Package 'JATSdecoder'

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Title Extract meta data, text and study characteristics from NISO-JATS coded XML files

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Description Provides a function collection to extract metadata, sectioned text and study characteristics from scientific articles. JATSdecoder() converts NISO-JATS-tagged XML files to a structured list with elements containing title, author, journal, history, link, abstract, sectioned text and references. Articles in PDF format can be easily converted to NISO-JATS with the open source software CER-MINE (<https://github.com/CeON/CERMINE>). study.character() extracts multiple study characteristics like number of included studies, statistical methods used, alpha error, power, statistical results, correction method for multiple testing, software used. An estimation of the involved sample size is performed based on reports within the abstract and the reported degrees of freedom within statistical results. In addition, the package contains some useful functions to process text (text2sentences(), text2num(), ngram(), strsplit2(), grep2()).

Depends R (>= 3.1.1)

Imports utils, stats, NLP, openNLP

License GPL-3

URL <https://github.com/ingmarboeschen/JATSdecoder>

Language en-US

Encoding UTF-8

LazyData true

RoxygenNote 7.2.1

R topics documented:

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allStats	*allStats*

Description

Extracts any statistical results from text string with some uniformizations.

Usage

```
allStats(x)
```

Arguments

x   a text string to extract statistical results from.

Examples

```
x<-c("The mean difference of scale A was significant (beta=12.9, t(18)=2.5, p<.05)",
"The ANOVA yielded significant results on factor A (F(2,18)=6, p<.05, eta(g)2<-.22).",
"The correlation of x and y was r=.37.")
allStats(x)
```

---

est.ss	*est.ss*

Description

Function to estimate studies sample size by maximizing different conservative estimates. Performs four different extraction heuristics for sample sizes mentioned in abstract, text and statistical results.

Usage

```
est.ss(
  abstract = NULL,
  text = NULL,
  stats = NULL,
  standardStats = NULL,
  quantileDF = 0.9,
  max.only = FALSE,
  max.parts = TRUE
)
```

Arguments

abstract : an abstract text string.

text : the main text string to process (usually method and result sections). If text has content, arguments "stats" and "standardStats" are deactivated and filled with results by get.stats(text).

stats : statistics extracted with get.stats(x)$stats (only active if no text is submitted).

standardStats : standard statistics extracted with get.stats(x)$standardStats (only active if no text is submitted).

quantileDF : quantile of (df1-1)+(df2+2) to extract.

max.only : Logical. If TRUE only the final estimate will be returned, if FALSE all sub estimates are returned as well.

max.parts : Logical. If FALSE outputs all captured sample sizes in sub inputs.

Details

Sample size extraction from abstract:
- Extracts N= from abstract text and performs POS search with list of synonyms of sample units

Sample size extraction from text:
- Unifies and extracts textlines with age descriptions, than computes sum of hits as nage - Unifies and extracts all "numeric male-female" patterns than computes sum of first male/female hit - Unifies and extracts textlines with participant description than computes sum of first three hits as ntext

Sample size extraction from statistical results:
- Extracts "N=" in statistical results extracted with allStats() that contain p-value: e.g.: chi(2, N=12)=15.2, p<.05

Sample size extraction by degrees of freedom with result of standardStats(allStats()):
- Extracts df1 and df2 if possible and neither containing a ".", than calculates specified quantile of (df1+1)+(df2+2) (at least 2 group comparison assumed)

Examples

```
a<-"One hundred twelve students participated in our study."
x<-"Our sample consists of three hundred twenty five undergraduate students.
    The F-test indicates significant differences in means F(2,102)=3.21, p<.05."
est.ss(abstract=a,text=x)
```

---

get.abstract *get.abstract*

Description

Extracts abstract tag from NISO-JATS coded XML file or text as vector of abstracts.

Usage

```
get.abstract(
  x,
  sentences = FALSE,
  remove.title = TRUE,
  letter.convert = TRUE,
  cermine = FALSE
)
```

Arguments

x	a NISO-JATS coded XML file or text.
sentences	Logical. If TRUE abstract is returned as vector of sentences.
remove.title	Logical. If TRUE removes section titles in abstract.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.
cermine	Logical. If TRUE and if 'letter.convert=TRUE' CERMINE specific letter correction is carried out (e.g. inserting of missing operators to statistical results).

