality score is calculated as:

$$F = 50\left(\frac{n_f - n_c}{N} + 1\right)$$

Where:

$$f = \{\textit{noun}, \textit{adjective}, \textit{preposition}, \textit{article}\}$$

$$c = \{\textit{pronoun}, \textit{verb}, \textit{adverb}, \textit{interjection}\}$$

$$N = \sum (f + c + \textit{conjunctions})$$

Value

A list containing at the following components:

text	The text variable
POStagged	Raw part of speech for every word of the text variable
POSprop	Part of speech proportion for every word of the text variable
POSfreq	Part of speech count for every word of the text variable
pos.by.freq	The part of speech count for every word of the text variable by grouping variable(s)
pos.by.prop	The part of speech proportion for every word of the text variable by grouping variable(s)
form.freq.by	The nine broad part of speech categories count for every word of the text variable by grouping variable(s)
form.prop.by	The nine broad part of speech categories proportion for every word of the text variable by grouping variable(s)
formality	Formality scores by grouping variable(s)
pos.resaped	An expanded formality scores output (grouping, word.count, pos & form.class) by word

Note

Heylighen & Dewaele(2002) say "At present, a sample would probably need to contain a few hundred words for the measure to be minimally reliable. For single sentences, the F-value should only be computed for purposes of illustration".

References

Heylighen, F., & Dewaele, J.M. (2002). Variation in the contextuality of language: An empirical measure. *Context in Context, Special issue of Foundations of Science*, 7 (3), 293-340.

Examples

```
## Not run:
with(DATA, formality(state, person))
with(DATA, formality(state, list(sex, adult), plot = TRUE))
rajDEM <- key_merge(raj, raj.demographics, 'person')
with(raj, formality(rajPOS, act, plot=TRUE))
with(raj, formality(rajPOS, person, plot=TRUE, bar.colors="Dark2"))
with(raj, formality(rajPOS, person, plot=TRUE, bar.colors=c("Dark2", "Set1")))
with(raj, formality(rajPOS, list(person, act), plot=TRUE, bar.colors="Set1"))
with(rajDEM, formality(rajPOS, sex, plot=TRUE, bar.colors="RdBu"))
with(rajDEM, formality(rajPOS, list(fam.aff, sex), plot=TRUE,
  bar.colors="RdBu"))
with(rajDEM, formality(rajPOS, list(died, fam.aff), plot=TRUE,
  bar.colors="RdBu", point.cex=2, point.pch = 3))
raj.form <- with(rajDEM, formality(rajPOS, list(died, sex), plot=TRUE,
  bar.colors="RdBu", point.cex=2, point.pch = "|"))
names(raj.form)
colsplit2df(raj.form$formality)

## End(Not run)
```

gantt	<i>Generate Unit Spans</i>
-------	----------------------------

Description

Generates start and end times of supplied text selections (i.e. text selections are determined by any number of grouping variables).

Usage

```
gantt(text.var, grouping.var, plot = TRUE,
      units = "words", sums = FALSE, plot.colors = NULL,
      box.color = NULL, col.sep = "_")
```

Arguments

text.var	The text variable
grouping.var	The grouping variables. Also takes a single grouping variable or a list of 1 or more grouping variables.
plot	logical. If TRUE plots the start-end times as a gantt plot.
units	The unit of measurement to analyze. One of the strings "character", "syllable", "word", or "sentence".
sums	logical. If TRUE reports and optionally plots the total units used by grouping variable(s).
plot.colors	The colors of the Gantt plot bars. Either a single color or a length equal to the number of grouping variable(s).
box.color	A single color of the box around the Gantt plot bars.

Value

Returns a data frame of start and end times by grouping variable(s) or optionally returns a list of two: (1) A data frame of the total units used by grouping variable(s) and (2) a data frame of start and end times by grouping variable(s). Optionally plots a gantt plot of the returned data.

Note

For repeated measures data output use `gantt_rep`; for a convenient wrapper that takes text and generates plots use `gantt_plot`; and for a flexible gantt plot that works with code matrix functions (cm) use `gantt_wrap`.

Author(s)

DigEmAll (stackoverflow.com) and Tyler Rinker <tyler.rinker@gmail.com>.

References

Clark, W. & Gantt, H. (1922) The Gantt chart, a working tool of management. New York, Ronald Press.

See Also

[gantt_rep](#), [gantt_wrap](#), [gantt_plot](#)

Examples

```
## Not run:
gantt(DATA$state, DATA$person)
gantt(DATA$state, DATA$person, sums = TRUE)
gantt(DATA$state, list(DATA$sex, DATA$adult))
gantt(mraja1$dialogue, mraja1$person) #hard to see without box color
gantt(mraja1$dialogue, mraja1$sex)
gantt(mraja1$dialogue, mraja1$person, box.col = "black")
gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), plot.colors = NULL)
gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), plot.colors = "black")
gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), plot = FALSE)
gantt(mraja1$dialogue, mraja1$person, units = "characters", box.color = "black")
gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), units = "characters")
with(mraja1, gantt(dialogue, list(fam.aff, sex, died),
  units = "characters", sums = TRUE))
gantt(mraja1$dialogue, mraja1$person, units = "syllables", box.color = "black", sums = TRUE)
gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), units = "syllables")

(dat <- gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex), units = "sentences",
  plot.colors = 'black', sums = TRUE, col.sep = "_")$gantt.df)
gantt_wrap(dat, fam.aff_sex, title = "Gantt Plot")

## End(Not run)
```

gantt_plot

Gantt Plot

Description

A convenience function that wraps `gantt`, `gantt_rm` and `gantt_wrap` into a single plotting function.

Usage

```
gantt_plot(text.var, grouping.var, rm.var = NULL,
  fill.var = NULL, xlab = "duration (in words)",
  units = "words", col.sep = "_", ...)
```

Arguments

<code>text.var</code>	The text variable.
<code>grouping.var</code>	The grouping variables. Also takes a single grouping variable or a list of 1 or more grouping variables.
<code>rm.var</code>	An optional single vector or list of 1 or 2 of repeated measures to facet by
<code>fill.var</code>	An optional variable to fill the code stips by.
<code>units</code>	The unit of measurement.
<code>col.sep</code>	The column separator.
<code>...</code>	Other arguments passed to gantt_wrap .

Value

Returns a Gantt style visualization. Invisibly returns the ggplot2 list object.

Note

For non repeated measures data/plotting use `gantt`; for repeated measures data output use `gantt_rep`; and for a flexible gantt plot that works with code matrix functions (cm) use `gantt_wrap`.

References

Clark, W. & Gantt, H. (1922) The Gantt chart, a working tool of management. New York, Ronald Press.

See Also

[gantt](#) [gantt_rep](#), [gantt_wrap](#),

Examples

```
## Not run:
with(rajSPLIT, gantt_plot(text.var = dialogue, grouping.var = person, size=4))
with(rajSPLIT, gantt_plot(text.var = dialogue, grouping.var =
  list(fam.aff, sex), rm.var = act,
  title = "Romeo and Juliet's dialogue"))
with(rajSPLIT, gantt_plot(dialogue, list(fam.aff, sex), act, transform=T))
rajSPLIT2 <- rajSPLIT
rajSPLIT2$newb <- as.factor(sample(LETTERS[1:2], nrow(rajSPLIT2),
  replace=TRUE))
z <- with(rajSPLIT2, gantt_plot(dialogue, list(fam.aff, sex),
  list(act, newb), size = 4))
z + theme(panel.margin = unit(1, "lines")) + scale_colour_grey()
z + scale_colour_brewer(palette="Dark2")

## End(Not run)
```

gantt_rep

Generate Unit Spans for Repeated Measures

Description

Produces start and end times for occurrences for each repeated measure condition.

Usage

```
gantt_rep(rm.var, text.var, grouping.var,
  units = "words", col.sep = "_")
```

Arguments

<code>rm.var</code>	An optional single vector or list of 1 or 2 of repeated measures to facet by.
<code>text.var</code>	The text variable.
<code>grouping.var</code>	The grouping variables. Also takes a single grouping variable or a list of 1 or more grouping variables.
<code>units</code>	The unit of measurement to analyze. One of the strings "character", "syllable", "word", or "sentence".

Value

Returns a data frame of start and end times by repeated measure and grouping variable(s)

Note

For non repeated measures data/plotting use `gantt`; for a convenient wrapper that takes text and generates plots use `gantt_plot`; and for a flexible gantt plot that works with code matrix functions (cm) use `gantt_wrap`.

References

Clark, W. & Gantt, H. (1922) The Gantt chart, a working tool of management. New York, Ronald Press.

See Also

[gantt](#), [gantt_wrap](#), [gantt_plot](#)

Examples

```
## Not run:
(dat3 <- with(rajSPLIT, gantt_rep(act, dialogue, list(fam.aff, sex), units = "words",
  col.sep = "_"))
gantt_wrap(dat3, fam.aff_sex, facet.vars = "act", title = "Repeated MeasuresGantt Plot",
  minor.line.freq = 25, major.line.freq = 100)

## End(Not run)
```

gantt_wrap

Gantt Plot

Description

A ggplot2 wrapper that produces a Gantt plot

Usage

```
gantt_wrap(dataframe, plot.var, facet.vars = NULL,
  fill.var = NULL, title = NULL,
  ylab = as.character(plot.var),
  xlab = "duration.default", rev.factor = TRUE,
  transform = FALSE, ncol = NULL, minor.line.freq = NULL,
  major.line.freq = NULL, sig.dig.line.freq = 1,
  hms.scale = NULL, scale = NULL, space = NULL, size = 3,
  rm.horiz.lines = FALSE, x.ticks = TRUE, y.ticks = TRUE,
  legend.position = NULL, bar.color = NULL,
  border.color = NULL, border.size = 2,
  border.width = 0.1, constrain = TRUE)
```

Arguments

<code>dataframe</code>	A data frame with plotting variable(s) and a column of start and end times.
<code>plot.var</code>	A factor plotting variable (y axis)
<code>facet.vars</code>	An optional single vector or list of 1 or 2 to facet by
<code>fill.var</code>	An optional variable to fill the code strips by.
<code>title</code>	An optional title for the plot.
<code>ylab</code>	An optional y label.
<code>xlab</code>	An optional x label.
<code>rev.factor</code>	logical. If TRUE reverse the current plotting order so the first element in the plotting variable's levels is plotted on top.
<code>ncol</code>	if an integer value is passed to this <code>gantt_wrap</code> uses <code>facet_wrap</code> rather than <code>facet_grid</code>
<code>transform</code>	logical. If TRUE the repeated facets will be transformed from stacked to side by side.
<code>minor.line.freq</code>	A numeric value for frequency of minor grid lines.
<code>major.line.freq</code>	A numeric value for frequency of major grid lines.
<code>sig.dig.line.freq</code>	An internal rounding factor for minor and major line freq. Generally, default value of 1 suffices for larger range of x scale may need to be set to -2..
<code>hms.scale</code>	logical. If TRUE converts scale to h:m:s format. Default NULL attempts to detect if object is a <code>cm_time2long</code> object
<code>scale</code>	Should scales be fixed ("fixed", the default), free ("free"), or free in one dimension ("free_x", "free_y")
<code>space</code>	If "fixed", the default, all panels have the same size. If "free_y" their height will be proportional to the length of the y scale; if "free_x" their width will be proportional to the length of the x scale; or if "free" both height and width will vary. This setting has no effect unless the appropriate scales also vary.
<code>size</code>	The width of the plot bars.
<code>rm.horiz.lines</code>	logical. If TRUE the horizontal lines will be removed.
<code>x.ticks</code>	logical. If TRUE the x ticks will be displayed.
<code>y.ticks</code>	logical. If TRUE the y ticks will be displayed.

<code>legend.position</code>	The position of legends. ("left", "right", "bottom", "top", or two-element numeric vector).
<code>bar.color</code>	Optional color to constrain all bars.
<code>border.color</code>	The color to plot border around Gantt bars (default is NULL).
<code>border.size</code>	An integer value for the size to plot borders around Gantt bars. Controls length (width also controlled if not specified).
<code>border.width</code>	Controls border width around Gantt bars. Use a numeric value in addition to border size if plot borders appear disproportional.
<code>constrain</code>	logical. If TRUE the Gantt bars touch the edge of the graph.

Value

Returns a Gantt style visualization. Invisibly returns the ggplot2 list object.

Note

For non repeated measures data/plotting use `gantt`; for repeated measures data output use `gantt_rep`; and for a convenient wrapper that takes text and generates plots use `gantt_plot`.

Author(s)

Andrie de Vries and Tyler Rinker <tyler.rinker@gmail.com>.

References

Clark, W. & Gantt, H. (1922) The Gantt chart, a working tool of management. New York, Ronald Press.

See Also

[gantt](#), [gantt_plot](#), [gantt_rep](#), [facet_grid](#), [facet_wrap](#)

Examples

```
## Not run:
(dat <- gantt(mraja1$dialogue, list(mraja1$fam.aff, mraja1$sex),
  units = "sentences", plot.colors = 'black', sums = TRUE,
  col.sep = "_")$gantt.df)
gantt_wrap(dat, fam.aff_sex, title = "Gantt Plot")
dat$codes <- sample(LETTERS[1:3], nrow(dat), TRUE)
gantt_wrap(dat, fam.aff_sex, fill.var = "codes", legend.position = "bottom")

(dat3 <- with(rajSPLIT, gantt_rep(act, dialogue, list(fam.aff, sex),
  units = "words", col.sep = "_"))
x <- gantt_wrap(dat3, fam.aff_sex, facet.vars = "act",
  title = "Repeated Measures Gantt Plot")
x + scale_color_manual(values=rep("black", length(levels(dat3$fam.aff_sex))))

## End(Not run)
```

hash

Hash/Dictionary Lookup

Description

Creates a new environemnt for quick hash style dictionary lookup.

Usage

```
hash(x)
```

Arguments

x A two column dataframe

Value

Creates a "hash table" or a two column data frame in its own environment.

Author(s)

Bryan Goodrich and Tyler Rinker <tyler.rinker@gmail.com>.

References

<http://www.talkstats.com/showthread.php/22754-Create-a-fast-dictionary>

See Also

[lookup](#), [environment](#)

Examples

```
## Not run:
new.hash <- hash(aggregate(conc~Plant, C02, sum))
new.hash <- hash(aggregate(conc~Plant, C02, sum))
sapply(as.character(C02$Plant), function(x) {
  if(exists(x, env = new.hash)) {
    get(x, e = new.hash)
  } else {
    NA
  }
})

## End(Not run)
```

htruncdf	<i>Dataframe Viewing</i>
----------	--------------------------

Description

htruncdf - Convenience function to view the head of a truncated dataframe.

truncdf - Convenience function to view a truncated dataframe.

qview - Convenience function to view a summary and head of a dataframe.

Usage

```
htruncdf(dataframe, n = 10, width = 10)
```

```
truncdf(dataframe, end = 10, begin = 1)
```

```
qview(dataframe, ...)
```

Arguments

dataframe	A data.frame object.
n	Number of rows to display.
width	The width of the columns to be displayed.
end	The last element to be displayed.
begin	The first element to be displayed.
...	Other arguments passed to head .

