

Package ‘Ecfun’

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Title Functions for Ecdat

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Description Functions to update datasets in Ecdat and to create,manipulate, plot and analyze those and similar datasets.

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R topics documented:

as.Date1970	2
asNumericDF	3
camelParse	5
financialCrisisFiles	6
grepNonStandardCharacters	7
match.data.frame	9
mergeUSHouse.senate	10
mergeVote	12
parseCommas	14
parseDollars	16
parseName	17
Ping	19
qqnorm2	21
qqnorm2s	24
read.testURLs	27
read.transpose	28
readCookPVI	30
readFinancialCrisisFiles	32
readNIPA	34
readUSHouse	35
readUSSenate	37
readUSStateAbbreviations	38
recode2	40
subNonStandardCharacters	41
subNonStandardNames	43
testURLs	44
USHouse.senate	47
USSenateClass	48
Index	51

as.Date1970	<i>Date from a number of days since the start of 1970.</i>
-------------	--

Description

as.Date.numeric requires origin to be specified. The present function assumes that this origin is January 1, 1970.

Usage

```
as.Date1970(x, ...)
```

Arguments

`x` a numeric vector of dates in days since the start of 1970.
`...` optional arguments to pass to `as.Date`.

Value

Returns a vector of Dates

Author(s)

Spencer Graves

See Also

[as.Date](#) [as.POSIXct1970](#)

Examples

```
days <- c(0, 1, 365)
Dates <- as.Date1970(days)
```

```
all.equal(c('1970-01-01', '1970-01-02', '1971-01-01'),
          as.character(Dates))
```

```
all.equal(days, as.numeric(Dates))
```

asNumericDF

Coerce to numeric dropping commas and info after a blank

Description

Delete commas (thousand separators) and drop information after a blank, then coerce to numeric and order the rows by the `orderBy`. Some Excel imports include commas as thousand separators; this replaces any commas with `char(0)`, ”. Also, some character data includes footnote references following the year. Table F-1 from the US Census Bureau needs all three of these features: It needs `orderBy`, because the most recent year appears first, just the opposite of most other data sets where the most recent year appears last. It has footnote references following a character string indicating the year. And it includes commas as thousand separators.

Usage

```
asNumericChar(x)
asNumericDF(x, keep=function(x)any(!is.na(x)), orderBy)
```

Arguments

x	For asNumericChar, this is a character vector to be converted to numeric after gsub(' ', '', x). For asNumericDF, this is a data.frame with all character columns to be converted to numerics.
keep	something to indicate which columns to keep
orderBy	Which columns to order the rows of x[, keep] by

Details

1. Replace commas by nothing
2. strsplit on ' ' and take only the first part, thereby eliminating the footnote references.
3. Replace any blanks with NAs
4. as.numeric
5. lapply(x, 1-4)
6. order the rows; by default, ascending on the first column

Value

all numeric data.frame

Author(s)

Spencer Graves

See Also

[scan gsub Quotes](#)

Examples

```
fakeF1 <- data.frame(yr=c('1948', '1947 (1)'),
                    q1=c('1,234', ''), duh=rep(NA, 2) )
nF1 <- asNumericDF(fakeF1)

nF1. <- data.frame(yr=asNumericChar(fakeF1$yr),
                  q1=asNumericChar(fakeF1$q1))[2:1,]

# correct answer
row.names(nF1.) <- 2:1

nF1c <- data.frame(yr=1947:1948, q1=c(NA, 1234))
row.names(nF1c) <- 2:1

all.equal(nF1, nF1.)
```

financialCrisisFiles *Files containing financial crisis data*

Description

FinancialCrisisFiles in Ecdat is an object of class financialCrisisFiles created by the financialCrisisFiles function to describe files containing data on financial crises downloadable from <http://www.reinhartandrogoff.com/data/browse-by-topic/topics/7/>.

Usage

```
financialCrisisFiles(files=c("22_data.xls", "23_data.xls",  
                          "Varieties_Part_III.xls", "25_data.xls"), ...)
```

Arguments

files	character vector of file names
...	arguments to pass with file and sheet name to read.xls when reading a sheet of an MS Excel file. This is assumed to be the same for all sheets of all files. If this is not the case, the resulting financialCrisisFiles object will have to be edited manually before using it to read the data.

Details

Reinhart and Rogoff (<http://www.reinhartandrogoff.com>) provide numerous data sets analyzed in their book, "This Time Is Different: Eight Centuries of Financial Folly". Of interest here are data on financial crises of various types for 70 countries spanning the years 1800 - 2010, downloadable from <http://www.reinhartandrogoff.com/data/browse-by-topic/topics/7/>.

The function financialCrisisFiles produces a list of class financialCrisisFiles describing four different Excel files in very similar formats with one sheet per Country and a few extra descriptor sheets. The data object FinancialCrisisFiles is the default output of that function.

It does this in several steps:

1. Read the first sheet of each file
2. Extract the names of the Countries from that first sheet.
3. Elimiate any blank spaces in the names to convert, e.g., "Costa Rica" to "CostaRica".
4. Find the sheets corresponding to each of the compressed names.
5. Construct the output list.

Value

The function financialCrisisFiles returns a list of class financialCrisisFiles. This is a list with components carrying the names of files to be read. Each component is a list of optional arguments to pass to `do.call(read.xls, ...)` to read the sheet with name = name of that component.

The default value returned by `financialCrisisFiles` is the data object `FinancialCrisisFiles`. This corresponds to the files downloaded from <http://www.reinhartandrogoff.com/data/browse-by-topic/topics/7/> in January 2013 (except for the fourth, which was not available there because of an error with the web site but instead was obtained directly from Prof. Reinhart).

Author(s)

Spencer Graves

Source

<http://www.reinhartandrogoff.com>

References

Carmen M. Reinhart and Kenneth S. Rogoff (2009) *This Time Is Different: Eight Centuries of Financial Folly*, Princeton U. Pr.

See Also

[read.xls](#)

Examples

```
Ecdat.demoFiles <- system.file('demoFiles', package='Ecdat')
Ecdat.xls <- dir(Ecdat.demoFiles, pattern='xls$',
               full.names=TRUE)
tst <- financialCrisisFiles(Ecdat.xls)

## Not run:
# check
\dontshow{stopifnot()}
all.equal(tst, data(FinancialCrisisFiles))
\dontrun{}}

## End(Not run)
```

grepNonStandardCharacters

grep for nonstandard characters

Description

Return the indices of elements of `x` containing characters that are not in `standardCharacters`.

Usage

```
grepNonStandardCharacters(x, value=FALSE,
  standardCharacters=c(letters, LETTERS, ' ', '.', ',', 0:9,
    '\"', "\'", '-', '_ ', '(', ')', '[', ']', '\n'),
  ... )
```

Arguments

x character vector in which it is desired to identify elements containing characters not in standardCharacters.

value logical: TRUE to return the values found in x, FALSE to return their indices.

standardCharacters Characters to overlook in x to identify anything not in standardCharacters.

... optional arguments for [regexpr](#)

Details

1. `x. <- strsplit(x, "")`: convert the input character vector to a list of vectors of character vectors with `nchar(x.[i]) == 1` for `i` in `1:length(x)`.
2. `sapply(x., ...)` to identify all elements for which any element of `x[[i]]` is not in `standardCharacters`.

Value

an integer vector identifying all elements of x containing a character not in standardCharacters.

Author(s)

Spencer Graves

See Also

[grep](#), [regexpr](#), [subNonStandardCharacters](#), [showNonASCII](#)

Examples

```
Names <- c('Raul', 'Ra`l', 'Torres,Raul', 'Torres, Raul')
# confusion in character sets can create
# names like Names[2]

chk <- grepNonStandardCharacters(Names)

all.equal(chk, 2)

chkv <- grepNonStandardCharacters(Names, TRUE)

all.equal(chkv, 'Ra`l')
```

match.data.frame	<i>Identify the row of y best matching each row of x</i>
------------------	--

Description

For each row of `x[, by.x]`, find the best matching row of `y[, by.y]`, with the best match defined by `grep.` and `split`.

`grep.` and `split` must either be `missing` or have the same length as `by.x` and `by.y`. If `grep.[i]` and `split[i]` are `NA`, do a complete match of `x[, by.x[i]]` and `y[, by.y[i]]`. Otherwise, for each row `j`, look for a match for `strsplit(x[j, by.x[i]], split[i])[[1]][1]` among `strsplit(y[, by.y[i]], split[i])`. See details.

Usage

```
match.data.frame(x, y, by, by.x=by, by.y=by, grep., split, sep=':')
```

Arguments

<code>x, y</code>	data.frames
<code>by, by.x, by.y</code>	names of columns of <code>x</code> and <code>y</code> to match.
<code>grep.</code>	a character vector of the type of match for each element of <code>by.x</code> and <code>by.y</code> . If <code>NA</code> , require a perfect match. Alternatives are <code>grep</code> and <code>agrep</code> to find a match for the first segment in <code>strsplit(x, split=split[i])</code> among any of the segments of <code>strsplit(y, split=split[i])</code> . Use <code>fixed=TRUE</code> with the calls to these functions. NOTE: These alternatives are not examined if a unique match is found between <code>x[, by.x[is.na(grep.) & is.na(split)]]</code> and the corresponding columns of <code>y</code> .
<code>split</code>	A character vector of split characters to pass to <code>strsplit</code> ; <code>strsplit</code> is not called if <code>is.na(split)</code> .
<code>sep</code>	a <code>sep</code> argument to use with <code>paste</code> to produce a matching key for the columns of <code>x</code> and <code>y</code> for which perfect matches are required. If <code>(missing(sep) && not(missing(grep.)))</code> <code>sep <- ' '</code> except where <code>grep. = NA</code> .

