Package ‘omnibus’

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Type Package
Title Helper Tools for Managing Data, Dates, Missing Values, and Text
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Maintainer Adam B. Smith <adam.smith@mobot.org>

Description An assortment of helper functions for managing data (e.g., rotating values in matrices by a user-defined angle, switching from row- to column-indexing), dates (e.g., intuiting year from messy date strings), handling missing values (e.g., removing elements/rows across multiple vectors or matrices if any have an NA), and text (e.g., flushing reports to the console in real-time).

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Author Adam B. Smith [cre, aut] (<https://orcid.org/0000-0002-6420-1659>)
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bracket

Identify values bracketing another value

Description

This function takes an ordered vector of numeric or character values and finds the pair that bracket a third value, \( x \). If \( x \) is exactly equal to one of the values in the vector, then a single value equal to \( x \) is returned. If \( x \) falls outside of the range of the vector, then the least/most extreme value of the vector is returned (depending on which side of the distribution of the vector \( x \) resides). Optionally, users can have the function return the index of the values that bracket \( x \).

Usage

\[
\text{bracket}(x, \text{by}, \text{index} = \text{FALSE}, \text{inner} = \text{TRUE}, \text{warn} = \text{FALSE})
\]

Arguments

- **x**: One or more numeric or character values.
- **by**: A vector of numeric or character values. These should be sorted (from high to low or low to high... if not, an error will result).
- **index**: Logical. If \text{FALSE} (default), then numeric values in \text{by} will be returned. If \text{TRUE}, then the index or indices of the bracketing value(s) will be returned.
- **inner**: Logical. If \text{TRUE} (default), then if \( x \) is surrounded by at least one series of repeating values, return the values (or indices) among the repeated sequence(s) closest to the value of \( x \). If \text{FALSE}, return the value(s) (or indices) among the repeated sequence(s) farthest from the value of \( x \). For example, if \text{index} = \text{TRUE}, \text{by} = c(1, 2, 2, 2, 3, 3), and \( x = 2.5 \), setting \text{inner} = \text{TRUE} will return the index of the third 2 and first 3. If \text{inner} = \text{FALSE}, then the function returns the index of the first 2 and second 3.
- **warn**: Logical. If \text{TRUE}, then warn if \( x \) is outside the range of \text{by}.

Value

If \( x \) is a single value, then the function will return a numeric vector of length 1 or 2, depending on how many values bracket \( x \). If all values of \text{by} are the same, then the median index (or value) of \text{by} is returned. If \( x \) is a vector, then the result will be a list with one element per item in \( x \) with each element having the same format as the case when \( x \) is a single value.

Examples

\[
\text{by} \leftarrow 2 \times (1:5) \\
\text{bracket}(4.2, \text{by}) \\
\text{bracket}(6.8, \text{by}) \\
\text{bracket}(3.2, \text{by}, \text{index}=\text{TRUE}) \\
\text{bracket}(c(3.2, 9.8, 4), \text{by})
\]
capIt

*Capitalize first letter of a string*

**Description**

This function capitalizes the first letter of a string or the first letters of a list of strings.

**Usage**

```r
capIt(x)
```

**Arguments**

- `x`  
  Character or character list.

**Value**

Character or character list.

**See Also**

`toupper, tolower`.

**Examples**

```r
x <- c('shots', 'were', 'exchanged at the ', 'hospital.')
capIt(x)
```
combineDf

Combine data frames with different fields using a crosswalk table

Description

This function combines multiple data frames, possibly with different column names, into a single data frame. Usually `merge` will be faster and easier to implement if the columns to be merged on have the same names, and `rbind` will always be faster and much easier if the column names match exactly.