Value

htruncdf - returns n number of rows of a truncated dataframe.

truncdf - returns a truncated dataframe.

qview - returns a dataframe head with summary statistics.

See Also

[head](#)

Examples

```
## Not run:  
htruncdf(raj)  
htruncdf(raj, 20)  
htruncdf(raj, ,20)  
truncdf(raj)  
truncdf(raj, 40)  
qview(raj)  
qview(CO2)  
  
## End(Not run)
```

imperative

Intuitively Remark Sentences as Imperative

Description

Automatic imperative remarking.

Usage

```
imperative(dataframe, person.var, text.var,
  lock.incomplete = FALSE, additional.names = NULL,
  warning = FALSE)
```

Arguments

dataframe	A data.frame object.
person.var	The person variable.
text.var	The text variable.
lock.incomplete	logical. If TRUE locks incomplete sentences (sentences ending with " ") from being marked as imperative.
additional.names	Additional names that may be used in a command (people in the context that do not speak).
warning	logical. If TRUE provides comma warnings (sentences that contain numerous commas that may be handled incorrectly by the algorithm).

Value

Returns a dataframe with a text variable indicating imperative sentences. Imperative sentences are marked with * followed by the original end mark.

Note

The algorithm used by imperative is sentive to English language dialects and types. Commas can indicate a choppy sentence and may indicate a false postive.

Examples

```
## Not run:
DATA3 <- data.frame(name=c('sue', rep(c('greg', 'tyler', 'phil', 'sue'), 2)),
  statement=c('go get it|', 'I hate to read.', 'Stop running!', 'I like it!',
    'You are terrible!', "Don't!", 'Greg, go to the red, brick office.',
    'Tyler go to the gym.', "Alex don't run."), stringsAsFactors = FALSE)
imperative(DATA3, 'name', 'statement', , c('Alex'))
imperative(DATA3, 'name', 'statement', lock.incomplete = TRUE, c('Alex'))
imperative(DATA3, 'name', 'statement', , c('Alex'), warning=TRUE)
X <- imperative(mrajalspl, 'person', 'dialogue', warning=FALSE)
truncdf(X[, -7], 60)
strwrap(X$dialogue)

## End(Not run)
```

incomplete.replace	<i>Denote Incomplete End Marks With " "</i>
--------------------	---

Description

Replaces incomplete sentence end marks (., ..., .?, ..?, en \& em dash etc.) with "|".

Usage

```
incomplete.replace(text.var, scan.mode = FALSE)
```

```
incomp(text.var, scan.mode = FALSE)
```

Arguments

text.var The text variable.

scan.mode logical. If TRUE only scans and reports incomplete sentences.

Value

Returns a text variable (character sting) with incomplete sentence marks (., ..., .?, ..?, en \& em dash etc. replaced with "|". If scan mode is TRUE returns a data frame with incomplete sentence location.

Examples

```
## Not run:
x <- c("the...", "I.?", "you.", "threw..", "we?")
incomplete.replace(x)
incomp(x)
incomp(x, TRUE)

## End(Not run)
```

increase.amplification.words	<i>Amplifying Words</i>
------------------------------	-------------------------

Description

A dataset containing a vector of words that amplify word meaning.

Format

A vector with 32 elements

Details

Valence shifters are words that alter or intensify the meaning of the polarized words and include negators and amplifiers. Negators are, generally, adverbs that negate sentence meaning; for example the word like in the sentence, "I do like pie.", is given the opposite meaning in the sentence, "I do not like pie.", now containing the negator not. Amplifiers are, generally, adverbs or adjectives that intensify sentence meaning. Using our previous example, the sentiment of the negator altered sentence, "I seriously do not like pie.", is heightened with addition of the amplifier seriously.

interjections	<i>Interjections</i>
---------------	----------------------

Description

A dataset containing a character vector of common interjections.

Format

A character vector with 139 elements

References

<http://www.vidarholen.net/contents/interjections/>

key_merge	<i>Merge Demographic Information with Person/Text Transcript</i>
-----------	--

Description

Wrapper function ([merge](#)) for merging demographic information with a person/text transcript.

Usage

```
key_merge(transcript.df, key.df, common.column = NULL,
          default.arrange = TRUE)
```

Arguments

transcript.df	The text/person transcript dataframe
key.df	The demographic dataframe.
common.column	The column(s) shared by transcript.df and key.df. If NULL function defaults to use any columns with the same name.
default.arrange	logical. If TRUE will arrange the columns with text to the far right.

Value

Outputs a merged transcript dataframe with demographic information.

See Also[merge](#)**Examples**

```
## Not run:
#First view transcript dataframe and demographics dataframe.
lapply(list(raj, raj.demographics), head)
merged.raj <- key_merge(raj, raj.demographics)
htruncdf(merged.raj, 10, 40)

## End(Not run)
```

kullback.leibler	<i>Kullback Leibler Statistic</i>
------------------	-----------------------------------

Description

A proximate measure between two probability distributions applied to speech.

Usage

```
kullback.leibler(x, y = NULL, digits = 3)
```

Arguments

<code>x</code>	A numeric vector, matrix or data frame.
<code>y</code>	A second numeric vector if <code>x</code> is also a vector. Default is <code>NULL</code> .
<code>digits</code>	Number of decimal places to round.

Details

Uses Kullback & Leibler's (1951) formula:

$$D_{KL}(P||Q) = \sum_i \ln \left(\frac{P_i}{Q_i} \right) P_i$$

Value

Returns a matrix of the Kullback Leibler measure between each vector of probabilities.

Note

The `kullback.leibler` function generally receives the output of either `wfm` or `wfdf` functions.

References

Kullback, S., & Leibler, R.A. (1951). On Information and sufficiency. *Annals of Mathematical Statistics* 22 (1): 79-86. doi:10.1214/aoms/1177729694

Examples

```
## Not run:
p.df <- wfdf(DATA$state, DATA$person)
p.mat <- wfm(text.var = DATA$state, grouping.var = DATA$person)

kullback.leibler(p.mat)
kullback.leibler(p.df)
kullback.leibler(p.df$greg, p.df$sam)

p.df2 <- wfdf(raj$dialogue, raj$person)
kullback.leibler(p.df2)

## End(Not run)
```

left.just

*Text Justification***Description**

left.just - Left justifies a text/character column.

right.just - A means of undoing a left justification.

Usage

```
left.just(dataframe, column = NULL, keep.class = FALSE)
```

```
right.just(dataframe)
```

Arguments

dataframe	A data.frame object with the text column.
column	The column to be justified. If NULL all columns are justified.
keep.class	logical. If TRUE will attempt to keep the original classes of the dataframe if the justification is not altered (i.e. numeric will not be honored but factor may be).

Value

Returns a dataframe with selected text column left/right justified.

Note

left.just inserts spaces to achieve the justification. This could interfere with analysis and therefore the output from left.just should only be used for visualization purposes, not analysis.

Examples

```
## Not run:
left.just(DATA)
left.just(DATA, "state")
left.just(CO2)
right.just(left.just(CO2))

## End(Not run)
```

lookup	<i>Hash Table/Dictionary Lookup</i>
--------	-------------------------------------

Description

Environment based hash table useful for large vector lookups.

Usage

```
lookup(terms, key.match, key.reassign = NULL,  
       missing = NA)
```

Arguments

terms	A vector of terms to undergo a lookup.
key.match	Either a two column data frame (if data frame supplied no key reassign needed) of a match key and reassignment column or a single vector match key.
key.reassign	A single reassignment vector supplied if key.match is not a two column data frame.
missing	Value to assign to terms not matching the key.match.

Value

Outputs A new vector with reassigned values.

See Also

[new.env](#)

Examples

```
## Not run:  
lookup(mtcars$carb, sort(unique(mtcars$carb)),  
       c('one', 'two', 'three', 'four', 'six', 'eight'))  
lookup(mtcars$carb, sort(unique(mtcars$carb)),  
       seq(10, 60, by=10))  
  
key <- data.frame(x=1:2, y=c("A", "B"))  
big.vec <- sample(1:2, 3000000, T)  
lookup(big.vec, key)  
  
lookup(1:5, data.frame(1:4, 11:14))  
lookup(LETTERS[1:5], data.frame(LETTERS[1:5], 100:104))  
  
## End(Not run)
```

mcsv_r

*Read/Write Multiple csv Files at a Time***Description**

mcsv_w - Read and assign multiple csv files at the same time.

mcsv_w - Write multiple csv files into a file at the same time.

Usage

```
mcsv_r(files, a.names = NULL, l.name = NULL, list = TRUE)
```

```
mcsv_w(..., dir = NULL, open = FALSE)
```

Arguments

files	csv file(s) to read.
a.names	object names to assign the csv file(s) to. If NULL assigns the csv to the name(s) of the csv file(s) in the global environment.
l.name	A character vector of names to assign to the csv files (dataframes) being read in. Default (NULL) uses the names of the files in the directory without the file extension.
list	A character vector of length one to name the list being read in. Default is "L1".
...	data.frame object(s) to write to a file
dir	optional directory names. If NULL a directory will be created in the working directory with the data and time stamp as the folder name.
open	logical. If TRUE opens the directory upon completion.

Details

mcsv is short for "multiple csv" and the suffix c(_r, _w) stands for "read" (r) or "write" (w).

Value

mcsv_r - reads in multiple csv files at once.

mcsv_w - creates a directory with multiple csv files. Silently returns the path of the directory.

Note

mcsv_r is useful for reading in multiple csv files from `cm_csv$temp` for interaction with `cm_range2long`.

See Also

[cm_range2long](#), [cm_df\\$temp](#)

Examples

```
## Not run:
#mcsv_r EXAMPLE:
mtcarsb <- mtcars; C02b <- C02
a <- mcsv_w(mtcarsb, C02b, dir="foo")
a
rm("mtcarsb", "C02b") # gone from .GlobalEnv
(nms <- dir(a))
mcsv_r(paste(a, nms, sep="/"))
mtcarsb; C02b
rm("mtcarsb", "C02b") # gone from .GlobalEnv
mcsv_r(paste(a, nms, sep="/"), paste0("foo.dat", 1:2))
foo.dat1; foo.dat2
rm("foo.dat1", "foo.dat2") # gone from .GlobalEnv
delete("foo")

#mcsv_w EXAMPLE:
a <- mcsv_w(mtcars, C02, dir="foo")
a
delete("foo")

## End(Not run)
```

merge.all

*Merge Multiple Data Sets***Description**

Merge multiple data sets together.

Usage

```
merge.all(frames, by, na.replace = NA)
```

Arguments

frames	Multiple dataframes to merge together.
by	Specifications of the common column(s).
na.replace	Value to replace missing values with.

Value

Returns a dataframe with multiple dataframes merged together.

References

<http://stackoverflow.com/questions/9551555/combine-a-series-of-data-frames-and-create-new-column>

See Also

[merge](#)

Examples

```
## Not run:
#Create three dataframe
Week_1_sheet <- read.table(text="ID Gender  DOB Absences Unexcused_Absences Lates
1 1      M 1997      5              1    14
2 2      F 1998      4              2     3", header=TRUE)

Week_2_sheet <- read.table(text="ID Gender  DOB Absences Unexcused_Absences Lates
1 1      M 1997      2              1    10
2 2      F 1998      8              2     2
3 3      M 1998      8              2     2", header=TRUE)

Week_3_sheet <- read.table(text="ID Gender  DOB Absences Unexcused_Absences Lates
1 1      M 1997      2              1    10
2 2      F 1998      8              2     2", header=TRUE)

#Consolidate them into a list
WEEKlist <- list(Week_1_sheet , Week_2_sheet , Week_3_sheet)

names(WEEKlist) <- LETTERS[1:3]

#change names of columns that may overlap with other data frame yet not have
#duplicate data
lapply(seq_along(WEEKlist), function(x) {
  y <- names(WEEKlist[[x]]) #do this to avoid repeating this 3 times
  names(WEEKlist[[x]]) <- c(y[1:3], paste(y[4:length(y)], ".", x, sep=""))
}) #notice the assignment to the enviroment

merge.all(frames=WEEKlist, by=c('ID', 'Gender', 'DOB'))
merge.all(frames=WEEKlist, by=1:3, na.replace = 0)

## End(Not run)
```

mrja1

Romeo and Juliet: Act 1 Dialogue Merged with Demographics

Description

A dataset containing act 1 of Romeo and Juliet with demographic information.

Format

A data frame with 235 rows and 5 variables

Details

- person. Character in the play
- sex. Gender
- fam.aff. Family affiliation of character
- died. Dummy coded death variable (0-no; 1-yes); if yes the character dies in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

mrja1spl	<i>Romeo and Juliet: Act 1 Dialogue Merged with Demographics and Split</i>
----------	--

Description

A dataset containing act 1 of Romeo and Juliet with demographic information and turns of talk split into sentences.

Format

A data frame with 508 rows and 7 variables

Details

- person. Character in the play
- tot.
- sex. Gender
- fam.aff. Family affiliation of character
- died. Dummy coded death variable (0-no; 1-yes); if yes the character dies in the play
- dialogue. The spoken dialogue
- stem.text.

References

http://shakespeare.mit.edu/romeo_juliet/full.html

multigsub	<i>Multiple gsub</i>
-----------	----------------------

Description

A wrapper for [gsub](#) that takes a vector of search terms and a vector or single value of replacements.

Usage

```
multigsub(pattern, replacement = NULL, text.var,
  leadspace = FALSE, trailspace = FALSE, fixed = TRUE,
  ...)

mgsub(pattern, replacement = NULL, text.var,
  leadspace = FALSE, trailspace = FALSE, fixed = TRUE,
  ...)
```

Arguments

pattern	Character string to be matched in the given character vector.
replacement	Character string equal in length to pattern or of length one which are a replacement for matched pattern.
text.var	The text variable.
leadspace	logical. If TRUE inserts a leading space in the repalcements.
trailspace	logical. If TRUE inserts a trailing space in the repalcements.
fixed	logical. If TRUE, pattern is a string to be matched as is. Overrides all conflicting arguments.
...	Additional arguments passed to gsub .

Value

Returns a vector with the pattern replaced.

Note

The replacements occur sequentially rather than all at once. This means a previous (first in pattern string) sub could alter a later sub.

See Also

[gsub](#)

Examples

```
## Not run:
multisub(c("it's", "I'm"), c("it is", "I am"), DATA$state)
mgsub(c("it's", "I'm"), c("it is", "I am"), DATA$state)
mgsub("[:punct:]", "PUNC", DATA$state, fixed = FALSE)

## End(Not run)
```

multiscale	<i>Nested Standardization</i>
------------	-------------------------------

Description

Standardize within a subgroup and then within a group.

Usage

```
multiscale(numeric.var, grouping.var,
  original_order = TRUE, digits = 2)
```

Arguments

numeric.var	A numeric variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
original_order	logical. IF TRUE orders by the original order. If FALSE orders by group.
digits	Integer; number of decimal places to round.

Value

Returns a list of two:

SCALED_OBSERVATIONS

A dataframe of scaled observations at level one and two of the nesting with possible outliers.

DESCRIPTIVES_BY_GROUP

A data frame of descriptives by group.