Details

1. Check `by.x`, `by.y`, `grep.` and `split`. If `((missing(by.x) | missing(by.y)) && missing(by))` `by <- names(x)`
2. `fullMatch <- (is.na(grep.) & is.na(split))`. Create `keyfx` and `keyfy` by pasting columns of `x[, by.x[fullMatch]]` and `y[, by.y[fullMatch]]`. Also create `x.` and `y.` = `strsplit` of `x[, by.x[!fullMatch]]`.
3. Iterate over rows of `x` looking for the best match. This includes an inner loop over columns of `x[, by.x[!fullMatch]]`, stopping on the first unique match. Return `(-1)` if no unique match is found.

Value

an integer vector of length nrow(x) containing the index of the best matching row of y or NA if no adequate match was found.

Author(s)

Spencer Graves

See Also

[strsplit](#), [is.na](#) [grep](#), [agrep](#) [match](#), [row.match](#), [join](#), [match_df](#) [classify](#)

Examples

```
newdata <- data.frame(state=c("AL", "MI", "NY"),
                     surname=c("Rogers", "Rogers", "Smith"),
                     givenName=c("Mike R.", "Mike K.", "Al"),
                     stringsAsFactors=FALSE)
reference <- data.frame(state=c("NY", "NY", "MI", "AL", "NY", "MI"),
                      surname=c("Smith", "Rogers", "Rogers (MI)",
                                "Rogers (AL)", "Smith", 'Jones'),
                      givenName=c("John", "Mike", "Mike", "Mike",
                                "T. Albert", 'Al Thomas'),
                      stringsAsFactors=FALSE)
newInRef <- match.data.frame(newdata, reference,
                             grep.=c(NA, 'agrep', 'agrep'))

all.equal(newInRef, c(4, 3, 5))
```

mergeUSHouse.senate *Expand a dataset on some members of the US Congress to the entire membership*

Description

Merge a [data.frame](#) regarding some members of the US Congress with a [data.frame](#) with general information on all members.

Usage

```
mergeUSHouse.senate(x, UScongress=USHouse.senate(),
                    newrows="amount0",
                    default=list(member=FALSE, amount=0, vote="notEligible",
                                incumbent=TRUE) )
```

Arguments

x	a data.frame to be merged with UScongress
UScongress	a data.frame to be merged with x.
newrows	name of a logical column to add that is TRUE for rows added to x and FALSE otherwise.
default	default values for columns of x identified by <code>regexr(names(default)[i], tolower(names(x)))</code> .

Details

1. `keyx <- with(x, paste(houseSenate, state, District, sep=":"))`
2. `keyy <- with(UScongress(houseSenate, state, District, sep=":"))`
3. `notx <- !is.element(keyy, keyx)`
4. `Y <- UScongress[notx,]`
5. add default columns to Y
6. if(!newrows is not in names(x))x <- cbind(x, newrows=FALSE)
7. `Y[, newrows] <- TRUE`
8. `xY <- rbind(x, Y[c(names(x))])`
9. replace 'Democrat' with 'Democratic' in `xY[['Party']]`
10. Look for NAs in "incumbent" who are nevertheless in UScongress; fix. Thus, if `x[['incumbent']]` is TRUE or FALSE, this value is not checked in UScongress; it's checked only if NA. The check consists of comparing names for a given Office:state:district between `strsplit(x[['surname']], ' ')[[1]][1]` and `strsplit(UScongress[['surname']], ' ')[[1]][1]` and similarly for givenName. This allows 'Rogers' in `x[['surname']]` to match 'Rogers (AL)' in `UScongress[['surname']]`, etc. The algorithm is not perfect, but errors should be rare – and could be fixed manually.

Value

a [data.frame](#) combining x and UScongress as desired

Author(s)

Spencer Graves

See Also

[merge USHouse.senate](#)

Examples

```
tst <- data.frame(Office=factor(rep(c('House', 'Senate'), c(4, 2))),
  State=factor(c('Missouri', 'Minnesota', 'Tennessee',
    'New York', rep('South Carolina', 2))),
  state=factor(c('MO', 'MN', 'TN', 'NY', 'SC', 'SC')),
  district=as.character(c(4, 1, 8, 18, 2, 3)),
  surname=c('Hartzler', 'Walz', 'Fincher', 'Maloney',
    'Graham', 'DeMint'),
```

```

givenName=c('Vicky', 'Timothy J.', 'Stephen Lee',
            'Sean Patrick', 'Lindsey', 'Jim'),
Party=c('Republican', 'Democrat', 'Republican', 'Democrat',
        'Republican', 'Democrat'),
CommitteeMember=rep(c(TRUE, FALSE), c(4, 2)),
amount=c(5000, 2000, 29500, 1000, 1000, 11500),
xvote=c('Y', 'N', 'Y', 'Y', 'notEligible', 'notEligible'),
incumbent=NA, stringsAsFactors=FALSE )
tst2 <- mergeUShouse.senate(tst)

# A couple of simple tests; don't test too much,
# because the results of UShouse.senate change,
# and we don't want this test to fail
# due to changes that don't affect Ecdat code

tst3 <- tst2[!tst2$amount0, c(1, 4:6, 8:10)]
row.names(tst) <- row.names(tst3)

## Not run:
all.equal(tst[c(1, 4:6, 8:10)], tst3)

## End(Not run)
# tst3[2] = state = factor with 56 levels,
# and tst[2] only has 5; compare without this

```

mergeVote

Merge Roll Call Vote

Description

Merge roll call vote record with a [data.frame](#) containing other information. The vote records are typically incomplete, so match first on houseSenate and surname. If this match is incomplete, try using givenName. If that fails, try state and district, which may not always be present in vote.

Usage

```
mergeVote(x, vote, Office="House", vote.x, check.x=TRUE)
```

Arguments

x	a data.frame whose columns include Office, surname, and givenName.
vote	a data.frame with column names which when forced tolower would match surname, givenname, and vote. However, the givenname may not be complete, so use it only if the surname is not sufficient.
Office	Either "House" or "Senate"; ignored if vote includes a column Office.

vote.x	name of a column of x containing a vote to be updated with the vote column of the vote <code>data.frame</code> . If <code>missing</code> and x has a column with a name matching "vote", then vote.x is that column. If <code>missing</code> but x has no such column, then append a column to x with the name of the vote column of the vote <code>data.frame</code> .
check.x	logical: If TRUE, check for rows of <code>x[, vote.x]</code> that are NOT in vote and throw an error if found.

Details

1. Parse `vote.x` to get the name of the column of x into which to write the vote column of the vote `data.frame`.
2. If the vote `data.frame` contains a column `Office`, ignore the `Office` argument. Otherwise, add the argument `houseSenate` as a column of vote.
3. Create `keyx <- with(x, paste(Office, surname, sep=":"))`, `keyx2 <- paste(keyx, givenName, sep=":"))`, `keyx. <- paste(houseSenate, state, district, sep=":"))`, and similarly `keyv`, `keyv2`, and `keyv.` from vote.
4. Look for `keyv` in `keyx`. When a unique match is found, transfer the vote the vote column of x. When no match is found, try for `keyv2` in `keyx2` or `keyv.` in `keyx`. If those fail, print an error message with the information from vote on all failures and ask the user to add state and district information.
5. `if(check.x)`, check for rows in `x[, vote.x]` that are NOT "notEligible" but are also not in vote: Throw an error if any are found.

Value

a `data.frame` with the same columns as x with its vote column modified per the vote argument.

Author(s)

Spencer Graves

See Also

[mergeUSHouse.senate](#)

Examples

```
##
## 1. Test good cases
##
votetst <- data.frame(
  surName=c('Smith', 'Jones', 'Graves', 'Jsn', 'Jsn', 'Gay'),
  givenName=c("Sam", "", "", "John", "John", ''),
  votex=factor(c('Y', 'N', 'abstain', 'Y', 'Y', 'Y')),
  State=factor(rep(c("CA", "", "SC", "NY"), c(1, 2, 1, 2))),
  district=rep(c("13", "1", "2", "1"), c(1, 2, 2, 1)),
  stringsAsFactors=FALSE )
```

```

x1 <- data.frame(
  Office=factor(rep(c("House", "Senate"), e=8)),
  state=rep(c("NY", "SC", "SD", "CA", "AK", "AR", "NY", "NJ"), 2),
  District=rep(c("2", "2", "At Large", "13", "1", "9", "1", "3"), 2),
  surname=rep(c('Jsn', 'Jsn', 'Smith', 'Smith', 'Jones',
    'Graves', 'Rx', 'Agnew'), 2),
  givenName=rep(c("John D.", "John J.",
    "Samual", "Samual", "Mary", "Mary", "Susan", 'Spiro'), 2),
  don=1:16, stringsAsFactors=FALSE)

x1. <- mergeVote(x1, votetst)

x2 <- cbind(x1, votex=factor( rep(
  c('Y', 'notEligible', 'Y', 'N', 'abstain', 'Y', 'notEligible'),
  c(2,1,1,1,1,1,9) ) ) )

all.equal(x1., x2)

##
## 2. Test a case with a vote error in x
##

x1a <- cbind(x1, voterr=rep(
  c('notEligible', 'Y', 'notEligible'), c(7, 1, 8)))

x1a. <- try(mergeVote(x1a, votetst))

class(x1a.)=='try-error'

```

parseCommas

Convert character string with Dollar signs and commas to numerics

Description

as.numeric of character strings after suppressing commas and dollar signs. This is a generalization of [parseDollars](#).

Usage

```

parseCommas(x, pattern='\\$', replacement='',
  acceptableErrorRate=0, ...)
## Default S3 method:
parseCommas(x, pattern='\\$', replacement='',
  acceptableErrorRate=0, ...)
## S3 method for class 'data.frame'

```

```
parseCommas(x, pattern='\\$|,', replacement='',
            acceptableErrorRate=0, ...)
```

Arguments

x vector of character strings to be converted to numerics

pattern regular expression to be replaced by replacement

replacement Character string to substitute for each occurrence of pattern

acceptableErrorRate number indicating the proportion of new NAs to that can be introduced and still assume it's numeric

... optional arguments to pass to [gsub](#)

Details

```
as.numeric(gsub(x, ...))
```

The `data.frame` method outputs another `data.frame` with character or factor columns converted to numerics using `parseDollars` whenever that can be done without creating NAs.