Examples

```
x<-"Some text <abstract>Some abstract</abstract> some text"
get.abstract(x)
x<-"Some text <abstract>Some abstract</abstract> TEXT <abstract with subsettings>
Some other abstract</abstract> Some text "
get.abstract(x)
```

---

get.aff	*get.aff*

Description

Extracts the affiliation tag information from NISO-JATS coded XML file or text as a vector of affiliations.

Usage

```
get.aff(x, remove.html = FALSE, letter.convert = TRUE)
```

Arguments

x	a NISO-JATS coded XML file or text.
remove.html	Logical. If TRUE removes all html tags.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.

Examples

```
x<-"Some text <aff>Some affiliation</aff> some text"
get.aff(x)
x<-"TEXT <aff>Some affiliation</aff> TEXT <aff>Some other affiliation</aff> TEXT"
get.aff(x)
```

---

get.alpha.error   *get.alpha.error*

Description

Extracts reported and corrected alpha error from text and 1-alpha confidence intervalls.

Usage

```
get.alpha.error(x, p2alpha = FALSE, output = "list")
```

Arguments

x	text string to process.
p2alpha	Logical. If TRUE detects and extracts alpha errors denoted with critical p-value (what may lead to some false positive detections).
output	One of c("list","vector"). If output="list" returns a list containing: alpha_error, corrected_alpha, alpha_from_CI, alpha_max, alpha_min. If output="vector" returns unique alpha errors but no distinction of types.

Examples

```
x<-c("The threshold for significance was adjusted to .05/2",
"Type 1 error rate was alpha=.05.")
get.alpha.error(x)
x<-c("We used p<.05 as level of significance.",
     "We display .95 CIs and use an adjusted alpha of .10/3.",
     "The effect was significant with p<.025.")
get.alpha.error(x)
```

---

get.assumptions   *get.assumptions*

Description

Extracts the mentioned statistical assumptions from a text string by a dictionary search of 22 common statistical assumptions.

Usage

```
get.assumptions(x, hits_only = TRUE)
```

Arguments

x	text string to process.
hits_only	Logical. If TRUE returns the detected assumtions only, else a hit matrix with all potential assumptions is returned.

Examples

```
x<-"Sphericity assumption and gaus-marcov was violated."
get.assumptions(x)
```

---

get.author   *get.author*

Description

Extracts author tag information from NISO-JATS coded XML file or text.

Usage

```
get.author(x, paste = "", short.names = FALSE, letter.convert = FALSE)
```

Arguments

x	a NISO-JATS coded XML file or text.
paste	if paste!="" author list is collapsed to one cell with seperator specified (e.g. paste=";").
short.names	Logical. If TRUE fully available first names will be reduced to single letter abbreviation.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.

---

get.category   *get.category*

Description

Extracts category tag/s from NISO-JATS coded XML file or text as vector of categories.

Usage

```
get.category(x)
```

Arguments

x   a NISO-JATS coded XML file or text.

Examples

```
x<-"Some text <article-categories>Some category</article-categories> some text"
get.category(x)
```

---

get.contrib   *get.contrib*

Description

Extracts contrib tag/s from NISO-JATS coded XML file or text as vector of contributers.

Usage

```
get.contrib(x, remove.html = FALSE, letter.convert = FALSE)
```

Arguments

x   a NISO-JATS coded XML file or text.

remove.html   Logical. If TRUE removes all HTML tags.

letter.convert   Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.

---

get.country   *get.country*

Description

Extracts country tag from NISO-JATS coded XML file or text as vector of unique countries.

Usage

```
get.country(x, unifyCountry = TRUE)
```

Arguments

x   a NISO-JATS coded XML file or text.

unifyCountry   Logical. If TRUE replaces country name with standardised country name.

Examples

```
x<-"Some text <country>UK</country> some text <country>England</country>
    Text<country>Berlin, Germany</country>"
get.country(x)
```

---

get.doi   *get.doi*

Description

Extracts articles doi from NISO-JATS coded XML file or text.

Usage

```
get.doi(x)
```

Arguments

x   a NISO-JATS coded XML file or text.

get.editor	*get.editor*

Description

Extracts editor tag from NISO-JATS coded XML file or text as vector of editors.