See Also

[scale](#)

Examples

```
## Not run:
dat <- with(mraja1spl, word_stats(dialogue, list(person, sex, fam.aff)))
head(colsplit2df(dat$ts))
with(colsplit2df(dat$ts), multiscale(word.count, person))
with(colsplit2df(dat$ts), multiscale(word.count, list(fam.aff, sex)))
with(colsplit2df(dat$ts), multiscale(word.count, list(fam.aff, sex),
  original_order = FALSE))

## End(Not run)
```

NAer

Replace Missing Values (NA)

Description

Replace missing values (NA) in a vector or dataframe.

Usage

```
NAer(x, replace = 0)
```

Arguments

x A vector or dataframe with missing values (NA).

replace The value to replace missing values (NA) with.

Value

Returns a vector or dataframe with missing values replaced.

Examples

```
## Not run:
set.seed(10)
x <- sample(c(rep(NA, 4), 1:10), 20, rep=T)
y <- data.frame(matrix(x, 5, 4))
names(y) <- paste('var', 1:4, sep="_")

NAer(x)
NAer(y)
NAer(y, "MISSING")

## End(Not run)
```

negation.words

*Negating Words***Description**

A dataset containing a vector of words that negate word meaning.

Format

A vector with 16 elements

Details

Valence shifters are words that alter or intensify the meaning of the polarized words and include negators and amplifiers. Negators are, generally, adverbs that negate sentence meaning; for example the word like in the sentence, "I do like pie.", is given the opposite meaning in the sentence, "I do not like pie.", now containing the negator not. Amplifiers are, generally, adverbs or adjectives that intensify sentence meaning. Using our previous example, the sentiment of the negator altered sentence, "I seriously do not like pie.", is heightened with addition of the amplifier seriously.

negative.words

*Negative Words***Description**

A dataset containing a vector of negative words.

Format

A vector with 4783 elements

Details

A sentence containing more negative words would be deemed a negative sentence, whereas a sentence containing more positive words would be considered positive.

References

Hu, M., & Liu, B. (2004). Mining opinion features in customer reviews. National Conference on Artificial Intelligence.

<http://www.cs.uic.edu/~liub/FBS/sentiment-analysis.html>

OnixTxtRetToolkitSWL1 *Onix Text Retrieval Toolkit Stopword List 1*

Description

A stopwords list containing a character vector of stopwords.

Format

A character vector with 404 elements

Details

From Onix Text Retrieval Toolkit API Reference: "This stopwords list is probably the most widely used stopwords list. It covers a wide number of stopwords without getting too aggressive and including too many words which a user might search upon."

Note

Reduced from the original 429 words to 404.

References

<http://www.lextek.com/manuals/onix/stopwords1.html>

outlier.detect *Detect Outliers in Text*

Description

Locate possible outliers for text variables given numeric word function.

Usage

```
outlier.detect(text.var, grouping.var = NULL,
              FUN = word.count, scale.by = "grouping")
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
FUN	A word function with a numeric vector output (e.g. syllable.sum, character.count or word.count).
scale.by	A character string indicating which dimensions to scale by. One of "all", "grouping", or "both". Default NULL scales by all.

Value

Returns a dataframe with possible outliers.

Examples

```
## Not run:
with(DATA, outlier.detect(state))
with(DATA, outlier.detect(state, FUN = character.count))
with(DATA, outlier.detect(state, person, FUN = character.count))
with(DATA, outlier.detect(state, list(sex, adult), FUN = character.count))
with(DATA, outlier.detect(state, FUN = syllable.sum))
htruncdf(with(raj, outlier.detect(dialogue, person)), 15, 45)

## End(Not run)
```

outlier.labeler

Locate Outliers in Numeric String

Description

Locate and label possible outliers in a string.

Usage

```
outlier.labeler(x, standardize = TRUE, ...)
```

Arguments

x	A numeric vector.
standardize	logical. If TRUE scales the vector first.
...	Other arguments passed to scale .

Value

Returns a matrix (one column) of possible outliers coded as "3sd", "2sd" and "1.5sd", corresponding to \geq to 3, 2, or 1.5 standard deviations.

See Also

[scale](#)

Examples

```
## Not run:
outlier.labeler(mtcars$hp)
by(mtcars$mpg, mtcars$cyl, outlier.labeler)
tapply(mtcars$mpg, mtcars$cyl, outlier.labeler)

## End(Not run)
```

paste2*Paste an Unspecified Number Of Text Columns*

Description

Paste unspecified columns or a list of vectors together.

Usage

```
paste2(multi.columns, sep = ".", handle.na = TRUE,  
       trim = TRUE)
```

Arguments

<code>multi.columns</code>	The multiple columns or a list of vectors to paste together.
<code>sep</code>	A character string to separate the terms.
<code>handle.na</code>	logical. If TRUE returns NA if any column/vector contains a missing value.
<code>trim</code>	logical. If TRUE leading/trailing white space is removed.

Value

Returns a vector with row-wise elements pasted together.

Note

[paste](#) differs from [paste2](#) because [paste](#) does not allow an unspecified number of columns to be pasted. This behavior can be convenient for inside of functions when the number of columns being pasted is unknown.

See Also

[paste](#)

Examples

```
## Not run:  
v <- rep(list(state.abb, state.name), 5)  
n <- sample(5:10, 1)  
paste(v[1:n]) #odd looking return  
paste2(v[1:n])  
paste2(v[1:n], sep="|")  
paste2(mtcars, sep="|")  
paste(mtcars, sep="|") #odd looking return  
paste2(CO2, sep="|-|")  
  
## End(Not run)
```

polarity.score *Polarity Score (Sentiment Analysis)*

Description

Aproximate the sentiment (polarity) of text by grouping variable(s).

Usage

```
polarity.score(text.var, grouping.var = NULL,
               positive.list = positive.words,
               negative.list = negative.words,
               negation.list = negation.words,
               amplification.list = increase.amplification.words,
               rm.incomplete = FALSE, digits = 3, ...)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
positive.list	A character vector of terms indicating positive reaction.
negative.list	A character vector of terms indicating negative reaction.
negation.list	A character vector of terms reversing the intent of a positive or negative word.
amplification.list	A character vector of terms that increases the intensity of a psoitive or negativbe word.
digits	Integer; number of decimal places to round.
...	Other arguments supplied to endf.

Details

The equation used by the algorithm to assign value to polarity to each sentence fist utilizes the sentiment dictionary (Hu and Liu, 2004) to tag each word as either positive (x_i^+), negative (x_i^-), neutral (x_i^0), negator(x_i^-), or amplifier (x_i^\uparrow). Neutral words hold no value in the equation but do affect word count (n). Each positive (x_i^+) and negative (x_i^-) word is then weighted by the amplifiers (x_i^\uparrow) directly proceeding the positive or negative word. Next, I consider amplification value, adding the assigned value $1/n - 1$ to increase the polarity relative to sentence length while ensuring that the polarity scores will remain between the values -1 and 1. This weighted value for each polarized word is then multiplied by -1 to the power of the number of negated (x_i^-) words directly proceeding the positive or negative word. Last, these values are then summed and divided by the word count (n) yielding a polarity score (δ) between -1 and 1.

$$\delta = \frac{\sum(x_i^0, \quad x_i^\uparrow + x_i^+ \cdot (-1)^{\sum(x_i^-)}, \quad x_i^\uparrow + x_i^- \cdot (-1)^{\sum(x_i^-)})}{n}$$

Where:

$$x_i^\uparrow = \frac{1}{n - 1}$$

Value

Returns a list of two dataframes:

all	<p>A dataframe of scores per row with:</p> <ul style="list-style-type: none"> • wc - word count • polarity - sentence polarity score • raw - raw polarity score (considering only positive and nagtive words) • negation.adj.raw - raw adjusted for negation words • amplification.adj.raw - raw adjusted for amplification words • pos.words - words considered positive • neg.words - words considered negative
group	A dataframe with the average polarity score by grouping variable.

Note

The polarity score is dependant upon the polarity dictionary used. This function defaults to the word polarity word dictionary used by Hu, M., & Liu, B. (2004), however, this may not be appropriate for the context of children in a classroom. The user may (is encouraged) to provide/augment the dictionary. For instance the word "sick" in a high school setting may mean that something is good, whereas "sick" used by a typical adult indicates something is not right or negative connotation.

Also note that `polarity.score` assumes you've run `sentSplit`.

References

Hu, M., & Liu, B. (2004). Mining opinion features in customer reviews. National Conference on Artificial Intellgence.

<http://www.slideshare.net/jeffreybreen/r-by-example-mining-twitter-for>

See Also

<https://github.com/trestletech/Sermon-Sentiment-Analysis>

Examples

```
## Not run:
(poldat <- with(DATA, polarity.score(state, person)))
with(DATA, polarity.score(state, list(sex, adult)))
names(poldat)
poldat$all
poldat$group
poldat2 <- with(mraja1spl, polarity.score(dialogue, list(sex, fam.aff, died)))
colsplit2df(poldat2$group)

## End(Not run)
```

pos *Parts of Speech Tagging*

Description

pos - Apply part of speech tagger to transcript(s).

pos.by - Apply part of speech tagger to transcript(s) by zero or more grouping variable(s).

pos.tags - Useful for interpreting the parts of speech tags created by pos and pos.by.

Usage

```
pos(text.var, parallel = FALSE, na.omit = FALSE,
    digits = 2, progress.bar = TRUE, gc.rate = 10)
```

```
pos.by(text.var, grouping.var = NULL, digits = 2, ...)
```

```
pos.tags(type = "pretty")
```

Arguments

text.var	The text variable
parallel	logical. If TRUE attempts to run the function on multiple cores. Note that this may not mean a speed boost if you have one core or if the data set is smaller as the cluster takes time to create.
na.omit	logical. If TRUE missing values (JcodeNA) will be omitted.
digits	integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed
progress.bar	logical. If TRUE attempts to provide a OS appropriate progress bar. If parallel is TRUE this argument is ignored. Note that setting this argument to TRUE may slow down the function.
gc.rate	An integer value. This is a necessary argument because of a problem with the garbage collection in the openNLP function that pos wraps. Consider adjusting this argument upward if the error <code>java.lang.OutOfMemoryError</code> occurs.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
...	Other argument supplied to pos.
type	An optional character string giving the output of the pos tags. This must be one of the strings "pretty" (a left justified version of the output optimized for viewing but not good for export), "matrix" (a matrix version of the output), "dataframe" or "df" (a dataframe version of the output), "all" (a list of all three of the previous output types).

Value

pos returns a list of 4:

text	The original text
POSTagged	The original words replaced with parts of speech in context.

POSprop Dataframe of the proportion of parts of speech by row.
 POSfreq Dataframe of the frequency of parts of speech by row.

pos.by returns a list of 6:

text The original text
 POStagged The original words replaced with parts of speech in context.
 POSprop Dataframe of the proportion of parts of speech by row.
 POSfreq Dataframe of the frequency of parts of speech by row.
 pos.by.prop Dataframe of the proportion of parts of speech by grouping variable.
 pos.by.freq Dataframe of the frequency of parts of speech by grouping variable.

References

openNLP <http://opennlp.apache.org>

See Also

[tagPOS](#)

Examples

```
## Not run:
posdat <- pos(DATA$state)
str(posdat)
names(posdat)
posdat$text        #original text
posdat$POSTagged   #words replaced with parts of speech
posdat$POSprop     #proportion of parts of speech by row
posdat$POSfreq     #frequency of parts of speech by row

pos(DATA$state, parallel = TRUE) # not always useful

#use pos.tags to interpret part of speech tags used by pos & pos.by
pos.tags()
pos.tags("matrix")
pos.tags("dataframe")
pos.tags("df")
pos.tags("all")

posbydat <- with(DATA, pos.by(state, sex))
names(posbydat)
posbydat
posbydat$pos.by.prop
with(DATA, pos.by(state, list(adult, sex)))
#or more quickly - reuse the output from before
with(DATA, pos.by(posbydat, list(adult, sex)))

## End(Not run)
```

positive.words	<i>Positive Words</i>
----------------	-----------------------

Description

A dataset containing a vector of positive words.

Format

A vector with 2006 elements

Details

A sentence containing more negative words would be deemed a negative sentence, whereas a sentence containing more positive words would be considered positive.

References

Hu, M., & Liu, B. (2004). Mining opinion features in customer reviews. National Conference on Artificial Intelligence.

<http://www.cs.uic.edu/~liub/FBS/sentiment-analysis.html>

potential_NA	<i>Search for Potential Missing Values</i>
--------------	--

Description

Search for potential missing values (i.e. sentences that are merely a punctuation mark) and optionally replace with missing value (NA). Useful in the initial cleaning process.

Usage

```
potential_NA(text.var, n = 3)
```

Arguments

text.var	The text variable.
n	Number of characters to consider for missing (default is 3).

Value

Returns a dataframe of potential missing values row numbers and text.

Examples

```
## Not run:
DATA$state[c(3, 7)] <- "."
potential_NA(DATA$state, 20)
potential_NA(DATA$state)
# USE TO SELECTIVELY REPLACE CELLS WITH MISSING VALUES
DATA$state[potential_NA(DATA$state, 20)$row[-c(3)]] <- NA
DATA
DATA <- qdap::DATA

## End(Not run)
```

preposition

*Preposition Words***Description**

A dataset containing a vector of common prepositions.

Format

A vector with 162 elements

print.adjacency.matrix

*Prints an adjacency.matrix***Description**

Prints an adjacency.matrix.

Usage

```
## S3 method for class 'adjacency.matrix'
print(x, ...)
```

Arguments

x	The adjacency.matrix object
...	ignored

print.cm.distance	<i>Prints a cm.distance object</i>
-------------------	------------------------------------

Description

Prints a cm.distance object

Usage

```
## S3 method for class 'cm.distance'  
print(x, ...)
```

Arguments

x	The cm.distance object
...	ignored

print.formality.measure	<i>Prints a formality.measure object</i>
-------------------------	--

Description

Prints a formality.measure object

Usage

```
## S3 method for class 'formality.measure'  
print(x, ...)
```

Arguments

x	The formality.measure object
...	ignored

print.polarity.score	<i>Prints a polarity.score object</i>
----------------------	---------------------------------------

Description

Prints a polarity.score object.

Usage

```
## S3 method for class 'polarity.score'  
print(x, ...)
```

Arguments

x	The polarity.score object
...	ignored

print.POS	<i>Prints a POS object</i>
-----------	----------------------------

Description

Prints a POS object.

Usage

```
## S3 method for class 'POS'  
print(x, ...)
```

Arguments

x	The POS object
...	ignored

print.POSby	<i>Prints a POSby object</i>
-------------	------------------------------

Description

Prints a POSby object.

Usage

```
## S3 method for class 'POSby'  
print(x, ...)
```

Arguments

x	The POSby object
...	ignored

print.termco_c	<i>Prints an termco_c object.</i>
----------------	-----------------------------------

Description

Prints an termco_c object.

Usage

```
## S3 method for class 'termco_c'  
print(x, ...)
```

Arguments

x	The termco_c object
...	ignored

print.termco_d	<i>Prints a termco_d object.</i>
----------------	----------------------------------

Description

Prints a termco_d object.