Value

Numeric vector converted from the character strings in `x` or a `data.frame` with columns that are obviously numbers in character format converted to numerics.

Author(s)

Spencer Graves

See Also

[gsub](#) [as.numeric](#) [parseDollars](#)

Examples

```
##
## 1. a character vector
##
X2 <- c('-2,500', '$5,000.50')
x2 <- parseDollars(X2)

all.equal(x2, c(-2500, 5000.5))

##
## A data.frame
##
chDF <- data.frame(let=letters[1:2], Dol=X2, dol=x2)
numDF <- parseCommas(chDF)
```

```
chkDF <- chDF
chkDF$Do1 <- x2

all.equal(numDF, chkDF)
```

parseDollars

Convert character string with Dollar signs and commas to numerics

Description

as.numeric of character strings after suppressing commas and dollar signs. This is a special case of [parseCommas](#).

Usage

```
parseDollars(x, pattern='\$|,', replacement='', ...)
```

Arguments

x	vector of character strings to be converted to numerics
pattern	regular expression to be replaced by replacement
replacement	Character string to substitute for each occurrence of pattern
...	optional arguments to pass to gsub

Details

as.numeric(gsub(x, ...)). See also [parseCommas](#).

Value

Numeric vector converted from x.

Author(s)

Spencer Graves

See Also

[gsub](#) [as.numeric](#) [parseCommas](#)

Examples

```
##
## 1. a character vector
##
X2 <- c('-$2,500', '$5,000.50')
x2 <- parseDollars(X2)

all.equal(x2, c(-2500, 5000.5))

##
## A data.frame
##
chDF <- data.frame(let=letters[1:2], Dol=X2, dol=x2)
numDF <- parseCommas(chDF)

chkDF <- chDF
chkDF$Dol <- x2

all.equal(numDF, chkDF)
```

parseName	<i>Parse surname and given name</i>
-----------	-------------------------------------

Description

Identify the presumed surname in a character string assumed to represent a name and return the result in a character matrix with "surname" followed by "givenName".

Usage

```
parseName(x, surnameFirst=(median(regexpr(',', x))>0),
          suffix=c('Jr.', 'I', 'II', 'III', 'IV', 'Sr.'),
          fixNonStandard=subNonStandardNames, ...)
```

Arguments

x	a character vector
surnameFirst	logical: If TRUE, the surname comes first followed by a comma (","), then the given name. If FALSE, parse the surname from a standard Western "John Smith, Jr." format. If missing(surnameFirst), use TRUE if half of the elements of x contain a comma.
suffix	character vector of strings that are NOT a surname but might appear at the end without a comma that would otherwise identify it as a suffix.

`fixNonStandard` function to look for and repair nonstandard names such as names containing characters with accent marks that are sometimes mangled by different software. Use [identity](#) if this is not desired.

... optional arguments passed to `fixNonStandard`

Details

If `surnameFirst` is FALSE:

1. If the last character is ")" and the matching "(" is 3 characters earlier, drop all that stuff. Thus, "John Smith (AL)" becomes "John Smith".
2. Look for commas to identify a suffix like Jr. or III; remove and call the rest x2.
3. `split <- strsplit(x2, " ")`
4. Take the last as the surname.
5. If the "surname" found per 3 is in suffix, save to append it to the `givenName` and recurse to get the actual surname.

NOTE: This gives the wrong answer with double surnames written without a hyphen in the Spanish tradition, in which, e.g., "Anastasio Somoza Debayle", "Somoza Debayle" give the (first) surnames of Anastasio's father and mother, respectively: The current algorithm would return "Debayle" as the surname, which is incorrect.

6. Recompose the rest with any suffix as the `givenName`.

Value

a character matrix with two columns: `surname` and `givenName`

Author(s)

Spencer Graves

See Also

[strsplit](#) [identity](#)

Examples

```
##
## 1. Parse standard first-last name format
##
tst <- c('Joe Smith (AL)', 'Teresa Angelica Sanchez de Gomez',
        'John Brown, Jr.', 'John Brown Jr.',
        'John W. Brown III', 'John Q. Brown,I',
        'Linda Rosa Smith-Johnson', 'Anastasio Somoza Debayle',
        'Ra_l Vel_zquez')
library(Ecdat)
parsed <- parseName(tst)

tst2 <- matrix(c('Smith', 'Joe', 'Gomez', 'Teresa Angelica Sanchez de',
                'Brown', 'John, Jr.', 'Brown', 'John, Jr.',
```

```

'Brown', 'John W., III', 'Brown', 'John Q., I',
'Smith-Johnson', 'Linda Rosa', 'Debayle', 'Anastasio Somoza',
'Velazquez', 'Raul'),
ncol=2, byrow=TRUE)
# NOTE: This second to last example is in the Spanish tradition
# and is handled incorrectly by the current algorithm.
# The correct answer should be "Somoza Debayle", "Anastasio".
# However, fixing that would complicate the algorithm excessively for now.
colnames(tst2) <- c("surname", 'givenName')

all.equal(parsed, tst2)

##
## 2. Parse "surname, given name" format
##
tst3 <- c('Smith (AL),Joe', 'Sanchez de Gomez, Teresa Angelica',
'Brown, John, Jr.', 'Brown, John W., III', 'Brown, John Q., I',
'Smith-Johnson, Linda Rosa', 'Somoza Debayle, Anastasio',
'Vel_zquez, Ra_1')
tst4 <- parseName(tst3)

tst5 <- matrix(c('Smith', 'Joe', 'Sanchez de Gomez', 'Teresa Angelica',
'Brown', 'John, Jr.', 'Brown', 'John W., III', 'Brown', 'John Q., I',
'Smith-Johnson', 'Linda Rosa', 'Somoza Debayle', 'Anastasio',
'Velazquez', 'Raul'),
ncol=2, byrow=TRUE)
colnames(tst5) <- c("surname", 'givenName')

all.equal(tst4, tst5)

```

Ping

ping a Uniform resource locator (URL)

Description

***NOTE: THIS IS A PRELIMINARY VERSION OF THIS FUNCTION; ***NOTE: IT MAY BE CHANGED OR REMOVED IN A FUTURE RELEASE.

ping a Uniform resource locator (URL) or Internet Protocol (IP) address.

NOTE: Some Internet Service Providers (ISPs) play games with "ping". That makes the results of Ping unreliable.

Usage

```

Ping(url, pingArgs='', warn=NA,
      show.output.on.console=FALSE)

```

Arguments

<code>url</code>	a character string of a URL or IP address to ping. If <code>url</code> is a vector of length greater than 1, only the first component is used.
<code>pingArgs</code>	arguments to pass to the ping command of typical operating systems via <code>pingResult <- system(paste('ping', pingArgs, url), intern=TRUE, ...)</code>
<code>warn</code>	value for <code>options('warn')</code> during the call to <code>system</code> . NA to not change <code>options('warn')</code> during this call.
<code>show.output.on.console</code>	argument for <code>system</code> .

Details

1. `urlSplit0 <- strsplit(url, '://')[[1]]`
2. `urlS0 <- urlSplit0[min(2, length(urlSplit0))]`
3. `host <- strsplit(urlS0, '/')[[1]][1]`
4. `pingCmd <- paste('ping', pingArgs, host)`
5. `system(pingCmd, intern=TRUE, ...)`

Value

list with the following components:

<code>rawResults</code>	character vector of the raw results from the ping command
<code>rawNumbers</code>	numeric vector of the times measured
<code>counts</code>	numeric vector of numbers of packets sent, received, and lost
<code>p.lost</code>	proportion lost = lost / sent
<code>stats</code>	numeric vector of min, avg (mean), max, and mdev (standard deviation) of the measured round trip times

Author(s)

Spencer Graves

See Also

[system](#), [options](#)

Examples

```
##
## Some ISPs play games with ping.
## Therefore, the results are not reliable.
##
## Not run:
##
## good
```

```
##
(google <- Ping('http://google.com/ping works on host not pages'))

\dontshow{stopifnot()}
with(google, (counts[1]>0) && (counts[3]<1))
\dontshow{}}

##
## ping oops <<-- at one time, this failed.
##     However, with some ISPs, it works, so don't test it.
##

##
(couldnotfindhost <- Ping('oops'))

\dontshow{stopifnot()}
with(couldnotfindhost,
      length(grep('could not find host', rawResults))>0)
\dontshow{}}

##
## impossible, but not so obvious
##
(requesttimedout <- Ping('requesttimedout.com'))

\dontshow{stopifnot()}
with(requesttimedout, (counts[1]>0) && (counts[2]<1) &&
                      (counts[3]>0))
\dontshow{}}

## End(Not run)
```

qqnorm2

Normal Probability Plot with Multiple Symbols

Description

Create a normal probability plot with different symbols for the values of another variable. qqnorm2 produces an object of class qqnorm2, whose plot method produces the plot.