Usage

```r
combineDf(
  ..., 
  crosswalk, 
  sep = "; ", 
  use = NULL, 
  classes = NULL, 
  verbose = FALSE
)
```

Arguments

- `...` A list of data frames. If ignored, then paths and file names of data frames can be specified in `crosswalk`.
- `crosswalk` Data frame. Column names are fields desired in the output data frame. Each row corresponds to a different data frame to join. If `...` is not used then the first column must have the paths and file names to CSV, RDS, or RData files representing data frames to join. Other than this column, the elements of each cell contain the name of the column in each data frame that coincides with the column name in the `crosswalk` table. For example, if the final output is to have a column by the name of "species" and "data frame #1" has a column named "Species" and "data frame #2" has a column named "scientificName", then the first value in `crosswalk` under its "species" column will be "Species" and the second "scientificName". More complex joining can be done using the following in cells of `crosswalk`:
  - `._` at start of value: indicates the value in the `crosswalk` table will be read as text and repeated in the output in each row (minus the initial "._"). For example, "._inspected" will repeat the string "inspected" in every row of the output corresponding to the respective data frame.
  - `c(~~~)`: This will paste together fields in source data frame named in `...` using the string specified in `sep` ("~~~" represents column names of the respective data frame). Note that the entire string must be inside a single or double quotes as in `c()` or "c()" and the columns named inside `c()` must be delineated by the other kind of quote (single if `c()` is delineated by double, and vice versa).
- NA: Repeats NA.

**sep**
Character, specifies the string to put between fields combined with the `c(~~~)` format in crosswalk.

**use**
Logical, Character, or NULL, if ... is used then this is a list of logical elements (TRUE or FALSE), or a column name of `crosswalk` with logical values indicating whether or not this particular data frame is to be collated, or NULL, in which case all data frames are used (default).

**classes**
Character or character list, specifies the classes (e.g., numeric, character) to be assigned to each column in the output table. If NULL, all classes will be assumed to be character. If just one value is listed, all columns will be set to this class. If a list, it must be the same length as the number of columns in `crosswalk` and specify the class of each column.

**verbose**
Logical, if TRUE prints extra information during execution. Useful for debugging the `crosswalk` table.

**Value**
A data frame.

**See Also**
`merge`, `rbind`

**Examples**
```r
df1 <- data.frame(x1=1:5, x2='valid', x3=letters[1:5], x4=LETTERS[1:5], x5='stuff')
df2 <- data.frame(y1=11:15, y3=rev(letters)[1:5])

crosswalk <- data.frame(
a=c('x1', 'y1'),
b=c('x2', '_valid'),
c=c('c("x3", "x4")', 'y3'),
d=c('x5', NA)
)

out <- combineDf(df1, df2, crosswalk=crosswalk)
out
```

---

**corner**

**Corner of a matrix or data frame**

**Description**

This function returns a corner of a matrix or data frame (i.e., upper left, upper right, lower left, lower right).
countDecDigits

**Usage**

`corner(x, corner = 1, size = 5)`

**Arguments**

- **x**: Data frame, matrix, or vector with same number of rows or elements as `into`.
- **corner**: Integer in the set 1, 2, 3, 4 or character in the set `topleft`, `topright`, `bottomleft`, `bottomright` or in the set `tl`, `tr`, `bl`, `br`. Indicates which corner to return. Integers 1, 2, 3 and 4 correspond to top left, top right, bottom left, and bottom right corners. The default is 1, the top left corner.
- **size**: Positive integer, number of rows and columns to return. If there are fewer columns/rows than indicated then all columns/rows are returned.

**Value**

A matrix or data frame.

**See Also**

`head`, `tail`

**Examples**

```r
x <- matrix(1:120, ncol=12, nrow=10)
x
corner(x, 1)
corner(x, 2)
corner(x, 3)
corner(x, 4)
```

---

### countDecDigits

**Calculate number of digits after a decimal place**

**Description**

This function calculates the number of digits after a decimal place. Note that trailing zeros will likely be ignored.

**Usage**

`countDecDigits(x)`

**Arguments**

- **x**: Numeric or numeric list.
cull

Force objects to have length or rows equal to the shortest

Description

This function takes a set of vectors, data frames, or matrices and removes the last values/rows so that they all have a length/number of rows equal to the shortest among them.

Usage

cull(...)

Arguments

... Vectors, matrices, or data frames.