Usage

```
## S3 method for class 'termco_d'  
print(x, ...)
```

Arguments

x	The termco_d object
...	ignored

print.word.list	<i>Prints a word.list object</i>
-----------------	----------------------------------

Description

Prints a word.list object.

Usage

```
## S3 method for class 'word.list'  
print(x, ...)
```

Arguments

x	The word.list object
...	ignored

print.word.stats	<i>Prints a word.stats object</i>
------------------	-----------------------------------

Description

Prints a word.stats object.

Usage

```
## S3 method for class 'word.stats'  
print(x, ...)
```

Arguments

x	The word.stats object
...	ignored

```
print.word_associate
```

Prints a word_associate object

Description

Prints a word_associate object.

Usage

```
## S3 method for class 'word_associate'
print(x, ...)
```

Arguments

x	The word_associate object
...	ignored

prop	<i>Convert Raw Numeric Matrix or Data Frame to Proportions</i>
------	--

Description

Convert a raw matrix or dataframe to proportions/percents. Divides each element of a column by the column sum.

Usage

```
prop(mat, digits = 2, percent = FALSE)
```

Arguments

mat	A numeric matrix or dataframe.
digits	Integer; number of decimal places to round.
percent	logical. If TRUE output given as percent. If FALSE the output is proportion.

Value

Returns a matrix with proportionally scaled values.

Examples

```
## Not run:
y <- wfdm(DATA$state, DATA$person, stopwords = c("your", "yours"), margins = TRUE)
prop(wfm(wfdm = y), 4)      #as a proportion
prop(wfm(wfdm = y), 4, TRUE) #as a percentage
heatmap(prop(wfm(wfdm = y), 4))
wdstraj <- word_stats(rajSPLIT$dialogue, rajSPLIT$person)
prop(wdstraj$gts[, -1], 5)

## End(Not run)
```

qcv

*Quick Character Vector***Description**

Create a character vector without the use of quotation marks.

Usage

```
qcv(..., terms = NULL, space.wrap = FALSE,
    trailing = FALSE, leading = FALSE, split = " ",
    rm.blank = TRUE)
```

Arguments

...	Character objects. Either ... or terms argument must be utilized.
terms	An optional argument to present the terms as one long character string. This is useful if the split (separator) is not a comma (e.g. spaces are the term separators).
space.wrap	logical. If TRUE wraps the vector of terms with a leading/trailing space.
trailing	logical. If TRUE wraps the vector of terms with a trailing space.
leading	logical. If TRUE wraps the vector of terms with a leading space.
split	Character vector of length one to use for splitting (i.e. the separator used in the vector). For use with the argument terms.
rm.blank	logical. If TRUE removes all blank spaces from the vector.

Value

Returns a character vector.

See Also

[C](#)

Examples

```
## Not run:
qcv(I, like, dogs)
qcv(terms = "I, like, dogs") #default separator is " "
qcv(terms = "I, like, dogs", split = ",")
qcv(terms = "I like dogs")
qcv(I, like, dogs, space.wrap = TRUE)
qcv(I, like, dogs, trailing = TRUE)
qcv(I, like, dogs, leading = TRUE)
exclude(Top25Words, qcv(the, of, and))
qcv(terms = "mpg cyl disp hp drat wt qsec vs am gear carb")

## End(Not run)
```

qdap

*qdap: Quantitative Discourse Analysis Package***Description**

This package automates many of the tasks associated with quantitative discourse analysis of transcripts containing discourse. The package provides parsing tools for preparing transcript data, coding tools and analysis tools for richer understanding of the data. Many functions allow the user to aggregate data by any number of grouping variables, providing analysis and seamless integration with other R packages which enable higher level analysis and visualization of text. This empowers the researcher with more flexible, efficient and targeted methods and tools.

qprep

*Quick Preparation of Text***Description**

Wrapper for `bracketX`, `replace_number`, `replace_symbol`, `replace_abbreviation` and `scrubber` to quickly prepare text for analysis. Care should be taken with this function to ensure data is properly formatted and complete.

Usage

```
qprep(text.var, rm.dash = TRUE, bracket = "all",
      missing = NULL, names = FALSE,
      abbreviation = qdap::abbreviations, replace = NULL,
      ignore.case = TRUE, num.paste = "separate")
```

Arguments

<code>text.var</code>	The text variable.
<code>rm.dash</code>	logical. If TRUE dashes will be removed.
<code>bracket</code>	The type of bracket (and encased text) to remove. This is one of the strings "curly", "square", "round", "angle" and "all". These strings correspond to: {, [, (, < or all four types.
<code>missing</code>	Value to assign to empty cells.
<code>missing</code>	Value to assign to empty cells.
<code>names</code>	logical. If TRUE the sentences are given as the names of the counts.
<code>abbreviation</code>	A two column key of abbreviations (column 1) and long form replacements (column 2) or a vector of abbreviations. Default is to use qdap's abbreviations data set.
<code>replace</code>	A vector of long form replacements if a data frame is not supplied to the abbreviation argument.
<code>ignore.case</code>	logical. If TRUE replaces without regard to capitalization.
<code>num.paste</code>	A character string c("separate", "combine"); "separate" will treat each word section as separate, "combine" will lump the sections together as one word.

Note

Care should taken with this function to ensure data is properly formatted and complete.

See Also

[bracketX](#), [replace_abbreviation](#), [replace_number](#), [replace_symbol](#)

Examples

```
## Not run:
x <- "I like 60 (laughter) #d-bot and $6 @ the store w/o 8p.m."
qprep(x)

## End(Not run)
```

raj

Romeo and Juliet (Unchanged & Complete)

Description

A dataset containing the original transcript from Romeo and Juliet as it was scraped from: http://shakespeare.mit.edu/romeo_juliet/full.html.

Format

A data frame with 840 rows and 3 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue
- act. The act (akin to repeated measures)

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.act.1

Romeo and Juliet: Act 1

Description

A dataset containing Romeo and Juliet: Act 1.

Format

A data frame with 235 rows and 2 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.act.2

Romeo and Juliet: Act 2

Description

A dataset containing Romeo and Juliet: Act 2.

Format

A data frame with 205 rows and 2 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.act.3

Romeo and Juliet: Act 3

Description

A dataset containing Romeo and Juliet: Act 3.

Format

A data frame with 197 rows and 2 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.act.4

Romeo and Juliet: Act 4

Description

A dataset containing Romeo and Juliet: Act 4.

Format

A data frame with 115 rows and 2 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.act.5

Romeo and Juliet: Act 5

Description

A dataset containing Romeo and Juliet: Act 5.

Format

A data frame with 88 rows and 2 variables

Details

- person. Character in the play
- dialogue. The spoken dialogue

References

http://shakespeare.mit.edu/romeo_juliet/full.html

raj.demographics*Romeo and Juliet Demographics*

Description

A dataset containing Romeo and Juliet demographic information for the characters.

Format

A data frame with 34 rows and 4 variables

Details

- person. Character in the play
- sex. Gender
- fam.aff. Family affiliation of character
- died. Dummy coded death variable (0-no; 1-yes); if yes the character dies in the play

References

http://shakespeare.mit.edu/romeo_juliet/full.html

rajPOS*Romeo and Juliet Split in Parts of Speech*

Description

A dataset containing a list from [pos](#) using the [raj](#) data set (see [pos](#) for more information).

Format

A list with 4 elements

Details

text The original text

POStagged The original words replaced with parts of speech in context.

POSprop Dataframe of the proportion of parts of speech by row.

POSfreq Dataframe of the frequency of parts of speech by row.

References

http://shakespeare.mit.edu/romeo_juliet/full.html

 rajSPLIT

Romeo and Juliet (Complete & Split)

Description

A dataset containing the complete dialogue of Romeo and Juliet with turns of talk split into sentences.

Format

A data frame with 2151 rows and 8 variables

Details

- person. Character in the play
- sex. Gender
- fam.aff. Family affiliation of character
- died. Dummy coded death variable (0-no; 1-yes); if yes the character dies in the play
- dialogue. The spoken dialogue
- act. The act (akin to repeated measures)
- stem.text. Text that has been stemmed

References

http://shakespeare.mit.edu/romeo_juliet/full.html

 rank_freq_mplot

Rank Frequency Plot

Description

rank_freq_mplot - Plot a faceted word rank versus frequencies by grouping variable(s).

rank_freq_plot - Plot word rank versus frequencies.

Usage

```
rank_freq_mplot(text.var, grouping.var = NULL, ncol = 4,
  jitter = 0.2, log.freq = TRUE, log.rank = TRUE,
  hap.col = "red", dis.col = "blue", alpha = 1,
  shape = 1, title = "Rank-Frequency Plot", digits = 2,
  plot = TRUE)
```

```
rank_freq_plot(words, frequencies, plot = TRUE,
  title.ext = NULL, jitter.ammount = 0.1,
  log.scale = TRUE, hap.col = "red", dis.col = "blue")
```


Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
ncol	integer value indicating the number of columns in the facet wrap.
jitter	Ammount of horizontal jitter to add to the points.
log.freq	logical. If TURE plots the frequencies in the natural log scale.
log.rank	logical. If TURE plots the ranks in the natural log scale.
hap.col	Color of the hapax legomenon points.
dis.col	Color of the dis legomenon points.
alpha	Transparency level of points (ranges between 0 and 1).
title	Optional plot title.
digits	Integer; number of dicimal places to round.
plot	logical. If TRUE provides a rank frequency plot.
words	A vector of words.
frequencies	A vector of frequencies corresponding to the words argument.
title.ext	The title extension that extends: "Rank-Frequency Plot ..."
jitter.ammount	Ammount of horizontal jitter to add to the points.
log.scale	logical. If TRUE plots the rank and frequency as a log scale.

Value

Returns a rank-frequency plot and a list of three dataframes:

WORD_COUNTS	The word frquencies supplied to rank_freq_plot or created by rank_freq_mplot.
RANK_AND_FREQUENCY_STATS	A dataframe of rank and frequencies for the words used in the text.
LEGOMENA_STATS	A dataframe displaying the percent hapax legomena and percent dis legomena of the text.

References

Zipf, G. K. (1949). Human behavior and the principle of least effort. Cambridge, Massachusetts: Addison-Wesley. p. 1.

Examples

```
## Not run:
#rank_freq_mplot EXAMPLES:
rank_freq_mplot(DATA$state, DATA$person, ncol = 2, jitter = 0)
rank_freq_mplot(mraja1spl$dialogue, mraja1spl$person, ncol = 5,
  hap.col = "purple")
rank_freq_mplot(mraja1spl$dialogue, mraja1spl$person, ncol = 5,
  log.freq = FALSE, log.rank = FALSE, jitter = .6)
rank_freq_mplot(raj$dialogue, jitter = .5, alpha = 1/15)
rank_freq_mplot(raj$dialogue, jitter = .5, shape = 19, alpha = 1/15)

#rank_freq_plot EXAMPLES:
```

```

mod <- with(mraja1spl , word_list(dialogue, person, cut.n = 10,
  cap.list=unique(DF$person)))
rank_freq_plot(mod$fwl$Romeo$WORD, mod$fwl$Romeo$FREQ, title.ext = 'Romeo')
rank_freq_plot(mod$fwl$Romeo$WORD, mod$fwl$Romeo$FREQ, plot = FALSE)
rank_freq_plot(mod$fwl$Romeo$WORD, mod$fwl$Romeo$FREQ, title.ext = 'Romeo',
  jitter.ammount = 0.15, hap.col = "darkgreen", dis.col = "purple")
rank_freq_plot(mod$fwl$Romeo$WORD, mod$fwl$Romeo$FREQ, title.ext = 'Romeo',
  jitter.ammount = 0.5, log.scale=FALSE)
lapply(seq_along(mod$fwl), function(i){
  dev.new()
  rank_freq_plot(mod$fwl[[i]]$WORD, mod$fwl[[i]]$FREQ,
    title.ext = names(mod$fwl)[i], jitter.ammount = 0.5, log.scale=FALSE)
})

## End(Not run)

```

read.transcript

Read Transcripts Into R

Description

Read a .docx, .csv or .xlsx files into R.

Usage

```

read.transcript(file, col.names = NULL, text.var = NULL,
  merge.broke.tot = TRUE, header = FALSE, dash = "",
  ellipsis = "...", quote2bracket = FALSE,
  rm.empty.rows = TRUE,
  na.strings = c("999", "NA", "", " "), sep = NULL,
  skip = 0, nontext2factor = TRUE, ...)

```

Arguments

file	The name of the file which the data are to be read from. Each row of the table appears as one line of the file. If it does not contain an absolute path, the file name is relative to the current working directory, getwd().
col.names	Supplies a vector of column names to the transcript columns.
text.var	specifying the name of the text variable will ensure that variable is classed as character. If NULL read.transcript attempts to guess the text.variable (dialogue).
merge.broke.tot	If the file being read in is .docx and the transcript is formatted to have broken space between a single turn of talk read.transcript will attempt to merge these into a single turn of talk.
header	logical. If TRUE the file contains the names of the variables as its first line.
dash	Character to replace the en and em dashes special characters (default is to remove).
ellipsis	Character to replace the ellipsis special characters (default is text ...).
quote2bracket	logical If TRUE replaces curly quotes with curly braces (default is FALSE). If FALSE curly quotes are removed.

<code>rm.empty.rows</code>	logical. If TRUE read.transcript attempts to remove empty rows.
<code>na.strings</code>	A character vector of strings which are to be interpreted as NA values.
<code>sep</code>	The field separator character. Values on each line of the file are separated by this character. The default of NULL instructs read.transcript to use a separator suitable for the file type being read in.
<code>skip</code>	Integer; the number of lines of the data file to skip before beginning to read data.
<code>nontext2factor</code>	logical. If TRUE attempts to convert any non text to a factor.
<code>...</code>	Further arguments to be passed to read.table.

Value

Returns a dataframe of dialogue and people.

Note

If a transcript is a .docx file read transcript expects two columns (generally person and dialogue) with some sort of separator (default is colon separator). .doc files must be converted to .docx before reading in.

Author(s)

Bryan Goodrich and Tyler Rinker <tyler.rinker@gmail.com>.

References

<https://github.com/trinker/qdap/wiki/Reading-Transcripts-into-R>

Examples

```
## Not run:
doc1 <- system.file("extdata/trans1.docx", package = "qdap")
doc2 <- system.file("extdata/trans2.docx", package = "qdap")
doc3 <- system.file("extdata/trans3.docx", package = "qdap")
doc4 <- system.file("extdata/trans4.xlsx", package = "qdap")

read.transcript(doc1)
dat <- read.transcript(doc1, col.names = c("person", "dialogue"))
dat
rm_row(dat, "person", "[C]") #remove bracket row

read.transcript(doc2) #throws an error
read.transcript(doc2, skip = 1)
read.transcript(doc3, skip = 1) #wrong sep
read.transcript(doc3, sep = "-", skip = 1)
read.transcript(doc4)

## End(Not run)
```

replacer	<i>Replace Cells in a Matrix or Data Frame</i>
----------	--

Description

Replace elements of a dataframe, matrix or vector with least restrictive class.

Usage

```
replacer(dat, replace = 0, with = "-")
```

Arguments

dat	Data; either a dataframe, matrix or vector.
replace	Element to replace.
with	Replacement element.

Value

Returns a dataframe, matrix or vector with the element replaced.