Usage

```
qqnorm2(y, z, plot.it=TRUE, datax=TRUE, pch=NULL, ...)
## S3 method for class 'qqnorm2'
plot(x, y, ...)
## S3 method for class 'qqnorm2'
lines(x, ...)
## S3 method for class 'qqnorm2'
points(x, ...)
```

Arguments

<code>y</code>	For <code>qqnorm2</code> , <code>y</code> is a numeric vector for which a normal probability plot is desired. For <code>plot.qqnorm2</code> , <code>y</code> is ignored; it is included, because the generic <code>plot</code> function requires it.
<code>z</code>	A variable to indicate different plotting symbols.
<code>plot.it</code>	logical: Should the result be plotted?
<code>datax</code>	The <code>datax</code> argument of <code>qqnorm</code> : If TRUE, the data are displayed on the horizontal rather than the vertical axis. (The default value for <code>datax</code> is the opposite of that for <code>qqnorm</code> .)
<code>x</code>	an object of class <code>qqnorm2</code> .
<code>pch</code>	a named vector of the plotting symbols to be used with names corresponding to the levels of <code>z</code> . By default, if <code>z</code> takes levels FALSE and TRUE (or 0 and 1), <code>pch=c(4, 1)</code> to plot a "x" for FALSE and "o" for TRUE. If <code>z</code> assumes integer values between 0 and 255, by default, the symbols are chosen as described with <code>points</code> . Otherwise, by default, <code>z</code> is coerced to <code>character</code> , and the result is plotted. If <code>pch</code> is provided, it must either have names corresponding to levels of <code>z</code> , or <code>z</code> must be integers between 1 and <code>length(pch)</code> .
<code>...</code>	Optional arguments. For <code>plot.qqnorm2</code> , they are passed to <code>plot</code> . For <code>qqnorm2</code> , they are passed to <code>qqnorm</code> and to <code>plot.qqnorm2</code> .

Details

For `qqnorm2`:

1. `q2 <- qqnorm(y, datax=datax, ...)`
2. `q2[["z"]] <- z`
3. `q2[["pch"]]` gets whatever `pch` decodes to.
4. Silently return(`list(x, y, z, pch)`), where "x" and "y" are as returned by `qqnorm` in step 1 above.

For `plot.qqnorm2`, `plot(x, y, pch=pch[z], ...)`. For `lines.qqnorm2`, `lines(x, y, pch=pch[z], ...)`. For `points.qqnorm2`, `points(x, y, pch=pch[z], ...)`.

Value

`qqnorm2` returns a list with components, `x`, `y`, `z`, and `pch`.

Author(s)

Spencer Graves

See Also

[qqnorm](#), [qqnorm2s](#), [plot](#) [points](#) [lines](#)

Examples

```

##
## a simple test data.frame to illustrate the plot
## but too small to illustrate qqnorm concepts
##
tstDF <- data.frame(y=1:3, z1=1:3, z2=c(TRUE, TRUE, FALSE),
                   z3=c('tell', 'me', 'why'), z4=c(1, 2.4, 3.69) )
# plotting symbols circle, triangle, and "+"
qn1 <- with(tstDF, qqnorm2(y, z1))

# plotting symbols "x" and "o"
qn2 <- with(tstDF, qqnorm2(y, z2))

# plotting with "-" and "+"
qn. <- with(tstDF, qqnorm2(y, z2, pch=c('FALSE'='-', 'TRUE'='+')))

# plotting with "tell", "me", "why"
qn3 <- with(tstDF, qqnorm2(y, z3))

# plotting with the numeric values
qn4 <- with(tstDF, qqnorm2(y, z4))

##
## test plot, lines, points
##
plot(qn4, type='n') # establish the scales
lines(qn4)         # add a line
points(qn4)        # add points

##
## Check the objects created above
##
# check qn1
qn1. <- qqnorm(1:3, datax=TRUE, plot.it=FALSE)
qn1.$xlab <- 'y'
qn1.$ylab <- 'Normal scores'
qn1.$z <- tstDF$z1
qn1.$pch <- 1:3
names(qn1.$pch) <- 1:3
qn11 <- qn1.[c(3:4, 1:2, 5:6)]
class(qn11) <- 'qqnorm2'

all.equal(qn1, qn11)

# check qn2
qn2. <- qqnorm(1:3, datax=TRUE, plot.it=FALSE)
qn2.$xlab <- 'y'
qn2.$ylab <- 'Normal scores'
qn2.$z <- tstDF$z2
qn2.$pch <- c('FALSE'=4, 'TRUE'=1)
qn22 <- qn2.[c(3:4, 1:2, 5:6)]

```

```

class(qn22) <- 'qqnorm2'

all.equal(qn2, qn22)

# check qn.
qn.. <- qqnorm(1:3, datax=TRUE, plot.it=FALSE)
qn..$xlab <- 'y'
qn..$ylab <- 'Normal scores'
qn..$z <- tstDF$z2
qn..$pch <- c('FALSE'='-', 'TRUE'='+')
qn.2 <- qn..[c(3:4, 1:2, 5:6)]
class(qn.2) <- 'qqnorm2'

all.equal(qn., qn.2)

# check qn3
qn3. <- qqnorm(1:3, datax=TRUE, plot.it=FALSE)
qn3.$xlab <- 'y'
qn3.$ylab <- 'Normal scores'
qn3.$z <- as.character(tstDF$z3)
qn3.$pch <- as.character(tstDF$z3)
names(qn3.$pch) <- qn3.$pch
qn33 <- qn3.[c(3:4, 1:2, 5:6)]
class(qn33) <- 'qqnorm2'

all.equal(qn3, qn33)

# check qn4
qn4. <- qqnorm(1:3, datax=TRUE, plot.it=FALSE)
qn4.$xlab <- 'y'
qn4.$ylab <- 'Normal scores'
qn4.$z <- tstDF$z4
qn44 <- qn4.[c(3:4, 1:2, 5)]
qn44$pch <- NULL
class(qn44) <- 'qqnorm2'

all.equal(qn4, qn44)

```

qqnorm2s

Normal Probability Plot with Multiple Lines and Multiple Symbols

Description

Create a normal probability plot with multiple lines for different variables and different symbols for the values of another variable. `qqnorm2s` produces an object of class `qqnorm2s`, whose `plot` method produces the plot.

Usage

```
qqnorm2s(y, z, data., plot.it=TRUE, datax=TRUE, outnames=y,
         pch=NULL, col=c(1:4, 6), legend.=NULL, ...)
## S3 method for class 'qqnorm2s'
plot(x, y, ...)
```

Arguments

y	For qqnorm2s, y is a character vector of names of columns of data. for which normal probability plots are desired. data. is either a data.frame or a list of data.frames of the same length as y, with y[i] being the name of a column of the data.frame data.[[i]]. z is a similar character vector of names of columns of data., which identify symbols for plotting different points in a normal probability plot. The lengths of y, and z must match the number of data.frames in data.; if not, the lengths of the shorter are replicated to the length of the longest before computations begin. For plot.qqnorm2, y is ignored; it is included, because the generic plot function requires it.
z	A character vector giving the names of columns of data. to indicate different plotting symbols. z should be the same length as y and must equal the number of data.frames in the list data. of data.frames. If not, the shorter are replicated to the length of the longer.
data.	a data.frame or a list of data.frames with columns named in y and z.
plot.it	logical: Should the result be plotted?
datax	The datax argument of qqnorm : If TRUE, the data are displayed on the horizontal rather than the vertical axis. (The default value for datax is the opposite of that for qqnorm .)
outnames	Names for the components of the qqnorm2s object returned by the qqnorm2s function.
pch	a named vector of the plotting symbols to be used with names corresponding to the levels of z. By default, if z takes levels FALSE and TRUE (or 0 and 1), pch=c(4, 1) to plot a "x" for FALSE and "o" for TRUE. If z assumes integer values between 0 and 255, by default, the symbols are chosen as described with points . Otherwise, by default, z is coerced to character , and the result is plotted. If pch is provided, it must either have names corresponding to levels of z, or z must be integers between 1 and length(pch).
col	A vector indicating the colors corresponding to each element of y. Defaults to rep(c(1:4, 6), length=length(y)), with 1:4 and 6 being black, red, green, blue, and pink.
x	an object of class qqnorm2.
legend.	A list with components pch and col providing information for legend to identify the plotting symbols (pch) and colors (col).

By default, `pch = list(x='right', legend=names(qq2s[[1]][['pch']], pch=qq2s[[1]][['pch']]))`, where `qq2s` is described below in details.

Similarly, by default, `lines = list(x='bottomright', legend=y, lty=1, pch=NA, col=qq2s[[1]][['col']])`.

...

Optional arguments.

For `plot.qqnorm2s`, they are passed to `plot`.

For `qqnorm2s`, they are passed to `qqnorm2` and to `plot.qqnorm2s`.

Details

For `qqnorm2s`:

1. Create `qq2s = a list of objects of class qqnorm2`
2. Add `legend.` to `qq2s`.
3. `class(qq2s) <- 'qqnorm2s'`
4. `if(plot.it)plot(qq2s, ...)`
5. Silently return(`qq2s`).

For `plot.qqnorm2s`, create a plot with one line for each variable named in `y`.

Value

`qqnorm2s` returns a named list with components of class `qqnorm2` with `names = y` with each component having an additional component `col` plus one called "legend".

Author(s)

Spencer Graves

See Also

[qqnorm2 plot](#)

Examples

```
##
## One data.frame
##
tstDF2 <- data.frame(y=1:3, y2=3:5, z2=c(TRUE, TRUE, FALSE),
                    z3=c('tell', 'me', 'why'), z4=c(1, 2.4, 3.69) )
# produce the object and plot it
Qn2 <- qqnorm2s(c('y', 'y2'), 'z2', tstDF2)

# plot the object previously created
plot(Qn2)

# Check the object
qy <- with(tstDF2, qqnorm2(y, z2, type='b'))
qy$col <- 1
qy2 <- with(tstDF2, qqnorm2(y2, z2, type='b'))
```

```

qy2$col <- 2
legend. <- list(pch=list(x='right', legend=c('FALSE', 'TRUE'),
                pch=c('FALSE'=4, 'TRUE'= 1)),
               col=list(x='bottomright', legend=c('y', 'y2'),
                lty=1, col=1:2))
Qn2. <- list(y=qy, y2=qy2, legend.=legend.)
class(Qn2.) <- 'qqnorm2s'

all.equal(Qn2, Qn2.)

##
## Two data.frames
##
tstDF2b <- tstDF2
tstDF2b$y <- c(0.1, 0.1, 9)
Qn2b <- qqnorm2s('y', 'z2', list(tstDF2, tstDF2b),
                outnames=c('ok', 'oops'), log='x' )

```

read.testURLs	<i>Read a file produced by testURLs</i>
---------------	---

Description

***NOTE: THIS IS A PRELIMINARY VERSION OF THIS FUNCTION; ***NOTE: IT MAY BE CHANGED OR REMOVED IN A FUTURE RELEASE.

read.table(file.) and return the result as an object of class c('testURLs', 'data.frame').