Value

List objects with one element per object supplied as an argument to the function.

Examples

a <- 1:10
b <- 1:20
c <- letters
ull(a, b, c)
x <- data.frame(x=1:10, y=letters[1:10])
y <- data.frame(x=1:26, y=letters)
cull(x, y)
**dirCreate**

*Replacement for dir.create()*

---

### Description

This function is a somewhat friendlier version of `dir.create` in that it automatically sets `recursive=TRUE` and `showWarnings=FALSE` arguments.

### Usage

```r
dirCreate(...)  
```

### Arguments

- `...` Character string(s). The path and name of the directory to create. Multiple strings will be pasted together into one path, although slashes will not be pasted between them.

### Value

Nothing (creates a directory on the storage system).

### See Also

- `dir.create`

---

**domLeap**

*Day of month for leap years*

---

### Description

Data frame of day of month for each month in a leap year.

### Usage

```r
domLeap
```

### Format

An object of class 'data.frame'.

### Examples

```r
data(domLeap)
domLeap
```
domNonLeap  Day of month for non-leap years

Description
Data frame of day of month for each month in a non-leap year.

Usage
domNonLeap

Format
An object of class 'data.frame'.

Examples
data(domNonLeap)
domNonLeap

doyLeap  Day of year for leap years

Description
Data frame of day of year for each month in a leap year.

Usage
doyLeap

Format
An object of class 'data.frame'.

Examples
data(doyLeap)
doyLeap
**doyNonLeap**

| doyNonLeap | Days of year for non-leap years |

**Description**

Data frame of days of year for each month in a non-leap year.

**Usage**

doyNonLea

**Format**

An object of class 'data.frame'.

**Examples**

data(doyNonLeap)
doyNonLeap

---

**ellipseNames**

*Get names of objects input as arguments in ellipse (...)* form

**Description**

This function returns the names of objects input into a function as ellipses. It is only useful if called inside a function.

**Usage**

ellipseNames(...)

**Arguments**

... Objects.

**Value**

Character list.
eps

The smallest machine-readable number

Description
This function returns the smallest machine-readable number (equal to .Machine$double.eps).

Usage
eps()

Value
Numeric value.

Examples
eps()

insertCol

Insert a column or row into a data frame or matrix

Description
This function inserts one or more columns or rows before or after another column or row in a data frame or matrix. It is similar to `cbind` except that the inserted column(s)/row(s) can be placed anywhere.

Usage
insertCol(x, into, at = NULL, before = TRUE)
insertRow(x, into, at = NULL, before = TRUE)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>x</td>
<td>Data frame, matrix, or vector with same number of columns or rows or elements as into.</td>
</tr>
<tr>
<td>into</td>
<td>Data frame or matrix into which x is to be inserted.</td>
</tr>
<tr>
<td>at</td>
<td>Character, integer, or NULL. Name of column or column number or name of row or row number at which to do insertion. If NULL (default), the result is exactly the same as <code>cbind(into, x</code> except that it retains row numbers or column names from into.</td>
</tr>
<tr>
<td>before</td>
<td>Logical, if TRUE (default) then the insertion will occur in front of the column or row named in at, if FALSE then after. Ignored if at is NULL.</td>
</tr>
</tbody>
</table>
**isLeapYear**

Value

A data frame.

Functions

- `insertRow`: Insert a column or row into a data frame or matrix

See Also

`merge`, `cbind`, `insertRow`

Examples

```
x <- data.frame(y1=11:15, y2=rev(letters)[1:5])
into <- data.frame(x1=1:5, x2='valid', x3=letters[1:5], x4=LETTERS[1:5], x5='stuff')

insertCol(x, into=into, at='x3')
insertCol(x, into=into, at='x3', before=FALSE)
insertCol(x, into)

x <- data.frame(x1=1:3, x2=LETTERS[1:3])
into <- data.frame(x1=11:15, x2='valid')
row.names(into) <- letters[1:5]

insertRow(x, into=into, at='b')
insertRow(x, into=into, at='b', before=FALSE)
insertRow(x, into)
```

---

**isLeapYear**

Is a year a leap year?