Usage

```
get.editor(x, role = FALSE, short.names = FALSE, letter.convert = FALSE)
```

Arguments

x	a NISO-JATS coded XML file or text.
role	Logical. If TRUE adds role to editor name, if available.
short.names	Logical. If TRUE reduces fully available first names to one letter abbreviation.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.

get.history	*get.history*

Description

Extracts available publishing history tags from NISO-JATS coded XML file or text and compute pubDate and pubyear.

Usage

```
get.history(x, remove.na = FALSE)
```

Arguments

x	a NISO-JATS coded XML file or text.
remove.na	Logical. If TRUE hides non available date stamps.

---

get.journal *get.journal*

Description

Extracts journal tag from NISO-JATS coded XML file or text.

Usage

```
get.journal(x)
```

Arguments

x   a NISO-JATS coded XML file or text.

Examples

```
x<-"Some text <journal-title>PLoS One</journal-title> some text"
get.journal(x)
```

---

get.keywords *get.keywords*

Description

Extracts keyword tag/s from NISO-JATS coded XML file or text as vector of keywords.

Usage

```
get.keywords(
  x,
  paste = "",
  letter.convert = TRUE,
  include.max = length(keyword)
)
```

Arguments

x   a NISO-JATS coded XML file or text.

paste   if paste!="" keyword list is collapsed to one cell with seperator specified (e.g.
                paste=";").

letter.convert Logical. If TRUE converts hexadecimal and HTML coded characters to Uni-
                code.

include.max   a maximum number of keywords to extract.

Examples

```
x<-"Some text <kwd>Keyword 1</kwd>, <kwd>Keyword 2</kwd> some text"
get.keywords(x)
get.keywords(x,paste(", "))
```

---

get.method   *get.method*

Description

Extracts statistical methods mentioned in text.

Usage

```
get.method(x, add = NULL, cermine = FALSE)
```

Arguments

x	text to extract statistical methods from.
add	possible new end words of method as vector.
cermine	Logical. If TRUE CERMINE specific letter conversion will be performed.

Examples

```
x<-"We used multiple regression analysis and
two sample t tests to evaluate our results."
get.method(x)
```

---

get.multi.comparison   *get.multi.comparison*

Description

Extracts alpha-/p-value correction method for multiple comparisons from list with 15 correction methods.

Usage

```
get.multi.comparison(x)
```

Arguments

x	text string to process.

Examples

```
x<-"We used Bonferroni corrected p-values."
get.multi.comparison(x)
```

---

get.n.studies   *get.n.studies*

Description

Extracts number of studies/experiments from text.

Usage

```
get.n.studies(x, tolower = TRUE)
```

Arguments

x   text string to process.

tolower   Logical. If TRUE lowerises text and search patterns for processing.

---

get.outlier.def   *get.outlier.def*

Description

Extracts outlier/extreme value definition/removal in standard deviations, if present in text.

Usage

```
get.outlier.def(x)
```

Arguments

x   text string to process.

Examples

```
x<-"We removed 4 extreme values that were 3 SD above mean."
get.outlier.def(x)
```

---

get.power   *get.power*

Description

Extracts a priori power and empirial power values from text.

Usage

```
get.power(x)
```

Arguments

x   text string to process.

Examples

```
x<-"We used G*Power 3 to calculate the needed sample with
beta error rate set to 12% and alpha error to .05."
get.power(x)
```

---

get.R.package   *get.R.package*

Description

Extracts mentioned R packages from text.

Usage

```
get.R.package(x, update.package.list = FALSE)
```

Arguments

x   text string to process.

update.package.list

           Logical. If TRUE update of list with available packages is downloaded from CRAN with available.packages().

Examples

```
get.R.package("We used the R Software packages lme4 (and psych).")
```

get.references	*get.references*

Description

Extracts reference list from NISO-JATS coded XML file or text as vector of references.

Usage

```
get.references(
  x,
  letter.convert = FALSE,
  remove.html = FALSE,
  extract = "full"
)
```

Arguments

x	a NISO-JATS coded XML file or text.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.
remove.html	Logical. If TRUE removes all HTML tags.
extract	part of refernces to extract (one of "full" or "title").

get.sentence.with.pattern	
	get.sentence.with.pattern

Description

Returns lines with search term patterns.

Usage

```
get.sentence.with.pattern(x, patterns = c(""), tolower = TRUE)
```

Arguments

x	sentence vector to process.
patterns	search terms.
tolower	Logical. If TRUE converts search terms and text to lower case.