Usage

```
read.testURLs(file.='testURLresults.csv', ...)
```

Arguments

file.	Name of a CSV file to read
...	optional arguments for read.csv .

Details

```

dat <- read.csv(file., ...)
class(dat) <- c('testURLsFile', 'data.frame')

```

Value

a [data.frame](#) from the file written by [testURLs](#), of the same format as the testResults attribute of the testURLs object returned by [testURLs](#).

Author(s)

Spencer Graves

See Also[read.csv](#)**Examples**

```
# Test only 2 web sites, not the default 4,
# and test only twice, not the default 10 times:
tst <- testURLs(c(
  PVI="http://en.wikipedia.org/wiki/Cook_Partisan_Voting_Index",
  house="http://house.gov/representatives"),
  n=2, maxFail=2)

# The above should have created a file 'testURLresults.csv'
# in the working directory. Read it.

dat <- read.testURLs()
```

read.transpose

*Read a data table in transpose form***Description**

Read a text (e.g., csv) file, find rows with more than 3 sep characters. Parse the initial contiguous block of those into a matrix. Add attributes headers, footers, and a summary.

The initial application for this function is to read Table 6.16. Income and employment by industry in the National Income and Product Account tables published by the Bureau of Economic Analysis of the United States Department of Commerce.

Usage

```
read.transpose(file, header=TRUE, sep=',',
  na.strings='---', ...)
```

Arguments

file	the name of a file from which the data are to be read.
header	Logical: Is the second column of the identified data matrix to be interpreted as variable names?
sep	The field space separator character.
na.strings	character string(s) that translate into NA
...	optional arguments for strsplit

Details

1. `txt <- readLines(file)`
2. Split into fields.
3. Identify headers, Data, footers.
4. Recombine the second component of each Data row if necessary so all have the same number of fields.
5. Extract variable names
6. Numbers?
7. return the transpose

Value

A matrix of the transpose of the rows with the max number of fields with attributes 'headers', 'footers', 'other', and 'summary'. If this matrix can be coerced to numeric with no NAs, it will be. Otherwise, it will be left as character.

Author(s)

Spencer Graves

References

Table 6.16. Income and employment by industry in the National Income and Product Account tables published by the Bureau of Economic Analysis of the United States Department of Commerce. To get this table from www.bea.gov, under "U.S. Economic Accounts", first select "Corporate Profits" under "National". Then next to "Interactive Tables", select, "National Income and Product Accounts Tables". From there, select "Begin using the data...". Under "Section 6 - income and employment by industry", select each of the tables starting "Table 6.16". As of February 2013, there were 4 such tables available: Table 6.16A, 6.16B, 6.16C and 6.16D. Each of the last three are available in annual and quarterly summaries. The `USFinanceIndustry` data combined the first 4 rows of the 4 annual summary tables.

See Also

[read.table](#) [readLines](#) [strsplit](#)

Examples

```
# Find demoFiles/*.csv
demoDir <- system.file('demoFiles', package='Ecdat')
(demoCsv <- dir(demoDir, pattern='csv$', full.names=TRUE))

# Use the fourth example
# to ensure the code will handle commas in a name
# and NAs
nipa6.16D <- read.transpose(demoCsv[4])
str(nipa6.16D)
```

readCookPVI *Read Cook Partisan Voting Index*

Description

Read tables of the Cook Partisan Voting Index and returns a list with components 'House' and 'Senate'. readCookPVI returns the tables with the names of the current incumbents per readUShouse and readUSSenate; readCookPVI tables do not include the names of the incumbents.

Usage

```
readCookPVI(url.=
"http://en.wikipedia.org/wiki/Cook_Partisan_Voting_Index")
readCookPVI(url.=
"http://en.wikipedia.org/wiki/Cook_Partisan_Voting_Index",
            UShouse=readUShouse(), USSenate=readUSSenate(), ...)
```

Arguments

url. Universal resource locator to be read and processed to obtain the desired lists.
UShouse, USSenate [data.frame](#)s as returned by readUShouse and readUSSenate, respectively.
... optional arguments passed to readUShouse and readUSSenate.

Details

The primary source for these data is the Cook Political Report web site. However, the current URL we have for these data on that web site includes "2012" in the title. If and when the numbers are updated, we would expect that file name to change.

To avoid that problem the code is currently set to read from the Wikipedia article on "Cook Partisan Voting Index".

The algorithm reads the web site into a list, finds the desired tables on the list, then parses and formats them as desired. Then it merges the results with UShouse and USSenate.

Value

A list with components "House" and "Senate". Each contains a [data.frame](#). The "House" data.frame returned by readCookPVI includes the following columns:

State	name of the state
District	District, e.g, 1st, 2nd, At-Large
PVInum	PVI as a number ranging from roughly 50 to 150. 100 means that the vote split in that district was within 0.5 percent of the national average. 101 means that it tilts 1 percent (after rounding) to Republican. 98 means that it tilts 1 percent to Democratic; 99 is not used.

PVIchar PVI rating in character format. For example, 'D+1' means that the vote tilted 1 percent toward Democratic more than the national average. 'R+1' means that it tilted 1 percent toward Republican.

PartyOfRepresentative Party of the incumbent, either 'Republican' or 'Democratic'

The 'Senate' data.frame includes the following columns:

State name of the state

PVInum PVI numeric, as for 'House'

PVIchar PVI rating in character format, as for 'House'

PartyOfGovernor Party of the Governor of the state

PartyInSenate party of the incumbent senators, either 'Republican', 'Democratic', or 'Both'.

houseBalanceNum House balance as a number with 0 = 100 percent Democratic, 99.9 = 100 percent Republican, and 500 for the same number of Republicans as Democrats.

houseBalanceChar Count by party in the house delegation for that state, e.g., '6R, 1D' for 6 Republicans and 1 Democrat.

readCookPVI. adds to the above the information returned by [readUShouse](#) and [readUSsenate](#).

Author(s)

Spencer Graves

Source

[Wikipedia, "Cook Partisan Voting Index" The Cook Political Report](#)

See Also

[readUShouse](#), [readUSsenate](#)

Examples

```
## Not run:
CookPVI <- readCookPVI()

## End(Not run)

CookPVI. <- readCookPVI.()
```

readFinancialCrisisFiles

banking crisis data and function to read financial crisis files

Description

Read financial crisis data in files described by an object of class `financialCrisisFiles`. This is designed to read Excel files describing financial crises since 1800 downloaded from <http://www.reinhartandrogoff.com/data/browse-by-topic/topics/7/>.

`bankingCrises` is a `data.frame` created by `readFinancialCrisisFiles()` using 3 files downloaded from <http://www.reinhartandrogoff.com> and 1 obtained from Prof. Reinhart in January 2013.

Usage

```
readFinancialCrisisFiles(files, crisisType=7, ...)
```

Arguments

<code>files</code>	an object of class <code>financialCrisisFiles</code> .
<code>crisisType</code>	an integer (vector) between 1 and 8 indicating the type of data to be retrieved: 1=independence year (not a crisis but an indicator), 2=currency, 3=inflation, 4=stock market, 5=domestic sovereign debt crisis, 6=external sovereign debt crisis, 7=banking, 8=tally. ("Type" 1 = year.) These are all 0 or 1 indicating the presence of the event in the given year. Type 8 = sum of types 2 through 7.
<code>...</code>	arguments to pass with file and sheet name to <code>read.xls</code> when reading a sheet of an MS Excel file. This is assumed to be the same for all sheets of all files. If this is not the case, the resulting <code>financialCrisisFiles</code> object will have to be edited manually before using it to read the data.

Details

Reinhart and Rogoff (<http://www.reinhartandrogoff.com>) provide numerous data sets analyzed in their book, "This Time Is Different: Eight Centuries of Financial Folly". Of interest here are data on financial crises of various types for 70 countries spanning the years 1800 - 2010, downloadable from <http://www.reinhartandrogoff.com/data/browse-by-topic/topics/7/>.

The function `financialCrisisFiles` produces a list of class `financialCrisisFiles` describing different Excel files in very similar formats with one sheet per Country and a few extra descriptor sheets. The data object `FinancialCrisisFiles` is the default output of that function.

`readFinancialCrisisFiles` reads the sheets for the individual countries.

Value

If `length(crisisType) == 1`, a `data.frame` is returned with the first column being year, and with one other column containing the data for that `crisisType` for each country.

If `length(crisisType) > 1`, a list is returned containing a `data.frame` for each country.

Author(s)

Spencer Graves

Source

<http://www.reinhartandrogoff.com>

References

Carmen M. Reinhart and Kenneth S. Rogoff (2009) This Time Is Different: Eight Centuries of Financial Folly, Princeton U. Pr.