Examples

```
## Not run:
replacer(mtcars, 0, "REP")
replacer(mtcars, 8, NA)
replacer(c("a", "b"), "a", "foo")

## End(Not run)
```

replace_abbreviation	<i>Replace Abbreviations</i>
----------------------	------------------------------

Description

This function replaces abbreviations with long form.

Usage

```
replace_abbreviation(text.var,
  abbreviation = qdap::abbreviations, replace = NULL,
  ignore.case = TRUE)
```

Arguments

text.var	The text variable.
abbreviation	A two column key of abbreviations (column 1) and long form replacements (column 2) or a vector of abbreviations. Default is to use qdap's abbreviations data set.
replace	A vector of long form replacements if a data frame is not supplied to the abbreviation argument.
ignore.case	logical. If TRUE replaces without regard to capitalization.

Value

Returns a vector with abbreviations replaced.

See Also

[bracketX](#), [qprep](#), [replace_number](#), [replace_symbol](#)

Examples

```
## Not run:
x <- c("Mr. Jones is here at 7:30 p.m.",
      "Check it out at www.github.com/trinker/qdap",
      "i.e. He's a sr. dr.; the best in 2012 A.D.",
      "the robot at t.s. is 10ft. 3in.")

replace_abbreviation(x)

#create abbreviation and replacement vectors
abv <- c("in.", "ft.", "t.s.")
repl <- c("inch", "feet", "talkstats")

replace_abbreviation(x, abv, repl)

KEY <- rbind(abbreviations, data.frame(abv = abv, rep = repl))
replace_abbreviation(x, KEY)

## End(Not run)
```

replace_number

Replace Numerbers With Text Representation

Description

Replaces numeric represented numbers with words (e.g. 1001 becomes one thousand one).

Usage

```
replace_number(text.var, num.paste = "separate")
```

Arguments

text.var	The text variable.
num.paste	A character vector of either "separate" or "combine". Of "separate" is specified the elements of larger numbers are separated with spaces. If "combine" is selected the elements will be joined without spaces.

Value

Returns a vector with abbreviations replaced.

References

Fox, J. (2005). Programmer's niche: How do you spell that number? R News. Vol. 5(1), pp. 51-55.

See Also

[bracketX](#), [replace_abbreviation](#), [qprep](#), [replace_symbol](#)

Examples

```
## Not run:
x <- c("I like 346,457 ice cream cones.", "They are 99 percent good")
y <- c("I like 346457 ice cream cones.", "They are 99 percent good")
replace_number(x)
replace_number(y)
replace_number(x, "combine")

## End(Not run)
```

replace_symbol

Replace Symbols With Word Equivalents

Description

This function replaces symbols with word equivalents (e'g' @ becomes "at").

Usage

```
replace_symbol(text.var, dollar = TRUE, percent = TRUE,
  pound = TRUE, at = TRUE, and = TRUE, with = TRUE)
```

Arguments

text.var	The text variable.
dollar	logical. If TRUE replaces dollar sign (\$) with "dollar".
percent	logical. If TRUE replaces percent sign (%) with "percent".
pound	logical. If TRUE replaces pound sign (#) with "number".
at	logical. If TRUE replaces at sign (@) with "at".
and	logical. If TRUE replaces and sign (&) with "and".
with	logical. If TRUE replaces with sign (w/) with "with".

Value

Returns a character vector with symbols replaced..

See Also

[bracketX](#), [replace_abbreviation](#), [replace_number](#), [qprep](#)

Examples

```
## Not run:
x <- c("I am @ Jon's & Jim's w/ Marry", "I owe $41 for food", "two is 10%
  of a #")
replace_symbol(x)

## End(Not run)
```

rm_row	<i>Remove Rows That Contain Markers</i>
--------	---

Description

rm_row - Remove rows from a data set that contain a given marker/term.

rm_empty_row - Removes the empty rows of a data set that are common in reading in data (default method in read.transcript).

Usage

```
rm_row(dataframe, search.column, terms)
```

```
rm_empty_row(dataframe)
```

Arguments

dataframe	A dataframe object.
search.column	Column name to search for markers/terms.
terms	Terms/markers of the rows that are to be removed from the dataframe. The term/marker must appear at the beginning of the string and is case sensitive.

Value

rm_row - returns a dataframe with the termed/marked rows removed.

rm_empty_row - returns a dataframe with empty rows removed.

Examples

```
## Not run:
#rm_row EXAMPLE:
rm_row(DATA, "person", c("sam", "greg"))
rm_row(DATA, 1, c("sam", "greg"))
rm_row(DATA, "state", c("Comp"))

#rm_empty_row EXAMPLE:
x <- matrix(rep(" ", 4), ncol = 2)
dat <- DATA[, c(1, 4)]
colnames(x) <- colnames(dat)
(dat <- data.frame(rbind(dat, x)))
rm_empty_row(dat)

## End(Not run)
```

scrubber	<i>Use to clean text variables when importing a new data set.</i>
----------	---

Description

Use to clean text variables when importing a new data set. Removes extra white spaces other textual anomalies that may cause errors.

Usage

```
scrubber(text.var, num2word = FALSE, rm.quote = TRUE,
         fix.comma = TRUE, ...)
```

Arguments

text.var	The text variable
num2word	logical If TRUE replaces a numbers with text representations.
fix.comma	logical If TRUE removes any spaces before a comma.
rm.quote	logical If TRUE removes and \".
...	Other arguments passed to replace_number.

Value

Returns a parsed character vector.

See Also

[strip](#)

Examples

```
## Not run:
x <- c("I like 456 dogs , don't you?")
scrubber(x)
scrubber(x, TRUE)

## End(Not run)
```

Search	<i>Search Columns of a Data Frame</i>
--------	---------------------------------------

Description

Find terms located in columns of a data frame.

Usage

```
Search(dataframe, term, column.name = NULL,
       max.distance = 0.02, ...)
```


Arguments

dataframe	A dataframe object to search.
term	A character vector term to search for.
column.name	Optional column of the data frame to search (nome or index).
max.distance	Maximum distance allowed for a match. Expressed either as integer, or as a fraction of the pattern length times the maximal transformation cost (will be replaced by the smallest integer not less than the corresponding fraction).
...	Other arguments passed to agrep.

Value

Returns the rows of the data frame that amtch the search term.

Examples

```
## Not run:
SampDF <- data.frame("islands"=names(islands)[1:32],mtcars)

Search(SampDF, "Cuba", "islands")
Search(SampDF, "New", "islands")
Search(SampDF, "Ho")
Search(SampDF, "Ho", max.distance = 0)
Search(SampDF, "Axel Heiberg")
Search(SampDF, 19) #too much tolerance in max.distance
Search(SampDF, 19, max.distance = 0)
Search(SampDF, 19, "qsec", max.distance = 0)

## End(Not run)
```

sentSplit

*Sentence Splitting***Description**

sentSplit - Splits turns of talk into individual sentences (provided proper punctuation is used). This procedure is usually done as part of the data read in and cleaning process.

sentCombine - Combines sentences by the same grouping variable together.

TOT - Convert the tot column from [sentSplit](#) to turn of talk index (no sub sentence). Generally, for internal use.

Usage

```
sentSplit(dataframe, text.var,
  endmarks = c("?", ".", "!", "|"),
  incomplete.sub = TRUE, rm.bracket = TRUE,
  stem.col = FALSE, text.place = "right", ...)

sentCombine(text.var, grouping.var = "person")

TOT(tot)
```

Arguments

<code>dataframe</code>	A dataframe that contains the person and text variable.
<code>text.var</code>	The text variable.
<code>endmarks</code>	A character vector of endmarks to split turns of talk into sentences.
<code>incomplete.sub</code>	logical. If TRUE detects incomplete sentences and replaces with " ".
<code>rm.bracket</code>	logical. If TRUE removes brackets from the text.
<code>stem.col</code>	logical. If TRUE stems the text as a new column.
<code>text.place</code>	A character string giving placement location of the text column. This must be one of the strings "original", "right" or "left".
<code>...</code>	Additional options passed to <code>stem2df</code> .
<code>grouping.var</code>	The grouping variable (usually "person"). Does not take multiple vectors as most <code>qdap</code> functions do.
<code>tot</code>	A tot column from a <code>sentSplit</code> output.

Value

`sentSplit` - returns a dataframe with turn of talk broken apart into sentences. Optionally a stemmed version of the text variable may be returned as well.

`sentCombine` - returns a list of vectors with the continuous sentences by `grouping.var` pasted together. returned as well.

`TOT` - returns a numeric vector of the turns of talk without sentence sub indexing (e.g. 3.2 become 3).

Author(s)

Dason Kurkiewicz and Tyler Rinker <tyler.rinker@gmail.com>.

See Also

`bracketX`, `incomplete.replace`, `stem2df`, `TOT`

Examples

```
## Not run:
#sentSplit EXAMPLE:
sentSplit(DATA, "state")
sentSplit(DATA, "state", stem.col = FALSE)
sentSplit(DATA, "state", text.place = "left")
sentSplit(DATA, "state", text.place = "original")
sentSplit(raj, "dialogue")

#sentCombine EXAMPLE:
dat <- sentSplit(DATA, "state", stem.col = FALSE)
sentCombine(dat$state, dat$person)
sentCombine(dat$state, dat$sex)

#TOT EXAMPLE:
dat <- sentSplit(DATA, "state", stem.col = FALSE)
TOT(dat$tot)

## End(Not run)
```

spaste	<i>Add Leading/Trailing Spaces</i>
--------	------------------------------------

Description

Adds trailing and/or leading spaces to a vector of terms.

Usage

```
spaste(terms, trailing = TRUE, leading = TRUE)
```

Arguments

terms	A character vector of terms to insert trailing and/or leading spaces.
leading	logical. If TRUE inserts a leading space in the terms.
trailing	logical. If TRUE inserts a trailing space in the terms.

Value

Returns a character vector with trailing and/or leading spaces.

Examples

```
## Not run:
spaste(Top25Words)
spaste(Top25Words, FALSE)
spaste(Top25Words, ,FALSE)

## End(Not run)
```

speakerSplit	<i>Break and Stretch if Multiple Persons per Cell</i>
--------------	---

Description

Look for cells with multiple people and create separate rows for each person.

Usage

```
speakerSplit(dataframe, person.var = 1,
  sep = c("and", "&", ", "), track.reps = FALSE)
```

Arguments

dataframe	A dataframe that contains the person variable.
person.var	The person variable to be stretched.
sep	The separator(s) to search for and break on. Default is: c(" ", "_", ";")
track.reps	logical. If TRUE leaves the row names of person variable cells that were repeated and stretched.

Value

Returns an expanded dataframe with person variable stretched and accompanying rows repeated.

Examples

```
## Not run:
DATA$person <- as.character(DATA$person)
DATA$person[c(1, 4, 6)] <- c("greg, sally, & sam",
  "greg, sally", "sam and sally")

speakerSplit(DATA)
speakerSplit(DATA, track.reps=TRUE)

DATA$person[c(1, 4, 6)] <- c("greg_sally_sam",
  "greg.sally", "sam; sally")

speakerSplit(DATA, sep = c(".", "_", ";"))

DATA <- qdap::DATA #reset DATA

## End(Not run)
```

stemmer	<i>Stem Text</i>
---------	------------------

Description

stemmer - Stems a vector of text strings.

stem.words - Wrapper for stemmer that stems a vector of words.

stem2df - Wrapper for stemmer that stems a vector of text strings and returns a dataframe with the vector added..

Usage

```
stemmer(text.var, rm.bracket = TRUE, capitalize = TRUE,
  warn = TRUE, ...)

stem.words(...)

stem2df(dataframe, text.var, stem.name = NULL, ...)
```

Arguments

text.var	The text variable. In stemmer this is a vector text string. For stem2df this is a character vector of length one naming the text column.
rm.bracket	logical. If TRUE brackets are removed from the text.
capitalize	logical. If TRUE selected terms are capitalized
warn	logical. If TRUE warns about rows not ending with standard qdap punctuation endmarks.

...	Various: stemmer - <i>Other arguments passed to</i> capitalizer stem.words - <i>Words or terms.</i> stem2df - <i>Other arguments passed to</i> stemmer
dataframe	A dataframe object.
stem.name	A character vector of length one for the stemmed column. If NULL defaults to "stem.text".

Value

stemmer - returns a character vector with stemmed text.

stem.words - returns a dataframe with a character vector with.

stem2df - returns a dataframe with a character vector with stemmed text.

See Also

[capitalizer](#)

Examples

```
## Not run:
#stemmer EXAMPLE:
stemmer(DATA$state)
stemmer(raj$dialogue)

#stem.words EXAMPLE:
stem.words(doggies, jumping, swims)

#stem2df EXAMPLE:
stem2df(DATA, "state", "new")

## End(Not run)
```

stopwords

Remove Stopwords

Description

Transcript apply the removal of stopwords.

Usage

```
stopwords(textString, stopwords = Top25Words,
  unlist = FALSE, separate = TRUE, strip = FALSE,
  unique = FALSE, char.keep = NULL, names = FALSE,
  ignore.case = TRUE, apostrophe.remove = FALSE, ...)
```

Arguments

<code>textString</code>	A character string of text or a vector of character strings.
<code>stopwords</code>	A character vector of words to remove from the text. <code>qdap</code> has a number of data sets that can be used as stopwords including: <code>Top200Words</code> , <code>Top100Words</code> , <code>Top25Words</code> . For the <code>tm</code> package's traditional English stop words use <code>tm::stopwords("english")</code>
<code>unlist</code>	logical. If <code>TRUE</code> unlists into one vector. General use intended for when <code>separate</code> is <code>FALSE</code> .
<code>separate</code>	logical. If <code>TRUE</code> separates sentences into words. If <code>FALSE</code> retains sentences.
<code>strip</code>	logical. If <code>TRUE</code> strips the text of all punctuation except apostrophes.
<code>unique</code>	logical. If <code>TRUE</code> keeps only unique words (if <code>unlist</code> is <code>TRUE</code>) or sentences (if <code>unlist</code> is <code>FALSE</code>). General use intended for when <code>unlist</code> is <code>TRUE</code> .
<code>char.keep</code>	If <code>strip</code> is <code>TRUE</code> this argument provides a means of retaining supplied character(s).
<code>names</code>	logical. If <code>TRUE</code> will name the elements of the vector or list with the original <code>textString</code> .
<code>ignore.case</code>	logical. If <code>TRUE</code> stop words will be removed regardless of case. Additionally, case will be stripped from the text. If <code>FALSE</code> stopwords removal is contingent upon case. Additionally, case is not stripped.
<code>apostrophe.remove</code>	logical. If <code>TRUE</code> removes apostrophe's from the output.
<code>...</code>	further arguments passed to <code>strip</code> function

Value

Returns a vector of sentences, vector of words, or (default) a list of vectors of words with stop words removed. Output depends on supplied arguments.