Examples

```
text<-c("This demo", "demonstrates how", "get.sentence.with.pattern works.")
get.sentence.with.pattern(text,c("Demo","example","work"))
get.sentence.with.pattern(text,c("Demo","example","work"),tolower=FALSE)
```

---

get.sig.adjectives   *get.sig.adjectives*

Description

Extracts adjectives used for in/significance out of list with 37 potential adjectives.

Usage

```
get.sig.adjectives(x, unique_only = FALSE)
```

Arguments

x   text string to process.

unique_only   Logical. If TRUE returns unique hits only.

Examples

```
get.sig.adjectives(
 x<-"We found very highly significance for type 1 effect"
)
```

---

get.software   *get.software*

Description

Extracts mentioned software from text by dictionary search for 63 software names (object: .software_names).

Usage

```
get.software(x, add.software = NULL)
```

Arguments

x   text string to process.

add.software   a text vector with additional software name patterns to search for.

Examples

```
get.software("We used the R Software and Excel 4.0 to analyse our data.")
```

---

get.stats   *get.stats*

Description

Extracts statistical results from text string, XML, CERMXML, HTML or DOCX files. The result is a list with a vector containing all identified sticked results and a matrix containing the reported standard statistics and recalculated p-values if computation is possible.

Usage

```
get.stats(
  x,
  output = "both",
  stats.mode = "all",
  recalculate.p = TRUE,
  alternative = "undirected",
  estimateZ = FALSE,
  T2t = FALSE,
  R2r = FALSE,
  select = NULL,
  rm.na.col = TRUE,
  cermine = FALSE
)
```

Arguments

x	DOCX file path, NISO-JATS coded XML file path or plain textual content
output	Select the desired output. One of c("both", "allStats", "standardStats").
stats.mode	Select subset of standardStats. One of: c("all", "checkable", "computable", "uncomputable").
recalculate.p	Logical. If TRUE recalculates p-values of standardStats if possible.
alternative	Character. Select sidedness of recomputed p-values from t-, r- and beta-values. One of c("undirected", "directed", "both").
estimateZ	Logical. If TRUE detected beta-/d-value is divided by reported standard error "SE" to estimate Z-value ("Zest") for observed beta/d and recompute p-value. Note: This is only valid, if Gauss-Marcov assumptions are met and a sufficiently large sample size is used. If a Z- or t-value is detected in a report of a beta-/d-coefficient with SE, no estimation will be performed, although set to TRUE.
T2t	Logical. If TRUE capital letter T is treated as t-statistic.
R2r	Logical. If TRUE capital letter R is treated as correlation.
select	Select specific standard statistics only (e.g.: c("t", "F", "Chi2")).
rm.na.col	Logical. If TRUE removes all columns with only NA from standardStats.
cermine	Logical. If TRUE CERMINE specific letter conversion will be peformed on allStats results.

Source

A minimal web application that extracts statistical results from single documents with JATSdecoder::get.stats() is hosted at: https://www.get-stats.app/

Statistical results from subsets of articles in the PubMed Central library can be analyzed and used to identify studies with specific measures and effect and sample sizes. Further, p-value checking is possible on selections of less than 20,000 articles. is hosted at: https://www.scianalyzer.com/

References

Böschen (2021). "Evaluation of JATSdecoder as an automated text extraction tool for statistical results in scientific reports." *Scientific Reports.* doi: 10.1038/s41598-021-98782-3.

Examples

```
x<-c("The mean difference of scale A was significant (beta=12.9, t(18)=2.5, p<.05).",
"The ANOVA yielded significant results on
 faktor A (F(2,18)=6, p<.05, eta(g)2<-.22)",
"the correlation of x and y was r=.37.")
get.stats(x)
```

---

get.subject   *get.subject*

Description

Extracts subject tag/s from NISO-JATS coded XML file or text as vector of subjects.

Usage

```
get.subject(x, letter.convert = TRUE, paste = "")
```

Arguments

x	a NISO-JATS coded XML file or text.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.
paste	if paste!="" subject list is collapsed to one cell with seperator specified (e.g. paste=";").

Examples

```
x<-"Some text <subject>Some subject</subject> some text"
get.subject(x)
x<-"Some text <subject>Some subject</subject> TEXT ...
<subject>Some other subject</subject> Some text "
get.subject(x)
get.subject(x,paste=", ")
```

---

get.tables *get.tables*

Description

Extracts HTML tables as vector of tables.