See Also

[read.xls financialCrisisFiles](#)

Examples

```
##
## Recreate / update the data object BankingCrises
##
library(Ecdat)

## Not run:
bankingCrises <- readFinancialCrisisFiles(FinancialCrisisFiles)

## End(Not run)

##
## Toy example using local data to check the code
## and illustrate returning all the data not just one crisisTpe
##
Ecdat.demoFiles <- system.file('demoFiles', package='Ecdat')
Ecdat.xls <- dir(Ecdat.demoFiles, pattern='xls$',
               full.names=TRUE)
tst <- financialCrisisFiles(Ecdat.xls)

bankingCrises.tst <- readFinancialCrisisFiles(tst)
allCrises.tst <- readFinancialCrisisFiles(tst, 1:8)

# Manually construct tst from allCrises.tst
tst2 <- data.frame(year=1800:1999)
tst2$Algeria <- as.numeric(allCrises.tst$Algeria[-(1:12), 8])
tst2$CentralAfricanRep <- as.numeric(
  allCrises.tst$CentralAfricanRep[-(1:12), 8])
tst2$Taiwan <- as.numeric(allCrises.tst$Taiwan[-(1:11), 8])
tst2$UK <- as.numeric(allCrises.tst$UK[-(1:11), 8])

all.equal(bankingCrises.tst, tst2)
```

```
# check
data(bankingCrises)

all.equal(bankingCrises.tst,
  bankingCrises[1:200, c('year', 'Algeria', 'CentralAfricanRep',
    'Taiwan', 'UK')])
```

readNIPA

Read a National Income and Product Accounts data table

Description

Read multiple files with data in rows using [read.transpose](#) and combine the initial columns.

Usage

```
readNIPA(files, sep.footnote='/', ...)
```

Arguments

files	A character vector of names of files from which the data are to be read using read.transpose .
sep.footnote	a single character to identify footnote references in the variable names in some but not all of files.
...	optional arguments for read.transpose

Details

This is written first and foremost to facilitate updating [USFinanceIndustry](#) from Table 6.16: Income and employment by industry in the National Income and Product Account tables published by the Bureau of Economic Analysis of the United States Department of Commerce. As of February 2013, this table can be obtained from <http://www.bea.gov>: Under "U.S. Economic Accounts", first select "Corporate Profits" under "National". Then next to "Interactive Tables", select, "National Income and Product Accounts Tables". From there, select "Begin using the data...". Under "Section 6 - income and employment by industry", select each of the tables starting "Table 6.16". As of February 2013, there were 4 such tables available: Table 6.16A, 6.16B, 6.16C and 6.16D. Each of the last three are available in annual and quarterly summaries. The [USFinanceIndustry](#) data combined the first 4 rows of the 4 annual summary tables.

This is available in 4 separate files, which must be downloaded and combined using `readNIPA`. The first three of these are historical data and are rarely revised. For convenience and for testing, they are provided in the `demoFiles` subdirectory of this `Ecdat` package.

It has not been tested on other data but should work for annual data with a sufficiently similar structure.

The algorithm proceeds as follows:

1. `Data <- lapply(files, read.transpose)`

2. Is Data a list of numeric matrices? If no, print an error.
3. cbind common initial variables, averaging overlapping years, reporting percent difference
4. attributes: stats from files and overlap. Stats include the first and last year and the last revision date for each file, plus the number of years overlap with the previous file and the relative change in the common files kept between those two files.

Value

a `matrix` of the common variables

Author(s)

Spencer Graves

References

[United States Department of Commerce Bureau of Economic Analysis National Income and Product Account tables](#)

See Also

`read.table` `readLines` `strsplit`

Examples

```
# Find demoFiles/*.csv
demoDir <- system.file('demoFiles', package='Ecdat')
(demoCsv <- dir(demoDir, pattern='csv$', full.names=TRUE))

nipa6.16 <- readNIPA(demoCsv)
str(nipa6.16)
```

readUShouse	<i>Read the list of representatives in the United States House of Representatives</i>
-------------	---

Description

Read the list of representatives in the United States House of Representatives.

Usage

```
readUShouse(url.="http://house.gov/representatives/",
  nonvoting=c('American Samoa', 'District of Columbia',
    'Guam', 'Northern Mariana Islands', 'Puerto Rico',
    'Virgin Islands'),
  fixNonStandard=subNonStandardNames, ...)
```

Arguments

<code>url</code>	Universal resource locator to be read and processed to obtain the desired list
<code>nonvoting</code>	Character vector of the names of US territories that send a nonvoting delegate to the US House.
<code>fixNonStandard</code>	function to look for and repair nonstandard names such as names containing characters with accent marks that are sometimes mangled by different software. Use <code>identity</code> if this is not desired.
<code>...</code>	optional arguments passed to <code>fixNonStandard</code>

Details

1. `House.gov <- readHTMLTable(url)`. As of April 2013, this is a list of 80 tables. The first 56 are for the 50 states and 6 territories. The remaining 24 are for the first letter of the last name of the representatives.
2. Use `rbind` to collapse these into 2 tables. The first has the district as a number without identifying the state (because that was with the names of the first 56 tables in `House.gov`). The second has the state names but with the district numbers in a form not easily parsed.
3. Obtain the state names from the second table to match the names of the representatives in the first.
4. Add a `nonvoting` column for those "States" in `nonvoting`.
5. Look for and fix `surname` and `givenName` with nonstandard characters using `fixNonStandard`.

Value

`readUShouse` returns a `data.frame` with the following columns:

<code>State</code>	A factor identifying the state or territory the person represents
<code>state</code>	2-letter US Postal Service abbreviation for the state or territory
<code>district</code>	the character vector identifying the district each person represents. This is either an integer in character format or 0 for "At Large".
<code>Name</code>	A character vector giving the name of each representative (in surname, given name format)
<code>party</code>	a factor identifying the party affiliation of each representative ("D" or "R").
<code>Room</code>	character vector identifying the room number of the office
<code>Phone</code>	character vector giving the phone number
<code>Committees</code>	a character vector giving the committee assignments of each representative
<code>surname</code>	character vector giving the surname of each representative
<code>givenName</code>	given name of each representative (possibly with middle name or initial, a nickname, and a suffix like "Jr.")

Author(s)

Spencer Graves

See Also

[getURL](#) [readHTMLTable](#) [readUSsenate](#) [UShouse.senate.parseName](#) [readUSstateAbbreviations](#)
[subNonStandardNames](#) [readCookPVI](#)

Examples

```
UShouse <- readUShouse()
```

readUSsenate	<i>Read the list of elected officials in the United States Senate</i>
--------------	---

Description

Read the list of elected officials in the United States Senate.

Usage

```
readUSsenate(url.=
  "http://en.wikipedia.org/wiki/List_of_current_United_States_Senators",
  stateAbbreviations=Ecdat::USstateAbbreviations,
  fixNonStandard=subNonStandardNames, ...)
```

Arguments

url.	Universal resource locator to be read and processed to obtain the desired list NOTE: On April 26, 2013 the obvious naive use of readHTMLTable worked with the Wikipedia article on the US Senate but did not work with "senate.gov".
stateAbbreviations	a data.frame giving names and alternative codes for US states and territories. This must have a column named "Name" giving the names of all 50 states as they appear on the url and another whose name includes "USPS" giving the corresponding 2-letter codes.
fixNonStandard	function to look for and repair nonstandard names such as names containing characters with accent marks that are sometimes mangled by different software.
...	optional arguments passed to fixNonStandard

Details

1. Senate <- readHTMLTable(url)
2. Use [camelParse](#) to remove duplication in Name.
3. Look for and fix surname and givenName with nonstandard characters using fixNonStandard.

Value

readUSsenate returns a data.frame with the following columns:

State	A factor identifying the state the person represents
state	A factor giving the 2-letter USPS code for the state represented
Class	"1", "2", or "3" for election in the 6-year cycle including 2008, 2010, or 2012, respectively.
Name	A character vector giving the name of each representative (in surname, given name format)
Party	a factor identifying the party affiliation of each representative ("Democrat", "Republican", or "Independent").
Experience	character vector highlighting prior experience.
assumedOffice	character vector giving the date assumed office
Born	a character vector giving the year of birth
endOffice	a character vector giving the last day in the present term.
surname	character vector giving the surname of each representative
givenName	given name of each representative (possibly with middle name or initial, a nickname, and a suffix like "Jr.")

Author(s)

Spencer Graves

See Also

[getUrl readHTMLTable camelParse](#) to remove duplication in Name [readUShouse UShouse.senate parseName subNonStandardNames](#)

Examples

```
USsenate <- readUSsenate()
```

```
readUSstateAbbreviations
```

Read a list of abbreviations of states and territories of the United States

Description

Read the list of abbreviations of states and territories of the United states from the relevant Wikipedia article

Usage

```
readUSstateAbbreviations(url.=  
"http://en.wikipedia.org/wiki/List_of_U.S._state_abbreviations",  
  clean=TRUE, Names=c('Name', 'Status', 'ISO', 'ANSI.letters',  
    'ANSI.digits', 'USPS', 'USCG', 'Old.GPO', 'AP', 'Other') )
```

Arguments

url.	Universal resource locator to be read and processed to obtain the desired list
clean	logical: If TRUE, clean the data using subNonStandardCharacters and <code>strsplit(x, "\\[")</code>
Names	names for the columns of the data matrix read; ignored with a warning if the lengths do not match

Details

Wrapper for [readHTMLTable](#).

NOTE: `readHTMLTable(url)` returns a list of length 7, only one of which is the table we want. Moreover, that table contains some duplicates, which are removed by `readUSstateAbbreviations`. For example, 'NB' is an "Obsolete postal code" for Nebraska. If you need this, please consult the Wikipedia article.

Value

`readUSstateAbbreviations` returns a `data.frame` from the table in [the Wikipedia article on "List of U.S. state abbreviations"](#).

Author(s)

Spencer Graves

See Also

[getURL](#) [readHTMLTable](#) [make.names](#) [USstateAbbreviations](#)

Examples

```
abbreviations <- readUSstateAbbreviations()
```

`recode2`*bivariate recode*

Description

Recode `x1` and `x2` per the lexical codes table.