See Also

[strip](#), [bag.o.words](#), [stopwords](#)

Examples

```
## Not run:
stopwords(DATA$state)
stopwords(DATA$state, tm::stopwords("english"))
stopwords(DATA$state, Top200Words)
stopwords(DATA$state, Top200Words, strip = TRUE)
stopwords(DATA$state, Top200Words, separate = FALSE)
stopwords(DATA$state, Top200Words, separate = FALSE, ignore.case = FALSE)
stopwords(DATA$state, Top200Words, unlist = TRUE)
stopwords(DATA$state, Top200Words, unlist = TRUE, strip=TRUE)
stopwords(DATA$state, Top200Words, unlist = TRUE, unique = TRUE)

## End(Not run)
```

strip*Strip Text*

Description

Strip text of unwanted characters.

Usage

```
strip(x, char.keep = NULL, digit.remove = TRUE,  
      apostrophe.remove = TRUE, lower.case = TRUE)
```

Arguments

<code>x</code>	The text variable.
<code>char.keep</code>	A character vector of symbol character (i.e. punctuation) that strip should keep. The default is to strip everything except apostrophes.
<code>digit.remove</code>	logical. If TRUE strips digits from the text.
<code>apostrophe.remove</code>	logical. If TRUE removes apostrophe's from the output.
<code>lower.case</code>	logical. If TRUE forces all alpha characters to lower case.

Value

Retruns a vector of text that has been stripped of unwanted characters.

See Also

[stopwords](#)

Examples

```
## Not run:  
strip(DATA$state)  
strip(DATA$state, apostrophe.remove=FALSE)  
strip(DATA$state, char.keep = c("?", "."))  
  
## End(Not run)
```

strWrap*Wrap Character Strings to Format Paragraphs*

Description

A wrapper for [as.character](#) that writes to the Mac/Windows clipboard.

Usage

```
strWrap(text = "clipboard", width = 70, copy2clip = TRUE)
```

Arguments

- text character vector, or an object which can be converted to a character vector by [as.character](#).
- width A positive integer giving the target column for wrapping lines in the output.
- copy2clip logical. If TRUE attempts to copy the output to the clipboard.

Value

Prints a wrapped text vector to the console and copies the wrapped text to the clipboard on a Mac or Windows machine.

See Also

[strwrap](#)

Examples

```
## Not run:
x <- paste2(DATA$state, sep = " ")
strWrap(x)
strWrap(x, 10)
#should be copied to the clipboard on a Mac or Windows machine.

## End(Not run)
```

syllable.sum	<i>Syllabication</i>
--------------	----------------------

Description

- syllable.sum - Count the number of syllables per row of text.
- syllable.count - Count the number of syllables in a single text string.
- polysyllable.sum - Count the number of polysyllables per row of text.
- combo_syllable.sum - Count the number of both syllables and polysyllables per row of text.

Usage

```
syllable.sum(text.var, parallel = FALSE)

syllable.count(text, remove.bracketed = TRUE,
  algorithm.report = FALSE)

polysyllable.sum(text.var, parallel = FALSE)

combo_syllable.sum(text.var, parallel = FALSE)
```


Arguments

<code>text.var</code>	The text variable
<code>parallel</code>	logical. If TRUE attempts to run the function on multiple cores. Note that this may not mean a speed boost if you have one core or if the data set is smaller as the cluster takes time to create.
<code>text</code>	A single character vector of text.
<code>remove.bracketed</code>	logical. If TRUE brackets are removed from the analysis.
<code>algorithm.report</code>	logical. If TRUE generates a report of words not found in the dictionary (i.e. syllables were calculated with an algorithm).

Value

`syllable.sum` - returns a vector of syllable counts per row.

`syllable.count` - returns a dataframe of syllable counts and algorithm/dictionary uses and, optionally, a report of words not found in the dictionary.

`polysyllable.sum` - returns a vector of polysyllable counts per row.

`combo_syllable.sum` - returns a dataframe of syllable and polysyllable counts per row.

Note

The worker of all the syllable functions is `syllable.count` though it is not intended for direct use on a transcript. This function relies on a combined dictionary lookup (based on the Nettekorp Corpus (Sejnowski & Rosenberg, 1987)) and backup algorithm method.

References

Sejnowski, T.J., and Rosenberg, C.R. (1987). "Parallel networks that learn to pronounce English text" in *Complex Systems*, 1, 145-168.

Examples

```
## Not run:
syllable.count("Robots like Dason lie.")
syllable.count("Robots like Dason lie.", algorithm.report = TRUE)
syllable.sum(DATA$state)
polysyllable.sum(DATA$state)
combo_syllable.sum(DATA$state)

## End(Not run)
```

termco.a

*Search For and Count Terms***Description**

termco.a - Search a transcript by any number of grouping variables for categories (themes) of grouped root terms. While there are other termco functions in the termco family (i.e. termco.d) termco.a is a wrapper for general use.

termco.d - Search a transcript by any number of grouping variables for root terms.

term.match - Search a transcript for words that exactly match term(s).

termco2mat - Convert a termco dataframe to a matrix for use with visualization functions (e.g. heatmap2 of the gplots package).

Usage

```
termco.a(text.var, grouping.var = NULL, match.list,
  short.term = TRUE, ignore.case = TRUE, elim.old = TRUE,
  output = "percent", digits = 2,
  apostrophe.remove = FALSE, char.keep = NULL,
  digit.remove = NULL, ...)
```

```
termco.d(text.var, grouping.var = NULL, match.string,
  short.term = FALSE, ignore.case = TRUE,
  zero.replace = 0, output = "percent", digits = 2,
  apostrophe.remove = FALSE, char.keep = NULL,
  digit.remove = TRUE, ...)
```

```
term.match(text.var, terms, return.list = TRUE,
  apostrophe.remove = FALSE)
```

```
termco2mat(dataframe, drop.wc = TRUE, short.terms = TRUE,
  rm.zerocol = FALSE, no.quote = TRUE, transform = TRUE,
  trim.terms = TRUE)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
match.list	a list of named character vectors
short.term	logical. If TRUE column names are trimmed versions of the match list, otherwise the terms are wrapped with 'term(phrase)'
ignore.case	logical. If TRUE case is ignored.
elim.old	logical. If TRUE eliminates the columns that are combined together by the named match.list.
output	Type of proportion output; either "proportion" (decimal format) or "percent". Default is "percent".

<code>digits</code>	integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed.
<code>apostrophe.remove</code>	logical. If TRUE removes apostrophes from the text before examining.
<code>char.keep</code>	A character vector of symbol character (i.e. punctuation) that strip should keep. The default is to strip everything except apostrophes.
<code>digit.remove</code>	logical. If TRUE strips digits from the text.
<code>...</code>	Other argument supplied to strip.
<code>match.string</code>	A vector of terms to search for. When using inside of <code>term.match</code> the term(s) must be words or partial words but do not have to be when using <code>termco.d</code> (i.e. they can be phrases, symbols etc.).
<code>zero.replace</code>	Value to replace 0 values with.
<code>term.match</code>	- return.list logical. If TRUE returns the output for multiple terms as a list by term rather than a vector.
<code>dataframe</code>	A termco.a (or termco.d) dataframe or object.
<code>drop.wc</code>	logical. If TRUE the word count column will be dropped.
<code>short.colnames</code>	logical. If TRUE the "term()" portion of column names will be dropped.
<code>no.quote</code>	logical. If TRUE the matrix will be printed without quotes if it's character.
<code>transform</code>	logical. If TRUE the matrix will be transformed.

Value

`termco.a` & `termco.d` - both return a list, of class "termco.d", of data frames and information regarding word counts:

<code>raw</code>	raw word counts by grouping variable
<code>prop</code>	proportional word counts by grouping variable; proportional to each individual's word use
<code>rnp</code>	a character combination data frame of raw and proportional
<code>zero_replace</code>	value to replace zeros with; mostly internal use
<code>output</code>	character value for output type (either "proportion" or "percent"; mostly internal use)
<code>digits</code>	integer value of number of digits to display; mostly internal use

`term.match` - returns a list or vector of possible words that match term(s).

`termco2mat` - returns a matrix of term counts.

Note

The `match.list/match.string` is (optionally) case and character sensitive. Spacing is an important way to grab specific words and requires careful thought. Using "read" will find the words "bread", "read" "reading", and "ready". If you want to search for just the word "read" you'd supply a vector of `c(" read ", " reads", " reading", " reader")`. To search for non character arguments (i.e. numbers and symbols) additional arguments from `strip` must be passed.

See Also

[termco.c](#)

Examples

```
## Not run:
#termco.a examples:

# General form for match.list
#
# ml <- list(
#   cat1 = c(),
#   cat2 = c(),
#   catn = c()
# )

ml <- list(
  cat1 = c(" the ", " a ", " an "),
  cat2 = c(" I' " ),
  "good",
  the = c("the", " the ", " the", "the")
)

(dat <- with(raj.act.1, termco.a(dialogue, person, ml)))
names(dat)
dat$rn timer #useful for presenting in tables
dat$raw timer #prop and raw are useful for performing calculations
dat$prop timer
dat <- with(raj.act.1, termco.a(dialogue, person, ml,
  short.term = FALSE, elim.old=FALSE))

dat2 <- data.frame(dialogue=c("@bryan is bryan good @br",
  "indeed", "@ brian"), person=qcv(A, B, A))

ml <- list(wrds=c("bryan", "indeed"), bryan=c("bryan", "@ br", "@br"))

with(dat2, termco.a(dialogue, person, match.list=ml, char.keep="@"))

with(dat2, termco.a(dialogue, person, match.list=ml,
  char.keep="@", output="proportion"))

DATA$state[1] <- "12 4 rgfr r0ffrg0"
termco.a(DATA$state, DATA$person, '0', digit.remove=FALSE)

#Using with term.match and exclude
exclude(term.match(DATA$state, qcv(th), FALSE), "truth")
termco.a(DATA$state, DATA$person, exclude(term.match(DATA$state, qcv(th),
  FALSE), "truth"))
MTCH.LST <- exclude(term.match(DATA$state, qcv(th, i)), qcv(truth, stinks))
termco.a(DATA$state, DATA$person, MTCH.LST)

#termco.d examples:
term.match(DATA$state, qcv(i, the))
termco.d(DATA$state, DATA$person, c(" the", " i'"))
termco.d(DATA$state, DATA$person, c(" the", " i'"), ignore.case=FALSE)
termco.d(DATA$state, DATA$person, c(" the ", " i'"))

# termco2mat example:
MTCH.LST <- exclude(term.match(DATA$state, qcv(a, i)), qcv(is, it, am, shall))
termco_obj <- termco.a(DATA$state, DATA$person, MTCH.LST)
```

```
termco2mat(termco_obj)

# as a visual
dat <- termco2mat(termco_obj)
library(gplots)
heatmap.2(dat, trace="none")

## End(Not run)
```

termco.c

*Combine Columns from a termco_ Object***Description**

Combines the columns of a termco_ object. Generally intended for internal use but documented for completeness.

Usage

```
termco.c(termco.d.object, combined.columns, new.name,
  short.term = FALSE, zero.replace = NULL,
  lazy.term = TRUE, elim.old = TRUE, output = "percent")
```

Arguments

termco.d.object	An object generated by either termco_a or termco_d.
combined.columns	The names/indexes of the columns to be combined.
new.name	A character vector of length one to name the new combined column.
short.term	logical. If TRUE column names are trimmed versions of the match list, otherwise the terms are wrapped with 'term(phrase)'
zero.replace	Value to replace zeros with.
lazy.term	logical. If TRUE is approximate amtching (generally for internal use).
elim.old	logical. If TRUE eliminates the columns that are combined together by the named match.list.
output	Type of proportion output; either "proportion" (decimal format) or "percent". Default is "percent".

Value

Returns a return a list, of class "termco.c", of data frames and information regarding word counts:

raw	raw word counts by grouping variable
prop	proportional word counts by grouping variable; proportional to each individual's word use
rnp	a character combination data frame of raw and proportional
zero_replace	value to replace zeros with; mostly internal use
output	character value for outpur type (either "proportion" or "percent"; mostly internal use
digits	integer value od number of digits to display; mostly internal use

See Also[termco.a](#)

text2color

Map Words to Colors

Description

A dictionary lookup that maps words to colors.

Usage

```
text2color(words, recode.words, colors)
```

Arguments

words	A vector of words.
recode.words	A vector of unique words or a list of unique word vectors that will be matched against corresponding colors.
colors	A vector of colors of equal in length to recode.words + 1 (the +1 is for unmatched words).

Value

Returns a vector of mapped colors equal in length to the words vector.

See Also[lookup](#)**Examples**

```
## Not run:
set.seed(10)
x <- data.frame(X1 = sample(Top25Words[1:10], 20, TRUE))
text2color(x$X1, qcv(the, and, it), qcv(red, green, blue)) #blue was recycled
text2color(x$X1, qcv(the, and, it), qcv(red, green, blue, white))
x$X2 <- text2color(x$X1, list(qcv(the, and, it), "that"), qcv(red, green,
  white))
x

## End(Not run)
```

Top100Words*Fry's 100 Most Commonly Used English Words*

Description

A stopword list containing a character vector of stopwords.

Format

A character vector with 100 elements

Details

Fry's Word List: The first 25 make up about one-third of all printed material in English. The first 100 makem up about one-half of all printed material in English. The first 300 makem up about 65% of all printed material in English."

References

Fry, E. B. (1997). Fry 1000 instant words. Lincolnwood, IL: Contemporary Books.

Top200Words*Fry's 200 Most Commonly Used English Words*

Description

A stopword list containing a character vector of stopwords.

Format

A character vector with 200 elements

Details

Fry's Word List: The first 25 make up about one-third of all printed material in English. The first 100 makem up about one-half of all printed material in English. The first 300 makem up about 65% of all printed material in English."

References

Fry, E. B. (1997). Fry 1000 instant words. Lincolnwood, IL: Contemporary Books.

Top25Words

Fry's 25 Most Commonly Used English Words

Description

A stopword list containing a character vector of stopwords.

Format

A character vector with 25 elements

Details

Fry's Word List: The first 25 make up about one-third of all printed material in English. The first 100 make up about one-half of all printed material in English. The first 300 make up about 65% of all printed material in English."

References

Fry, E. B. (1997). Fry 1000 instant words. Lincolnwood, IL: Contemporary Books.

trans.cloud

Word Clouds by Grouping Variable

Description

Produces word clouds with optional theme coloring by grouping variable.

Usage