Usage

```
get.tables(x)
```

Arguments

x   HTML file or html text.

---

get.test.direction *get.test.direction*

Description

Extracts mentioned test direction/s (one sided, two sided, one and two sided) from text.

Usage

```
get.test.direction(x)
```

Arguments

x   text string to process.

---

get.text *get.text*

Description

Extracts main textual content from NISO-JATS coded XML file or text as sectioned text.

Usage

```
get.text(
  x,
  sectionsplit = "",
  grepsection = "",
  letter.convert = TRUE,
  greek2text = FALSE,
  sentences = FALSE,
  paragraph = FALSE,
  cermine = "auto",
  rm.table = TRUE,
  rm.formula = TRUE,
  rm.xref = TRUE,
  rm.media = TRUE,
  rm.graphic = TRUE,
  rm.ext_link = TRUE
)
```

Arguments

x	a NISO-JATS coded XML file or text.
sectionsplit	search patterns for section split (forced to lower case), e.g. c("intro", "method", "result", "discus").
grepsection	search pattern to reduce text to specific section namings only.
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.
greek2text	Logical. If TRUE some greek letters and special characters will be unified to textual representation (important to extract stats).
sentences	Logical. IF TRUE text is returned as sectioned list with sentences.
paragraph	Logical. IF TRUE "<New paragraph>" is added at the end of each paragraph to enable manual splitting at paragraphs.
cermine	Logical. If TRUE CERMINE specific error handling and letter conversion will be applied. If set to "auto" file name ending with 'cermxml$' will set cermine=TRUE.
rm.table	Logical. If TRUE removes <table> tag from text.
rm.formula	Logical. If TRUE removes <formula> tags.
rm.xref	Logical. If TRUE removes <xref> tag (citing) from text.
rm.media	Logical. If TRUE removes <media> tag from text.
rm.graphic	Logical. If TRUE removes <graphic> and <fig> tag from text.
rm.ext_link	Logical. If TRUE removes <ext link> tag from text.

---

get.title *get.title*

Description

Extracts article title from NISO-JATS coded XML file or text.

Usage

```
get.title(x)
```

Arguments

x   a NISO-JATS coded XML file or text.

---

get.type *get.type*

Description

Extracts article type from NISO-JATS coded XML file or text.

Usage

```
get.type(x)
```

Arguments

x   a NISO-JATS coded XML file or text.

---

get.vol *get.vol*

Description

Extracts volume, first and last page from NISO-JATS coded XML file or text.

Usage

```
get.vol(x)
```

Arguments

x   a NISO-JATS XML coded file or text.

---

grep2   *grep2*

Description

Extension of grep(). Allows to identify and extract cells with/without multiple search patterns that are connected with AND.

Usage

```
grep2(pattern, x, value = TRUE, invert = FALSE, perl = FALSE)
```

Arguments

pattern	Character vector containing regular expression as cells to be matched in the given character vector.
x	A character vector where matches are sought, or an object which can be coerced by as.character to a character vector. Long vectors are supported.
value	Logical. if FALSE, a vector containing the (integer) indices of the matches determined by grep2 is returned, and if TRUE, a vector containing the matching elements themselves is returned.
invert	Logical. If TRUE return indices or values for elements that do not match.
perl	Logical. Should Perl-compatible regexps be used?

Examples

```
x<-c("ab","ac","ad","bc","bad")
grep2(c("a","b"),x)
grep2(c("a","b"),x,invert=TRUE)
grep2(c("a","b"),x,value=FALSE)
```

---

has.interaction   *has.interaction*

Description

Identifies mentiones of interaction/moderator/mediator effect in text.

Usage

```
has.interaction(x)
```

Arguments

x	text string to process.

JATSdecoder   *JATSdecoder*

Description

Function to extract and restructure NISO-JATS coded XML file or text into a list with metadata and text as selectable elements. Use CERMINE to convert PDF to CERMXML files.