Usage

```
recode2(x1, x2, codes)
```

Arguments

<code>x1, x2</code>	vectors of the same length assuming a discrete number of levels
<code>codes</code>	a 2-dimensional matrix indexed by the levels of <code>x1</code> and <code>x2</code> . If <code>dimnames(codes)</code> are not provided, they are assumed to <code>unique(x1)</code> (or <code>unique(x2)</code>).

Details

1. If `length(x1) != length(x2)`, complain.
2. `if(is.logical(x1)) l1 <- c(FALSE, TRUE) else l1 <- unique(x1)`; ditto for `x2`.
3. `if(missing(codes)) codes <- outer(unique(x1), unique(x2))`
4. `if(is.null(dim(codes))) dim(codes) <- c(length(unique(x1)), length(unique(x2)))`
5. If `is.null(rownames(codes))`, set as follows: If `nrow(codes) == length(unique(x1))`, `rownames(codes) <- unique(x1)`. Else, if `nrow(codes) = max(x1)`, set `rownames(codes) <- seq(1, max(x1))`. Else throw an error. Ditto for `colnames, ncol, and x2`.
6. `codes[x1, x2]`

Value

a vector of the same length as `x1` and `x2`.

Author(s)

Spencer Graves

See Also

[dim rownames link{colnames}](#)

Examples

```

contrib <- c(-1, 0, 0, 1)
contrib0 <- c(FALSE, FALSE, TRUE, FALSE)

contribCodes <- recode2(contrib>0, contrib0,
  c('returned', 'received', '0', 'ERR') )

cC <- c('returned', 'returned', '0', 'received')

all.equal(contribCodes, cC)

```

subNonStandardCharacters

sub nonstandard characters with replacement

Description

Find the first and last character not in standardCharacters and replace all between them with replacement. For example, a string like "Ruben" where "e" carries an accent and is mangled by some software would become something like "Rub_n" using the default values for standardCharacters and replacement.

Usage

```

subNonStandardCharacters(x,
  standardCharacters=c(letters, LETTERS, ' ', '.', ',', '0:9',
    '\'", "\"", '-', '_', '(', ')', '[', ']', '\n'),
  replacement='_',
  gsubList=list(list(pattern='\\\\\\\\\\\\\\\\|\\\\\\\\\\\\\\\\',
    replacement='\"')),
  ... )

```

Arguments

x	character vector in which it is desired to find the first and last character not in standardCharacters and replace that substring by replacement.
standardCharacters	a character vector of acceptable characters to keep.
replacement	a character to replace the substring starting and ending with characters not in standardCharacters.
gsubList	list of lists of pattern and replacement arguments to be called in succession before looking for nonStandardCharacters
...	optional arguments passed to strsplit

Details

1. for(il in 1:length(gsubList))x <- gsub(gsubList[[il]][["pattern"]], gsublist[[il]][['replacement']], x)
2. nx <- length(x)
3. x. <- strsplit(x, "", ...)
4. for(ix in 1:nx) find the first and last standardCharacters in x.[ix] and substitute replacement for everything in between.

Value

a character vector with everthing between the first and last character not in standardCharacters replaced by replacement.

Author(s)

Spencer Graves

See Also

[sub](#), [strsplit](#), [grepNonStandardCharacters](#), [subNonStandardNames](#) [encoded_text_to_latex](#)
[subNonStandardNames](#)

Examples

```
# Consider Names = Ruben, Avila and Jose, where "e" and "A" in
# these examples carry an accent. With the default values
# for standardCharacters and replacement, these would become
# Rub_en, _vila, and Jos_.
# (The standard checks for R packages complains about
# non-standard characters, so none are included here.)
#
Names <- c('Ra`l', 'Ra`', '`l', 'Torres, Raul',
           "Robert C. \\Bobby\\\\"")
# confusion in character sets can create
# names like Names[2]
Name2 <- subNonStandardCharacters(Names)

Name2. <- c('Ra_l', 'Ra_', '_l', Names[4],
           'Robert C. "Bobby"')

all.equal(Name2, Name2.)
```

subNonStandardNames *sub for nonstandard names*

Description

sub(nonEnglishData["nonEnglish"], nonEnglishData["English"], x)

Usage

```
subNonStandardNames(x,
  standardCharacters=c(letters, LETTERS, ' ', '.', ',', ' ', 0:9,
    '\\"', "\\'", '-', '_ ', '(', ')', '[', ']', '\\n'),
  replacement='_ ',
  gsubList=list(list(pattern='\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\',
    replacement='\\'')),
  removeSecondLine=TRUE,
  nonStandardNames=Ecdat::nonEnglishNames, ...)
```

Arguments

x character vector in which it is desired replace nonStandardNames[, 1] in subNonStandardCharacters(x, ...) with the corresponding element of nonStandardNames[, 2].

standardCharacters, replacement, gsubList, ...
arguments passed to [subNonStandardCharacters](#)

removeSecondLine
logical: If TRUE, delete anything following "\\n" and return it as an attribute "secondLine"

nonStandardNames
data.frame or character matrix with two columns: Replace any substring of x matching nonStandardNames[, 1] with the corresponding element of nonStandardNames[, 2]

Details

1. removeSecondLine
2. x <- subNonStandardCharacters(x, standardCharacters, replacement, ...)
3. Loop over all rows of nonStandardNames substituting anything matching nonEnglishData[i, 1] with nonEnglishData[i, 2].
4. Eliminate leading and trailing blanks.

Value

a character vector with all nonStandardCharacters replaced first by replacement and then by the second column of nonStandardNames for any that match the first column.

Author(s)

Spencer Graves

See Also

[sub nonEnglishNames](#) [subNonStandardCharacters](#)

Examples

```
Names <- c('Raul', 'Ra`l', 'Torres,Raul', 'Torres, Raul',
           "Robert C. \\Bobby\\\\" , 'Ed \n --Vacancy')
# confusion in character sets can create
# names like Names[2]

library(Ecdat)

Name2 <- subNonStandardNames(Names)
Name2

Name2. <- c('Raul', 'Rau1', Names[3:4],
            'Robert C. "Bobby"', 'Ed')
attr(Name2., 'secondLine') <- c(rep(NA, 5), ' --Vacancy')
Name2.

all.equal(Name2, Name2.)
```

testURLs

Test URLs for intermittent download problems

Description

***NOTE: THIS IS A PRELIMINARY VERSION OF THIS FUNCTION; ***NOTE: IT MAY BE CHANGED OR REMOVED IN A FUTURE RELEASE.

try(getURL(...)) to read each element of `urls`. After each try, write a row to `file`. indicating which of `urls` was tested, the test time in seconds, and any error message. Repeat any failures up to `maxFail` times. After testing each element of `urls` once, repeat `n` times.

If(`ping`), precede each test with "ping `url[i]`". NOTE: Some Internet Service Providers seem to block some attempts to use "ping" or return fraudulent replies to "ping". It is included in the code, because it seemed like an obvious test. However, it is not executed by default because the results do not necessarily reflect what people might expect from "ping".

Return a list of the last successful version read if any from each element of `urls` with two attributes: (1) "urls" containing the `urls` argument. (2) "testResults" being an object of class `c('testURLs', 'data.frame')` of the test results written to `file`.

This function was written to diagnose a download problem with a particular Internet Service Provider (ISP). For other tools for testing an ISP, see measurementlab.net or the "Test your ISP" software discussion by the Electronic Frontier Foundation at the URL mentioned in references below.

Usage

```
testURLs(urls=c(
  wiki="http://en.wikipedia.org",
  wiki.PVI="http://en.wikipedia.org/wiki/Cook_Partisan_Voting_Index",
  house="http://house.gov",
  house.reps="http://house.gov/representatives"),
  file.='testURLresults.csv',
  n=10, maxFail=10, warn=-1, tzzone='GMT', ping=FALSE, ...)
```

Arguments

urls	a character vector assumed to be universal resource locators to pass to getURL for testing. The default was selected to provide a 2 x 2 experiment with two different web sites (en.wikipedia.org and house.gov) vs. the landing page and a subordinate page for each site.
file.	Name of a CSV file to which to write the results. If the file already exists, new results are appended to it.
n	number of times to repeat the cycle testing each member of urls.
maxFail	max tests for a continually failing URL. This is designed to make it relatively easy to determine dependencies between failures. If the failure rate is constant, the number of consecutive failures will follow a Poisson distribution. Otherwise, it may be possible to evaluate various effects using, e.g., state space techniques for non-normal time series. This could include daily and weekly cycles possibly with holiday effects and trends as well as drifts suggesting abnormal drifts in web traffic congestion.
warn	warn argument to pass to Ping .
tzzone	Time zone for Time. Defaults to GMT (UTC). tzzone=NULL will use the current locale.
ping	logical: TRUE to include Ping , FALSE otherwise.
...	optional arguments for Ping .

Details

```
for(i in 1:n):
```

```
1. pingi <- Ping(urls[i], ...)
```

```
2. The time for each call to getURL is computed by computing start.time <- proc.time() before calling try(getURL(.)), then computing the following after:
```

```
elapsed.time <- max(proc.time() - start.time, na.rm=TRUE)
```

After each of the urls is tested, a summary of the results is appended to file.. This includes the ping[['stats']], elapsed.time and the error message if the download failed.

The Electronic Frontier Foundation provides a table of existing software to "Test your ISP"; see the references below. This table includes a column noting whether the software is "active" (sending test traffic) or "passive" (observing the way the network treats natural traffic). The current testURLs function is "active", because it asks for a copy of the code at the indicated URL.

Value

an object of class `testURLs`, which in this case is a list of the last successful result returned by `getURL` for each element of `urls` with the following attributes:

<code>urls</code>	the <code>urls</code> argument used for this call
<code>testURLresults</code>	an object of class <code>c('testURLs', 'data.frame')</code> of the data written to file.. This has the following columns: <ul style="list-style-type: none"> • <code>Time date()</code> for the time a particular test started • <code>URL</code> the name in <code>urls</code> of the URL tested • <code>ping statistics</code> several columns with the count and stats returned by <code>Ping</code>. • <code>readTime</code> time in seconds for the attempt to read the URL (<code>getURL(urls[j])</code>) to complete. • <code>error character:</code> " if the read attempt was successful; the error message if not.