```
trans.cloud(text.var = NULL, grouping.var = NULL,
  word.list = NULL, stem = FALSE, target.words = NULL,
  expand.target = TRUE, target.exclude = NULL,
  stopwords = NULL, min.freq = 1, caps = TRUE,
  caps.list = NULL, random.order = FALSE, rot.per = 0,
  cloud.colors = NULL, title = TRUE, cloud.font = NULL,
  title.font = NULL, title.color = "black",
  title.padj = -4.5, title.location = 3,
  title.cex = NULL, title.names = NULL,
  proportional = FALSE, max.word.size = NULL,
  min.word.size = 0.5, legend = NULL, legend.cex = 0.8,
  legend.location = c(-0.03, 1.03), char.keep = NULL,
  char2space = NULL, ...)
```


Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
word.list	A frequency word list passed from word_list .
stem	logical. If TRUE the text.var will be stemmed.
target.words	A named list of vectors of words whose length corresponds to cloud.colors (+1 length in cloud colors for non matched terms).
expand.target	logical. If TRUE agrep will be used to expand the target.words.
target.exclude	A vector of words to exclude from the target.words.
stopwords	Words to exclude from the cloud.
min.freq	An integer value indicating the minimum frequency a word must appear to be included.
caps	logical. If TRUE selected words will be capitalized.
caps.list	A vector of words to capitalize (caps must be TRUE).
random.order	Plot words in random order. If false, they will be plotted in decreasing frequency.
rot.per	Proportion words with 90 degree rotation.
cloud.colors	A vector of colors equal to the length of target words +1.
title	logical. IF TRUE adds a title corresponding to the grouping.var.
cloud.font	The font family of the cloud text.
title.font	The font family of the cloud title.
title.color	A character vector of length one corresponding to the color of the title.
title.padj	Adjustment for the title. For strings parallel to the axes, padj = 0 means right or top alignment, and padj = 1 means left or bottom alignment.
title.location	On which side of the plot (1=bottom, 2=left, 3=top, 4=right).
title.cex	Character expansion factor for the title. NULL and NA are equivalent to 1.0.
title.names	Optional vector of title names equal in length to the grouping.var that will override the default use of the grouping.var names.
proportional	logical. If TRUE scales the word clouds across grouping.var to allow cloud to cloud comparisons.
max.word.size	A size argument to control the minimum size of the words.
min.word.size	A size argument to control the maximum size of the words.
legend	A character vector of names corresponding to the number of vectors in target.words.
legend.cex	Character expansion factor for the legend. NULL and NA are equivalent to 1.0.
legend.location	The x and y co-ordinates to be used to position the legend.
char.keep	A character vector of symbol character (i.e. punctuation) that strip should keep. The default is to strip everything except apostrophes. This enables the use of special characters to be turned into spaces or for characters to be retained.
char2space	A vector of characters to be turned into spaces. If char.keep is NULL, char2space will activate this argument.

Value

Returns a series of word cloud plots with target words (themes) colored.

See Also

[wordcloud](#)

Examples

```
## Not run:
terms <- list(
  I=c("i", "i'm"),
  mal=qcv(stinks, dumb, distrust),
  articles=qcv(the, a, an),
  pronoun=qcv(we, you)
)

with(DATA, trans.cloud(state, person, target.words=terms,
  cloud.colors=qcv(red, green, blue, black, gray65),
  expand.target=FALSE, proportional=TRUE))

with(DATA, trans.cloud(state, person, target.words=terms,
  stopwords=exclude(with(DATA, unique(bag.o.words(state))),
    unique(unlist(terms))),
  cloud.colors=qcv(red, green, blue, black, gray65),
  expand.target=FALSE, proportional=TRUE))

## End(Not run)
```

trans.venn	<i>Venn Diagram by Grouping Variable</i>
------------	--

Description

Produce a venn diagram by grouping variable.

Usage

```
trans.venn(text.var, grouping.var, stopwords = NULL,
  rm.duplicates = TRUE, title = TRUE, title.font = NULL,
  title.color = "black", title.cex = NULL,
  title.name = NULL, legend = TRUE, legend.cex = 0.8,
  legend.location = "bottomleft",
  legend.text.col = "black", legend.horiz = FALSE, ...)
```

Arguments

- text.var The text variable.
- grouping.var The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
- stopwords Words to exclude from the analysis.

rm.duplicates	logical. IF TRUE removes the duplicated words from the analysis (only single usage is considered).
title	logical. IF TRUE adds a title corresponding to the grouping.var.
title.font	The font family of the cloud title.
title.color	A character vector of length one corresponding to the color of the title.
title.cex	Character expansion factor for the title. NULL and NA are equivalent to 1.0
title.name	A title for the plot.
legend	logical. If TRUE uses the names from the target.words list corresponding to cloud.colors.
legend.cex	Character expansion factor for the legend. NULL and NA are equivalent to 1.0.
legend.location	The x and y co-ordinates to be used to position the legend. The location may also be specified by setting x to a single keyword from the list "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright", "right" and "center". This places the legend on the inside of the plot frame at the given location.
legend.text.col	The color used for the legend text.
legend.horiz	logical; if TRUE, set the legend horizontally rather than vertically.
...	Other arguments passed to plot.

Value

Returns a venn plot by grouping variable.

Note

The algorithm used to overlap the venn circles becomes increasingly overburdened and less accurate with increased grouping variables. An alternative is to use a network plot with dissimilarity measures labeling the edges between nodes (grouping variables).

See Also

[venneuler](#)

Examples

```
## Not run:
with(DATA , trans.venn(state, person, legend.location = "topright"))
#the plot below will take a considerable ammount of time to plot
with(raj.act.1 , trans.venn(dialogue, person, legend.location = "topleft"))

## End(Not run)
```

Trim	<i>Remove Leading/Trailing White Space</i>
------	--

Description

Remove leading/trailing white space.

Usage

```
Trim(x)
```

Arguments

x	The text variable.
---	--------------------

Value

Returns a vector with the leading/trailing white spaces removed.

Examples

```
## Not run:
(x <- c(" talkstats.com ", " really? ", " yeah"))
Trim(x)

## End(Not run)
```

url_dl	<i>Download Instructional Documents</i>
--------	---

Description

This function enables downloading documents for future instructional training.

Usage

```
url_dl(..., url = "http://dl.dropbox.com/u/61803503/")
```

Arguments

...	Document names to download.
url	The download url.

Value

Places a copy of the downloaded document in the users wordking directory.

Note

Not intended for general use.

Examples

```
## Not run:
# download transcript of the debate to working directory
url_dl(pres.deb1.docx, pres.deb2.docx, pres.deb3.docx)

# load multiple files with read transcript and assign to working directory
dat1 <- read.transcript("pres.deb1.docx", c("person", "dialogue"))
dat2 <- read.transcript("pres.deb2.docx", c("person", "dialogue"))
dat3 <- read.transcript("pres.deb3.docx", c("person", "dialogue"))

docs <- qcv(pres.deb1.docx, pres.deb2.docx, pres.deb3.docx)
dir() %in% docs
delete(docs) #remove the documents
dir() %in% docs

## End(Not run)
```

v.outer

Vectorized Version of outer

Description

Vectorized [outer](#).

Usage

```
v.outer(x, FUN, digits = 3, ...)
```

Arguments

x	A matrix, dataframe or equal length list of vectors.
FUN	A vectorized function.
digits	Integer; number of decimal places to round.
...	Other arguments passed to the function supplied to FUN.

Value

Returns a matrix with the vectorized [outer](#) function.

See Also

[outer](#), [cor](#)

Examples

```
## Not run:
pooled.sd <- function(x, y) {
  n1 <- length(x)
  n2 <- length(y)
  s1 <- sd(x)
  s2 <- sd(y)
  sqrt(((n1-1)*s1 + (n2-1)*s2)/((n1-1) + (n2-1)))
}
```

```

}

euc.dist <- function(x,y) sqrt(sum((x - y) ^ 2))
sum2 <- function(x, y) sum(x, y)

v.outer(mtcars, cor)
v.outer(mtcars, pooled.sd)
v.outer(mtcars, euc.dist)
v.outer(mtcars, sum2)

mtcars2 <- lapply(mtcars, function(x) x)
v.outer(mtcars2, cor)
v.outer(mtcars2, cor, method = "spearman")
v.outer(mtcars2, pooled.sd)
v.outer(mtcars2, euc.dist)
v.outer(mtcars2, sum2)

wc3 <- function(x, y) sum(sapply(list(x, y), wc, byrow = FALSE))
L1 <- word_list(DATA$state, DATA$person)$cwl
v.outer(L1, wc3)

## End(Not run)

```

wfm

*Word Frequency Matrix***Description**

wfm - Generate a word frequency matrix by grouping variable(s).

wfdf - Generate a word frequency data frame by grouping variable.

wfm.expanded - Expand a word frequency matrix to have multiple rows for each word.

wf.combine - Combines words (rows) of a word frequency data frame (wfdf) together.

Usage

```
wfm(text.var = NULL, grouping.var = NULL, wfdf = NULL,
    output = "raw", stopwords = NULL, digits = 2)
```

```
wfdf(text.var, grouping.var = NULL, stopwords = NULL,
    margins = FALSE, output = "raw", digits = 2)
```

```
wfm.expanded(text.var, grouping.var = NULL, ...)
```

```
wf.combine(wf.obj, word.lists, matrix = FALSE)
```

Arguments

text.var The text variable

grouping.var The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.

wfdf	A word frequency data frame given instead of raw text.var and optional grouping.var. Basically converts a word frequency dataframe (wfdf) to a word frequency matrix (wfm). Default is NULL.
output	Output type (either "proportion", "proportion" or "percent").
stopwords	A vector of stop words to remove.
digits	An integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed
margins	logical. If TRUE provides grouping.var and word variable totals.
...	Other arguments supplied to wfm.
wf.obj	A wfm or wfdf object.
word.lists	A list of character vectors of words to pass to wf.combine
matrix	logical. If TRUE returns the output as a wfm rather than a wfdf object

Value

wfm - returns a word frequency of the class matrix.

wfdf - returns a word frequency of the class data.frame with a words column and optional margin sums.

wfm.expanded - returns a matrix similar to a word frequency matrix (wfm) but the rows are expanded to represent the maximum usages of the word and cells are dummy coded to indicate that number of uses.

wf.combine - returns a word frequency matrix (wfm) or dataframe (wfdf) with counts for the combined word.lists merged and remaining terms(else).

Examples

```
## Not run:
#word frequency matrix (wfm) example:
with(DATA, wfm(state, list(sex, adult)))
dat <- with(DATA, wfm(state, person))

#word frequency dataframe (wfdf) example:
with(DATA, wfdf(state, list(sex, adult)))
with(DATA, wfdf(state, person))

#wfm.expanded example:
z <- wfm(DATA$state, DATA$person)
wfm.expanded(z)
wfm.expanded(DATA$state, DATA$person)
wfm.expanded(DATA$state, list(DATA$sex, DATA$adult))

#wf.combine example:
#raw no margins (will work)
x <- wfm(DATA$state, DATA$person)

#raw with margin (will work)
y <- wfdf(DATA$state, DATA$person, margins = TRUE)

#porportion (will not work)
z <- wfdf(DATA$state, DATA$person, output = "proportion")

WL1 <- c(y[, 1])
```

```

WL2 <- list(c("read", "the", "a"), c("you", "your", "your're"))
WL3 <- list(bob = c("read", "the", "a"), yous = c("you", "your", "your're"))
WL4 <- list(bob = c("read", "the", "a"), yous = c("a", "you", "your", "your're"))
WL5 <- list(yous = c("you", "your", "your're"))
WL6 <- list(c("you", "your", "your're")) #no name so will be called words 1
WL7 <- c("you", "your", "your're")

wf.combine(z, WL2) #Won't work not a raw frequency matrix
wf.combine(x, WL2) #Works (raw and no margins)
wf.combine(y, WL2) #Works (raw with margins)
wf.combine(y, c("you", "your", "your're"))
wf.combine(y, WL1)
wf.combine(y, WL3)
wf.combine(y, WL4) #Error b/c there's overlapping words in the word lists
wf.combine(y, WL5)
wf.combine(y, WL6)
wf.combine(y, WL7)

worlis <- c("you", "it", "it's", "no", "not", "we")
y <- wfdf(DATA$state, list(DATA$sex, DATA$adult), margins = TRUE)
z <- wf.combine(y, worlis, matrix = TRUE)

chisq.test(z)
chisq.test(wfm(wfdf = y))

## End(Not run)

```

word.associate

Find Associated Words.

Description

Find words associated with a given word(s) or a phrase(s). Results can be output as a network graph and/or wordcloud.

Usage

```

word.associate(text.var, grouping.var = NULL,
  match.string, text.unit = "sentence",
  extra.terms = NULL, target.exclude = NULL,
  stopwords = NULL, network.plot = FALSE,
  wordcloud = FALSE, cloud.colors = c("black", "gray55"),
  title.color = "blue", nw.label.cex = 0.8,
  title.padj = -4.5, nw.label.colors = NULL,
  nw.layout = NULL, nw.edge.color = "gray90",
  nw.label.proportional = TRUE, nw.title.padj = NULL,
  nw.title.location = NULL, title.font = NULL,
  title.cex = NULL, nw.edge.curved = TRUE,
  cloud.legend = NULL, cloud.legend.cex = 0.8,
  cloud.legend.location = c(-0.03, 1.03),
  nw.legend = NULL, nw.legend.cex = 0.8,
  nw.legend.location = c(-1.54, 1.41),
  legend.override = FALSE, char2space = NULL, ...)

```


Arguments

<code>text.var</code>	The text variable.
<code>grouping.var</code>	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
<code>match.string</code>	A list of vectors or vector of terms to associate in the text.
<code>text.unit</code>	The text unit (either "sentence" or "tot". This argument determines what unit to find the match string words within. For example if "sentence" is chosen the function pulls all text for sentences the match string terms are found in.
<code>extra.terms</code>	Other terms to color beyond the match string.
<code>target.exclude</code>	A vector of words to exclude from the <code>match.string</code> .
<code>stopwords</code>	Words to exclude from the analysis.
<code>network.plot</code>	logical. If TRUE plots a network plot of the words.
<code>wordcloud</code>	logical. If TRUE plots a wordcloud plot of the words.
<code>cloud.colors</code>	A vector of colors equal to the length of <code>match.string</code> +1.
<code>title.color</code>	A character vector of length one corresponding to the color of the title.
<code>nw.label.cex</code>	The magnification to be used for network plot labels relative to the current setting of <code>cex</code> . Default is .8.
<code>title.padj</code>	Adjustment for the title. For strings parallel to the axes, <code>padj</code> = 0 means right or top alignment, and <code>padj</code> = 1 means left or bottom alignment.
<code>nw.label.colors</code>	A vector of colors equal to the length of <code>match.string</code> +1.
<code>nw.layout</code>	layout types supported by <code>igraph</code> . See layout .
<code>nw.edge.color</code>	A character vector of length one corresponding to the color of the plot edges.
<code>nw.label.proportional</code>	logical. If TRUE scales the network plots across <code>grouping.var</code> to allow plot to plot comparisons.
<code>nw.title.padj</code>	Adjustment for the network plot title. For strings parallel to the axes, <code>padj</code> = 0 means right or top alignment, and <code>padj</code> = 1 means left or bottom alignment.
<code>nw.title.location</code>	On which side of the network plot (1=bottom, 2=left, 3=top, 4=right).
<code>title.font</code>	The font family of the cloud title.
<code>title.cex</code>	Character expansion factor for the title. NULL and NA are equivalent to 1.0.
<code>nw.edge.curved</code>	logical. If TRUE edges will be curved rather than straight paths.
<code>cloud.legend</code>	A character vector of names corresponding to the number of vectors in <code>match.string</code> . Both <code>nw.legend</code> and <code>cloud.legend</code> can be set separately; or one may be set and by default the other will assume those legend labels. If the user does not desire this behavior use the <code>legend.override</code> argument.
<code>cloud.legend.cex</code>	Character expansion factor for the wordcloud legend. NULL and NA are equivalent to 1.0.
<code>cloud.legend.location</code>	The x and y co-ordinates to be used to position the wordcloud legend. The location may also be specified by setting x to a single keyword from the list "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright", "right" and "center". This places the legend on the inside of the plot frame at the given location.

<code>nw.legend</code>	A character vector of names corresponding to the number of vectors in <code>match.string</code> . Both <code>nw.legend</code> and <code>cloud.legend</code> can be set separately; or one may be set and by default the other will assume those legend labels. If the user does not desire this behavior use the <code>legend.override</code> argument.
<code>nw.legend.cex</code>	Character expansion factor for the network plot legend. NULL and NA are equivalent to 1.0.
<code>nw.legend.location</code>	The x and y co-ordinates to be used to position the network plot legend. The location may also be specified by setting x to a single keyword from the list "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright", "right" and "center". This places the legend on the inside of the plot frame at the given location.
<code>legend.override</code>	By default if legend labels are supplied to either <code>cloud.legend</code> or <code>nw.legend</code> may be set and if the other remains NULL it will assume the supplied vector to the previous legend argument. If this behavior is not desired <code>legend.override</code> should be set to TRUE.
<code>char2space</code>	Currently a rode to no where. Eventually this will allow the retention of characters as is allowed in <code>trans.cloud</code> already.
<code>...</code>	Other arguments supplied to trans.cloud .

Value

Returns a list:

`word frequency matrices`

Word frequency matrices for each grouping variable.

`dialogue`

A list of dataframes for each word list (each vector supplied to `match.string`) and a final dataframe of all combined text units that contain any match string.

`match.terms`

A list of vectors of word lists (each vector supplied to `match.string`).

Optionally, returns a word cloud and/or a network plot of the text unit containing the `match.string` terms.

See Also

[trans.cloud](#), [word.network.plot](#), [wordcloud](#), [graph.adjacency](#)

Examples