Usage

```
JATSdecoder(
  x,
  sectionsplit = c("intro", "method", "result", "study", "experiment", "conclu",
    "implica", "discussion"),
  grepsection = "",
  sentences = FALSE,
  paragraph = FALSE,
  abstract2sentences = TRUE,
  output = "all",
  letter.convert = TRUE,
  unify.country.name = TRUE,
  greek2text = FALSE,
  warning = TRUE,
  countryconnection = FALSE,
  authorconnection = FALSE
)
```

Arguments

x	a NISO-JATS coded XML file or text.
sectionsplit	search patterns for section split of text parts (forced to lower case), e.g. c("intro", "method", "result", "discus").
grepsection	search pattern in regex to reduce text to specific section only.
sentences	Logical. IF TRUE text is returned as sectioned list with sentences.
paragraph	Logical. IF TRUE "<New paragraph>" is added at the end of each paragraph to enable manual splitting at paragraphs.
abstract2sentences	
	Logical. IF TRUE abstract is returned as vector with sentences.
output	selection of specific results to output c("all", "title", "author", "affiliation", "journal", "volume", "editor", "doi", "type", "history", "country", "subject", "keywords", "abstract", "sections", "text", "tables", "captions", "references").
letter.convert	Logical. If TRUE converts hexadecimal and HTML coded characters to Unicode.
unify.country.name	
	Logical. If TRUE tries to unify country name/s with list of country names from worldmap().

greek2text   Logical. If TRUE converts and unifies several greek letters to textual representation, e.g.: "alpha".

warning   Logical. If TRUE outputs a warning if processing CERMINE converted PDF files.

countryconnection
                Logical. If TRUE outputs country connections as vector c("A - B","A - C", ...).

authorconnection
                Logical. If TRUE outputs connections of a maximum of 50 involved authors as vector c("A - B","A - C", ...).

Note

A short tutorial on how to work with JATSdecoder and the generated outputs can be found at: https://github.com/ingmarboeschen/JATSdecoder

Source

An interactive web application for selecting and analyzing extracted article metadata and study characteristics for articles linked to PubMed Central is hosted at: https://www.scianalyzer.com/

The XML version of PubMed Central database articles can be downloaded in bulk from: https://ftp.ncbi.nlm.nih.gov/pub/pmc/oa_bulk/

References

Böschen (2021). "Software review: The JATSdecoder package - extract metadata, abstract and sectioned text from NISO-JATS coded XML documents; Insights to PubMed Central's open access database." *Scientometrics.* doi: 10.1007/s1119202104162z.

---

letter.convert   *letter.convert*

Description

Converts and unifies most hexadecimal and some HTML coded letters to Unicode characters. Performs CERMINE specific error correction (inserting operators, where these got lost while conversion).

Usage

```
letter.convert(x, cermine = FALSE, greek2text = FALSE, warning = TRUE)
```

Arguments

x	text string to process.
cermine	Logical. If TRUE CERMINE specific error handling and letter conversion will be applied.
greek2text	Logical. If TRUE some greek letters and special characters will be unified to textual representation (important to extract stats).
warning	Logical. If TRUE prints warning massage if CERMINE specific letter conversion was performed.

Examples

```
x<-c("five &#x0003c; ten","five &lt; ten")
letter.convert(x)
```

---

ngram	*ngram*

Description

Extracts ngram bag of words around words that match a search pattern. Note: If an input contains the search pattern twice, only the ngram bag of words of the last hit is detected. Consider individual text splitting with text2sentences() or strsplit2() before applying ngram().

Usage

```
ngram(
  x,
  pattern,
  ngram = c(-3, 3),
  tolower = FALSE,
  split = FALSE,
  exact = FALSE
)
```

Arguments

x	vector of text strings to process.
pattern	a search term pattern to extract the ngram bag of words.
ngram	a vector of length=2 that defines the number of words to extract from left and right side of pattern match.
tolower	Logical. If TRUE converts text and pattern to lower case.
split	Logical. If TRUE splits text input at "[.,;:] " before processing. Note: You may consider other text splits before.
exact	Logical. If TRUE only exact word matches will be processes

Examples

```
text<-"One hundred twenty-eight students participated in our Study,
that was administred in thirteen clinics."
ngram(text,pattern="study",ngram=c(-1,2))
```

---

standardStats   *standardStats*

Description

Extracts and restructures statistical standard results like Z, t, Cohen's d, F, eta^2, r, R^2, chi^2, BF_10, Q, U, H, OR, RR, beta values to a matrix. Performs a recomputation of p-values if possible.