Author(s)

Spencer Graves

References

[measurementlab.net "Test your ISP" software discussion by the Electronic Frontier Foundation "active" \(sending test traffic\) or "passive" \(observing the way the network treats natural traffic\).](#)

See Also

[try getURL Ping](#)

Examples

```
# Test only 2 web sites, not the default 4,
# and test only twice, not the default 10 times:
tst <- testURLs(c(
  PVI="http://en.wikipedia.org/wiki/Cook_Partisan_Voting_Index",
  house="http://house.gov/representatives"),
  n=2, maxFail=2)

(class(tst) == 'testURLs') &&
all(names(tst) == c('PVI', 'house')) &&
all(names(attributes(tst)) ==
  c('names', 'urls', 'testURLresults', 'class'))
```

UShouse.senate *Create a list of members of the US House and Senate*

Description

Combine the output of [readUShouse](#) and [readUSsenate](#).

Usage

```
UShouse.senate(house=readUShouse(), senate=readUSsenate())
```

Arguments

house, senate [data.frames](#) as returned by the functions [readUShouse](#) and [readUSsenate](#), respectively.

Details

Convert the two into a common format and rbind.

Value

a [data.frame](#) with the following columns:

Office	A factor identifying "House" vs. "Senate", indicating whether the person is in the US House or Senate
state	A factor identifying the state using the USPS 2-letter state code (all caps)
district	"0" or "At-Large" for members of the US House representing an entire state or integers in character format indicating the district. For the Senate, this contains the "class", which codes the year of the next election for that seat is an integer multiple of 6 years after 2012, 2008, or 2010 for class "1", "2", or "3", respectively.
Party	a factor identifying the party affiliation of each representative, e.g., 'Democratic', 'Republican', 'Democratic-Farmer-Labor', 'Independent'.
surname	family name
givenname	first name with possibly a middle name, nickname, and suffix (e.g., Jr., III).

Author(s)

Spencer Graves

See Also

[readUShouse](#) [readUSsenate](#)

Examples

```
house <- readUSHouse()

USreps <- USHouse.senate(house)
```

USsenateClass

Election Class given state and surname of a US Senator

Description

For all individuals in `x` with `houseSenate == "Senate"`, look up their state and surname in the reference table `senate` and return their Class. For individuals not found in `senate`, return `x[[district]]`.

Senate classes 1, 2 and 3 have their normal elections in 6-year cycles including 2000, 2002, and 2004 (or 2012, 2008, and 2010), respectively. When vacancies occur out of cycle, the vacancy is first filled with appointment by the governor of the state, and an election to fill that seat occurs in the next even-numbered year; the class of that seat does not change.

For example, **South Carolina Senator Jim DeMint** resigned effective January 1, 2013. **South Carolina Governor Nikki Haley** appointed **Tim Scott** to serve until a special election in 2014. This is a Class 3 seat, which means that another election for that seat will occur in 2016.

Usage

```
USsenateClass(x, senate=readUSsenate(),
             Office='Office', state='state',
             surname='surname', district='district', senatePattern='^Senate')
```

Arguments

<code>x</code>	<code>data.frame</code> with character or factor columns <code>Office</code> , <code>state</code> , <code>surname</code> , and <code>district</code> .
<code>senate</code>	<code>data.frame</code> as returned by <code>readUSsenate</code> .
<code>Office</code>	name of a character or factor variable <code>x</code> in which the members of the US Senate can be identified by <code>grep(senatePattern, x[, Office])</code> .
<code>state</code>	Standard 2-letter abbreviation for the state of the US
<code>surname</code>	the name of a column of <code>x</code> containing the surname
<code>district</code>	name of a column of <code>x</code> containing the number of the district in the US House. For states with only one representative, this may be 0.
<code>senatePattern</code>	a regular expression for identifying the senators from <code>x[, Office]</code> .

Details

The current algorithm may fail if both senators in a state have the same surname.

Value

a `data.frame` with one row for each row of `x` and the following columns:

<code>incumbent</code>	logical vector: NA if <code>Office == 'house'</code> . If <code>Office == 'senate'</code> , then TRUE if <code>state:surname</code> found in senate and FALSE otherwise.
<code>District</code>	a character vector containing the desired Class for all US Senators found in senate or a guess at the Class for non-incumbents. For members of the House, this returned the previous content of <code>x[[District]]</code> .

NOTES:

1. Incumbents can be missed if the spelling of the surname is different between `x` and senate. This can occur with, for example, Spanish surnames containing an accent.
2. If one but not two incumbents is found, others are currently assigned to the class of an incumbent not found. This could be a mistake, because the person could be a previous incumbent or could have lost to the incumbent in the last election.

Author(s)

Spencer Graves

See Also

[readUSsenate](#)

Examples

```
tst <- data.frame(Office=factor(c("House", "Senate", "Senate", 'Senate')),
                 state=factor(c('SC', 'SC', 'SC', 'NY')),
                 surname=c("Jones", "DeMint", "Graham", 'Smith'),
                 district=c("9", NA, NA, NA),
                 stringsAsFactors=FALSE)
tst. <- USsenateClass(tst)

chk <- data.frame(incumbent=c(NA, FALSE, TRUE, FALSE),
                 district=c("9", "3", "2", "1 or 3"),
                 stringsAsFactors=FALSE)

all.equal(tst., chk)

##
## test with names different from the default
##
tst2 <- tst
names(tst2) <- letters[1:4]
tst2. <- USsenateClass(tst2, Office='a',
                      state='b', surname='c', district='d')

all.equal(tst., tst2.)
```


Index

*Topic **IO**

- financialCrisisFiles, 6
- Ping, 19
- read.testURLs, 27
- read.transpose, 28
- readCookPVI, 30
- readFinancialCrisisFiles, 32
- readNIPA, 34
- readUShouse, 35
- readUSSenate, 37
- readUSstateAbbreviations, 38
- testURLs, 44
- UShouse.senate, 47

*Topic **datasets**

- readFinancialCrisisFiles, 32
- readUSstateAbbreviations, 38
- UShouse.senate, 47

*Topic **manip**

- as.Date1970, 2
- asNumericDF, 3
- camelParse, 5
- grepNonStandardCharacters, 7
- match.data.frame, 9
- mergeUShouse.senate, 10
- mergeVote, 12
- parseCommas, 14
- parseDollars, 16
- parseName, 17
- recode2, 40
- subNonStandardCharacters, 41
- subNonStandardNames, 43
- USSenateClass, 48

*Topic **plot**

- qqnorm2, 21
- qqnorm2s, 24

- agrep, 9, 10
- as.Date, 3
- as.Date1970, 2
- as.numeric, 15, 16

- as.POSIXct1970, 3
- asNumericChar (asNumericDF), 3
- asNumericDF, 3
- camelParse, 5, 37, 38
- character, 22, 25
- classify, 10
- data.frame, 10–13, 15, 25, 27, 30, 37, 47–49
- dim, 40
- encoded_text_to_latex, 42
- financialCrisisFiles, 6, 32, 33
- getURL, 37–39, 45, 46
- grep, 8–10
- grepNonStandardCharacters, 7, 42
- gsub, 4, 15, 16
- identity, 18, 36
- is.na, 10
- join, 10
- legend, 25
- lines, 22
- lines.qqnorm2 (qqnorm2), 21
- make.names, 39
- match, 10
- match.data.frame, 9
- match_df, 10
- matrix, 35
- merge, 11
- mergeUShouse.senate, 10, 13
- mergeVote, 12
- missing, 9, 13
- nonEnglishNames, 44
- options, 20

parseCommas, [14](#), [16](#)
parseDollars, [14](#), [15](#), [16](#)
parseName, [17](#), [37](#), [38](#)
paste, [9](#)
Ping, [19](#), [45](#), [46](#)
plot, [22](#), [25](#), [26](#)
plot.qqnorm2 (qqnorm2), [21](#)
plot.qqnorm2s (qqnorm2s), [24](#)
points, [22](#), [25](#)
points.qqnorm2 (qqnorm2), [21](#)

qqnorm, [22](#), [25](#)
qqnorm2, [21](#), [26](#)
qqnorm2s, [22](#), [24](#)
Quotes, [4](#)

rbind, [36](#)
read.csv, [27](#), [28](#)
read.table, [29](#), [35](#)
read.testURLs, [27](#)
read.transpose, [28](#), [34](#)
read.xls, [6](#), [7](#), [32](#), [33](#)
readCookPVI, [30](#), [37](#)
readFinancialCrisisFiles, [32](#)
readHTMLTable, [37–39](#)
readLines, [29](#), [35](#)
readNIPA, [34](#)
readUShouse, [30](#), [31](#), [35](#), [38](#), [47](#)
readUSsenate, [30](#), [31](#), [37](#), [37](#), [47–49](#)
readUSstateAbbreviations, [37](#), [38](#)
recode2, [40](#)
regexpr, [8](#)
row.match, [10](#)
rownames, [40](#)

scan, [4](#)
showNonASCII, [8](#)
strsplit, [5](#), [9](#), [10](#), [18](#), [28](#), [29](#), [35](#), [41](#), [42](#)
sub, [42](#), [44](#)
subNonStandardCharacters, [8](#), [39](#), [41](#), [43](#),
[44](#)
subNonStandardNames, [37](#), [38](#), [42](#), [43](#)
system, [20](#)

testURLs, [27](#), [44](#)
tolower, [12](#)
try, [46](#)

USFinanceIndustry, [34](#)
UShouse.senate, [11](#), [37](#), [38](#), [47](#)
USSenateClass, [48](#)
USStateAbbreviations, [39](#)