```
## Not run:
ms <- c(" I", "you")
et <- c(" it", " no")
word.associate(DATA2$state, DATA2$person, match.string = ms,
  wordcloud = TRUE, proportional = TRUE,
  network.plot = TRUE, nw.label.proportional = TRUE, extra.terms = et,
  cloud.legend = c("A", "B", "C", "D"),
  title.color = "blue", cloud.colors = c("red", "blue", "purple", "gray70"))

#=====
#Note: You don't have to name the vectors in the lists but I do for clarity
ms <- list(
  list1 = c(" I ", " you"),
```

```

    list2 = c(" wh")
  )

  et <- list(
    B = c(" the", " on"),
    C = c(" it", " no")
  )

word.associate(DATA2$state, DATA2$person, match.string = ms,
  wordcloud = TRUE, proportional = TRUE,
  network.plot = TRUE, nw.label.proportional = TRUE, extra.terms = et,
  cloud.legend = c("A", "B", "C", "D"),
  title.color = "blue", cloud.colors = c("red", "blue", "purple", "gray70"))

word.associate(DATA2$state, list(DATA2$day, DATA2$person), match.string = ms)

#####
m <- list(
  A1 = c("you", "in"), #list 1
  A2 = c(" wh")        #list 2
)

n <- list(
  B = c(" the", " on"),
  C = c(" it", " no")
)

word.associate(DATA2$state, list(DATA2$day, DATA2$person), match.string = m)
word.associate(raj.act.1$dialogue, list(raj.act.1$person), match.string = m)
(out <- with(mraja1spl, word.associate(dialogue, list(fam.aff, sex), match.string = m)))
names(out)
lapply(out$dialogue, htruncdf, n = 20, w = 20)
out$cap.f

## End(Not run)

```

word.count

*Word Counts***Description**

word.count - Transcript Apply Word Counts

character.count - Transcript Apply Character Counts

character.table - Computes a table of character counts by grouping variable(s).

Usage

```

word.count(text.var, byrow = TRUE, missing = NA,
  digit.remove = TRUE, names = FALSE)

wc(text.var, byrow = TRUE, missing = NA,
  digit.remove = TRUE, names = FALSE)

```

```
character.count(text.var, byrow = TRUE, missing = NA,
  apostrophe.remove = TRUE, digit.remove = TRUE,
  count.space = FALSE)
```

```
character.table(text.var, grouping.var)
```

```
char.table(text.var, grouping.var)
```

Arguments

<code>text.var</code>	The text variable
<code>byrow</code>	logical. If TRUE counts by row, if FALSE counts all words.
<code>missing</code>	Value to insert for missing values (empty cells).
<code>digit.remove</code>	logical. If TRUE removes digits before counting words.
<code>names</code>	logical. If TRUE the sentences are given as the names of the counts.
<code>apostrophe.remove</code>	= TRUE logical. If TRUE apostrophes will be counted in the character count.
<code>count.space</code>	logical. If TRUE spaces are counted as characters.

Value

`word.count` - returns a word count by row or total.

`character.count` - returns a character count by row or total.

`character.table` - returns a dataframe of character counts by grouping variable.

Note

`wc` is a convenient short hand for `word.count`.

See Also

[syllable.count](#)

Examples

```
## Not run:
# WORD COUNT
word.count(DATA$state)
wc(DATA$state)
word.count(DATA$state, names = TRUE)
word.count(DATA$state, byrow=FALSE, names = TRUE)
sum(word.count(DATA$state))

# CHARACTER COUNTS
character.count(DATA$state)
character.count(DATA$state, byrow=FALSE)
sum(character.count(DATA$state))

library(ggplot2)
library(reshape2)
dat <- character.table(DATA$state, list(DATA$sex, DATA$adult))
(dat2 <- colsplit2df(melt(dat), keep.orig = TRUE))
head(dat2)
```

```

dat3 <- dat2[rep(seq_len(dim(dat2)[1]), dat2[, 5]), -5]

ggplot(data = dat2, aes(y = variable, x = value, colour=sex)) +
  facet_grid(adult~.) +
  geom_line(size=1, aes(group =variable), colour = "black") +
  geom_point()

ggplot(data = dat3, aes(x = variable, fill = variable)) +
  geom_bar() +
  facet_grid(sex ~ adult, margins = TRUE) +
  theme(legend.position="none")

# CHARACTER TABLE
character.table(DATA$state, DATA$person)
char.table(DATA$state, DATA$person)
character.table(DATA$state, list(DATA$sex, DATA$adult))
colsplit2df(character.table(DATA$state, list(DATA$sex, DATA$adult)))

## End(Not run)

```

word.network.plot

*Word Network Plot***Description**

A network plot of words. Shows the interconnected and supporting use of words between textual units containing key terms.

Usage

```

word.network.plot(text.var, grouping.var = NULL,
  target.words = NULL, stopwords = Top100Words,
  label.cex = 0.8, label.size = 0.5, edge.curved = TRUE,
  vertex.shape = "circle", edge.color = "gray70",
  label.colors = "black", layout = NULL,
  title.name = NULL, title.padj = -4.5,
  title.location = 3, title.font = NULL, title.cex = 0.8,
  log.labels = FALSE, title.color = "black",
  legend = NULL, legend.cex = 0.8,
  legend.location = c(-1.54, 1.41), plot = TRUE)

```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
target.words	A named list of vectors of words whose length corresponds to label.colors (+1 length in cloud colors for non matched terms).
stopwords	Words to exclude from the analysis (default is Top100Words).
label.cex	The magnification to be used for network plot labels relative to the current setting of cex. Default is .8.

log.labels	logical. If TRUE uses a proportional log label for more readable labels. The formula is: $\log(\text{SUMS})/\max(\log(\text{SUMS}))$. label.size adds more control over the label sizes.
label.size	An optional sizing constant to add to labels if log.labels is TRUE.
edge.curved	logical. If TRUE edges will be curved rather than straight paths.
vertex.shape	The shape of the vertices (see igraph.vertex.shapes for more).
edge.color	A character vector of length one corresponding to the color of the plot edges.
label.colors	A character vector of length one corresponding to the color of the labels.
layout	layout types supported by igraph. See layout .
title.name	The title of the plot.
title.padj	Adjustment for the network plot title. For strings parallel to the axes, padj = 0 means right or top alignment, and padj = 1 means left or bottom alignment.
title.location	On which side of the network plot (1=bottom, 2=left, 3=top, 4=right).
title.font	The font family of the cloud title.
title.cex	Character expansion factor for the title. NULL and NA are equivalent to 1.0.
title.color	A character vector of length one corresponding to the color of the title.
legend	A character vector of names corresponding to the number of vectors in match.string.
legend.cex	Character expansion factor for the network plot legend. NULL and NA are equivalent to 1.0.
legend.location	The x and y co-ordinates to be used to position the network plot legend. The location may also be specified by setting x to a single keyword from the list "bottomright", "bottom", "bottomleft", "left", "topleft", "top", "topright", "right" and "center". This places the legend on the inside of the plot frame at the given location.
plot	logical. If TRUE plots a network plot of the words.

Value

Silently returns a list of igraph parameters. Optionally, plots the output.

See Also

[word.network.plot](#), [graph.adjacency](#)

Examples

```
## Not run:
word.network.plot(text.var=DATA$state, grouping.var=DATA$person)
word.network.plot(text.var=DATA$state, grouping.var=list(DATA$sex,
DATA$adult))
word.network.plot(text.var=DATA$state, grouping.var=DATA$person,
  title.name = "TITLE", log.labels=TRUE)
word.network.plot(text.var=raj.act.1, grouping.var=raj.act.1$person,
  stopwords = Top200Words)

## End(Not run)
```

word_diff_list	<i>Differences In Word Use Between Groups</i>
----------------	---

Description

Look at the differences in word uses between grouping variable(s). Look at all possible "a" vs. "b" combinations or "a" vs. all others.

Usage

```
word_diff_list(text.var, grouping.var, vs.all = FALSE,
               vs.all.cut = 1, stopwords = NULL, alphabetical = FALSE,
               digits = 2)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Takes a single grouping variable or a list of 1 or more grouping variables.
vs.all	logical. If TRUE looks at each grouping variable against all others ("a" vs. all comparison). If FALSE looks at each "a" vs. "b", comparison (e.g. for groups "a", "b", and "c"; "a" vs. "b", "a" vs. "c" and "b" vs. "c" will be considered).
vs.all.cut	If vs.all.cut = TRUE this argument controls the number of other groups that may share a word (default is 1).
stopwords	A vector of stop words to remove.
alphabetical	logical. If TRUE orders the word lists alphabetized by word. If FALSE order first by frequency and then by word.
digits	the number of digits to be displayed in the proportion column (default is 3).

Value

An list of word data frames comparing grouping variables word use against one another. Each data frame contains three columns:

word	The words unique to that group
freq	The number of times that group used that word
prop	The proportion of that group's over all word use dedicated to that particular word

Examples

```
## Not run:
with(DATA, word_diff_list(text.var = state, grouping.var = list(sex, adult)))
with(DATA, word_diff_list(state, person))
with(DATA, word_diff_list(state, grouping.var = list(sex, adult),
                          vs.all=TRUE, vs.all.cut=2))

with(mrja1, word_diff_list(text.var = dialogue,
                           grouping.var = list(mrja1$sex, mrja1$fam.aff)))
word_diff_list(mrja1$dialogue, mrja1$person)
word_diff_list(mrja1$dialogue, mrja1$fam.aff, stopwords = Top25Words)
```

```
word_diff_list(mraja1$dialogue, mraja1$fam.aff, vs.all=TRUE, vs.all.cut=2)

## End(Not run)
```

word_list

Raw Word Lists/Frequency Counts

Description

Transcript Apply Raw Word Lists and Frequency Counts by grouping variable(s).

Usage

```
word_list(text.var, grouping.var = NULL,
  stopwords = NULL, alphabetical = FALSE, cut.n = 20,
  cap = TRUE, cap.list = NULL, cap.I = TRUE,
  rm.bracket = TRUE, char.keep = NULL,
  apostrophe.remove = FALSE, ...)
```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one word list for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
stopwords	A vector of stop words to remove.
cut.n	Cut off point for reduced frequency stop word list (rfswl).
cap	logical. If TRUE capitalizes words from the cap.list.
cap.list	Vector of words to capitalize.
cap.I	logical. If TRUE capitalizes words containing the personal pronoun I.

Value

An object of class "word.list" is a list containing at the following components:

cwl	complete word list; raw words
swl	stop word list; same as rwl with stop words removed
fwl	frequency word list; a data frame of words and corresponding frequency counts
fswl	frequency stopword word list; same as fwl but with stopwords removed
rfswl	reduced frequency stopword word list; same as fswl but truncated to n rows

Examples

```
## Not run:
XX <- word_list(raj.act.1$dialogue)
names(XX)
XX$cwl
XX$swl
XX$fwl
XX$fswl
XX$rfswl
```



```

with(raj, word_list(text.var = dialogue, grouping.var = list(person, act)))
with(DATA, word_list(state, person))
with(DATA, word_list(state, person, stopwords = Top25Words))
with(DATA, word_list(state, person, cap = FALSE, cap.list=c("do", "we")))

## End(Not run)

```

word_stats

*Descriptive Word Statistics***Description**

Transcript apply descriptive word statistics.

Usage

```

word_stats(text.var, grouping.var = NULL, tot = NULL,
  parallel = FALSE, rm.incomplete = FALSE,
  digit.remove = FALSE, apostrophe.remove = FALSE,
  digits = 3, ...)

```

Arguments

text.var	The text variable.
grouping.var	The grouping variables. Default NULL generates one output for all text. Also takes a single grouping variable or a list of 1 or more grouping variables.
tot	Optional toutn of talk variable.
parallel	logical. If TRUE attempts to run the function on multiple cores. Note that this may not mean a speed boost if you have one core or if the data set is smaller as the cluster takes time to create (parallel is slower until approximately 10,000 rows).
rm.incomplete	logical. If TRUE incomplete statments are removed from calculating the output.
digit.remove	logical. If TRUE removes digits from calculating the output.
apostrophe.remove	logical. If TRUE removes apostrophes from calculating the output.
digits	Integer; number of decimal places to round.
...	Any other arguments passed to endf

Value

Returns a list of three descriptive word statistics:

ts	A data frame of descriptive word statistics by row
gts	A data frame of word statistics per grouping variable: <ul style="list-style-type: none"> • n.tot - number of turns of talk • n.sent - number of sentences • n.words - number of words • n.char - number of characters

- n.syl - number of syllables
- n.poly - number of polysyllables
- sptot - syllables per turn of talk
- wps - words per sentence
- cps - characters per sentence
- sps - syllables per sentence
- psps - polysyllables per sentence
- cpw - characters per word
- spw - syllables per word
- n.state - number of statements
- n.quest - number of questions
- n.exclm - number of exclamations
- n.incom - number of incomplete statements
- n.hapax - number of hapax legomenon
- n.dis - number of dis legomenon
- grow.rate - proportion of hapax legomenon to words
- prop.dis - proportion of dis legomenon to words

mpun An account of sentences with improper end mark

Examples

```
## Not run:
word_stats(mraja1spl$dialogue, mraja1spl$person)
(desc_wrds <- with(mraja1spl, word_stats(dialogue, person, tot = tot)))
names(desc_wrds)
desc_wrds$ts
desc_wrds$gts
desc_wrds$pun
with(mraja1spl, word_stats(dialogue, list(sex, died, fam.aff)))

## End(Not run)
```

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