Usage

```
standardStats(
  x,
  stats.mode = "all",
  recalculate.p = TRUE,
  alternative = "undirected",
  estimateZ = FALSE,
  T2t = FALSE,
  R2r = FALSE,
  select = NULL,
  rm.na.col = TRUE
)
```

Arguments

x	result of allStats().
stats.mode	Select subset of standard stats. One of: c("all", "checkable", "computable", "uncomputable").
recalculate.p	Logical. If TRUE recalculates p values (for 2 sided test) if possible.
alternative	Character. Select sidedness of recomputed p-values from t-, r- and beta-values. One of c("undirected", "directed", "both").
estimateZ	Logical. If TRUE detected beta-/d-value is divided by reported standard error "SE" to estimate Z-value ("Zest") for observed beta/d and recompute p-value. Note: This is only valid, if Gauss-Marcov assumptions are met and a sufficiently large sample size is used. If a Z- or t-value is detected in a report of a beta-/d-coefficient with SE, no estimation will be performed, although set to TRUE.
T2t	Logical. If TRUE capital letter T is treated as t-statistic.
R2r	Logical. If TRUE capital letter R is treated as correlation.
select	Select specific standard statistics only (e.g.: c("t", "F", "Chi2")).
rm.na.col	Logical. If TRUE removes all columns with only NA.

Examples

```
x<-c("t(38.8)<=>1.96, p=.002","F(2,39)<=>4, p<=>.05",
"U(2)=200, p>.25","Z<=>2.1, F(20.8,22.6)=200, p<.005,
BF(01)<=>4","chi=3.2, r(34)<=>-.7, p<.01, R2=76%.")
standardStats(x)
```

---

strsplit2 *strsplit2*

Description

Extension of strsplit(). Makes it possible to split lines before or after a pattern match without removing the pattern.

Usage

```
strsplit2(x, split, type = "remove", perl = FALSE)
```

Arguments

x	text string to process.
split	pattern to split text at.
type	one out of c("remove", "before", "after").
perl	Logical. If TRUE uses perl expressions.

Examples

```
x<-"This is some text, where text is the split pattern of the text."
strsplit2(x,"text","after")
```

---

study.character *study.character*

Description

Extracts study characteristics out of a NISO-JATS coded XML file. Use CERMINE to convert PDF to CERMXML files.

Usage

```
study.character(
  x,
  stats.mode = "all",
  recalculate.p = TRUE,
  alternative = "auto",
  estimateZ = FALSE,
  T2t = FALSE,
  R2r = FALSE,
  selectStandardStats = NULL,
  p2alpha = FALSE,
  alpha_output = "list",
  captions = TRUE,
  text.mode = 1,
  update.package.list = FALSE,
  add.software = NULL,
  quantileDF = 0.9,
  N.max.only = FALSE,
  output = "all",
  rm.na.col = TRUE
)
```

Arguments

x	JATS coded XML file.
stats.mode	Character. Select subset of standard stats. One of: c("all", "checkable", "computable").
recalculate.p	Logical. If TRUE recalculates p values (for 2 sided test) if possible.
alternative	Character. Select sidedness of recomputed p-values for t-, r- and Z-values. One of c("auto", "undirected", "directed", "both"). If set to "auto" 'alternative' will be be set to 'both' if get.test.direction() detects one-directional hypotheses/tests in text. If no directional hypotheses/tests are dtected only "undirected" recomputed p-values will be returned.
estimateZ	Logical. If TRUE detected beta-/d-value is divided by reported standard error "SE" to estimate Z-value ("Zest") for observed beta/d and recompute p-value. Note: This is only valid, if Gauss-Marcov assumptions are met and a sufficiently large sample size is used. If a Z- or t-value is detected in a report of a beta-/d-coefficient with SE, no estimation will be performed, although set to TRUE.
T2t	Logical. If TRUE capital letter T is treated as t-statistic when extracting statistics with get.stats().
R2r	Logical. If TRUE capital letter R is treated as correlation when extracting statistics with get.stats().
selectStandardStats	
	Select specific standard statistics only (e.g.: c("t", "F", "Chi2")).
p2alpha	Logical. If TRUE detects and extracts alpha errors denoted with critical p-value (what may lead to some false positive detections).

alpha_output	One of c("list", "vector"). If alpha_output = "list" a list with elements: alpha_error, corrected_alpha, alpha_from_CI, alpha_max, alpha_min is returned. If alpha_output = "vector" unique alpha errors without a distinction of types is returned.
captions	Logical. If TRUE captions text will be scanned for statistical results.
text.mode	Numeric. Defines text parts to extract statistical results from (text.mode=1: abstract and full text, text.mode=2: method and result section, text.mode=3: result section only).
update.package.list	
	Logical. If TRUE updates available R packages with utils::available.packages() function.
add.software	additional software names to detect as vector.
quantileDF	quantile of (df1+1)+(df2+1) to extract for estimating sample size.
N.max.only	return only maximum of estimated sample sizes.
output	output selection of specific results c("doi", "title", "year", "Nstudies", "methods", "alpha_error", "power", "multi_comparison_correction", "assumptions", "OutlierRemovalInSD", "InteractionModeratorMediatorEffect", "test_direction", "sig_adjectives", "software", "Rpackage", "stats", "standardStats", "estimated_sample_size").
rm.na.col	Logical. If TRUE removes all columns with only NA in extracted standard statistics.

Source

An interactive web application for selecting and analyzing extracted article metadata and study characteristics for articles linked to PubMed Central is hosted at: https://www.scianalyzer.com/

The XML version of PubMed Central database articles can be downloaded in bulk from: https://ftp.ncbi.nlm.nih.gov/pub/pmc/oa_bulk/

References

Böschen (2021). "Evaluation of JATSdecoder as an automated text extraction tool for statistical results in scientific reports." *Scientific Reports.* doi: 10.1038/s41598-021-98782-3.

---

text2num	*text2num*

Description

Converts special annotated number and written numbers in a text string to a fully digit representation. Can handle numbers with exponent, fraction, percent, e+num, products and written representation (e.g. 'fourtys-one') of all absolut numbers up to 99,999 (Note: gives wrong output for higher spelled numbers). Process is performed in the same order as its arguments.

Usage

```
text2num(
  x,
  exponent = TRUE,
  percentage = TRUE,
  fraction = TRUE,
  e = TRUE,
  product = TRUE,
  words = TRUE
)
```

Arguments

x	text string to process.
exponent	Logical. If TRUE values with exponent are converted to a digit representation.
percentage	Logical. If TRUE percentages are converted to a digit representation.
fraction	Logical. If TRUE fractions are converted to a digit representation.
e	Logical. If TRUE values denoted with 'number e+number' (e.g. '2e+2') or number*10^number are converted to a digit representation.
product	Logical. If TRUE values products are converted to a digit representation.
words	Logical. If TRUE written numbers are converted to a digit representation.

Examples

```
x<-c("numbers with exponent: -2^3, .2^-2, -.3^.2, 49^-.5, 2^10.",
     "numbers with percentage: 2%, 15 %, 25 percent.",
     "numbers with fractions: 1/100, -2/5, -7/-.1",
     "numbers with e: 10e+2, -20e3, .2E-2, 2e4",
     "numbers as products: 100*2, -20*.1, 2*10^3",
     "written numbers: twenty-two, one hundred fourty five",
     "mix: one hundred ten is not 1/10 is not 10^2 nor 10%/5")
text2num(x)
```

---

text2sentences   *text2sentences*

Description

Converts floating text to a vector with sentences via fine-tuned regular expressions.

Usage

```
text2sentences(x)
```

Arguments

x	text string to process.

Examples

```
x<-"Some text with result (t(18)=1.2, p<.05). This shows how text2sentences works."
text2sentences(x)
```

---

vectorize.text   *vectorize.text*

Description

Converts vector of text to a list of vectors with words within each cell. Note: punctuation will be removed.

Usage

```
vectorize.text(x)
```

Arguments

x   text string to vectorize.

Examples

```
text<-"One hundred twenty-eight students participated in our Study,
that was administred in thirteen clinics."
vectorize.text(text)
```

---

which.term   *which.term*

Description

Returns search element/s from vector that is/are present in text or returns search term hit vector for all terms.

Usage

```
which.term(x, terms, tolower = TRUE, hits_only = FALSE)
```

Arguments

x	text string to process.
terms	search term vector.
tolower	Logical. If TRUE converts search terms and text to lower case.
hits_only	Logical. If TRUE returns search pattern/s, that were found in text and not a search term hit vector.

Examples

```
text<-c("This demo demonstrates how which.term works.",
        "The result is a simple 0, 1 coded vector for all search patterns or
         a vector including the identified patterns only.")
which.term(text,c("Demo","example","work"))
which.term(text,c("Demo","example","work"),tolower=TRUE,hits_only=TRUE)
```

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