Description

Returns `TRUE` if the year is a leap year. You can use "negative" years for BCE.

Usage

`isLeapYear(x)`

Arguments

- `x` Integer or vector of integers representing years.

Value

Vector of logical values.
Examples

isLeapYear(1990:2004) # note 2000 *was* not a leap year
isLeapYear(1896:1904) # 1900 was *not* a leap year

isTRUENA: Vectorized test for truth robust to NA

Description

These functions work exactly the same as `x == TRUE` and `x == FALSE` but by default return FALSE for cases that are NA.

Usage

isTRUENA(x, ifNA = FALSE)

isFALSENA(x, ifNA = FALSE)

Arguments

- **x**: Single value or condition or a vector of values to evaluate.
- **ifNA**: Logical, value to return if the result of evaluating `x` is NA. Note that this can be anything (i.e., TRUE, FALSE, a number, etc.).

Value

Logical or value specified in `ifNA`.

Functions

- **isFALSENA**: Vectorized test for truth robust to NA

See Also

isTRUE, isFALSE, TRUE, logical

Examples

```r
x <- c(TRUE, TRUE, FALSE, NA)
x == TRUE
isTRUENA(x)
x == FALSE
isFALSENA(x)
isTRUENA(x, ifNA = Inf)
# note that isTRUE and isFALSE are not vectorized
isTRUE(x)
isFALSE(x)
```
listFiles

Replacement for list.files()

Description
This function is a slightly friendlier version of list.files in that it automatically includes the full.names=TRUE argument.

Usage
listFiles(x, ...)

Arguments

x 	Path name of folder containing files to list.
...	Arguments to pass to list.files (other than full.names).

Value
Character list.

See Also
list.files

Examples
# list files in location where R is installed
listFiles(R.home())
listFiles(R.home(), pattern='README')

longRun

Length of the longest run of a particular value in a numeric vector

Description
This function returns the length of the longest run of a particular numeric value in a numeric vector. A "run" is an uninterrupted sequence of the same number. Runs can be "wrapped" so that if the sequence starts and ends with the target value then it is considered as a consecutive run.

Usage
longRun(x, val, wrap = FALSE, na.rm = FALSE)
Arguments

- **x**: Numeric vector.
- **val**: Numeric. Value of the elements of `x` of which to calculate length of longest run.
- **wrap**: Logical. If `TRUE` then runs can "wrap" from the end of `x` to the start of `x` if the first and last elements of `x` are equal to `val`.
- **na.rm**: Logical. If `TRUE` then remove NAs first.

Value

Integer.

See Also

- `base::rle()`

Examples

```r
x <- c(1, 1, 1, 2, 2, 3, 4, 5, 6, 1, 1, 1, 1, 1)
longRun(x, 2)
longRun(x, 1)
longRun(x, 1, wrap=TRUE)
```

Description

Consider an ordered set of values, say 1, 4, 0, 0, 0, 2, 0, 10. We can ask, what is the number of times in which zeroes appear successively? In this example, we have one set of three continuous zeros, and one set of a single zero. So the number of runs with zero is 2, and the maximum run length is 3. This function calculates the number of runs based on a certain condition for defining the run. The condition is stated as a function that returns a logical value. The function for this example would be `function(x) x == 0`.

Usage

```r
maxRuns(x, fx, args = NULL, failIfAllNA = FALSE)
```

Arguments

- **x**: Vector of numeric, character, or other values.
- **fx**: A function that returns `TRUE`, `FALSE`, or (optionally) `NA`. The function must use `x` as its first argument. For example, `function(x) x == 0` is allowable, but `function(y) y == 0` is not. Values that count as `TRUE` will be counted toward a run.
- **args**: A `list` object with additional arguments to supply to the function `fx`.
- **failIfAllNA**: If `TRUE`, fail if all values are `NA` after being evaluated by `fx`. 
**Value**

Lengths of successive runs of elements that meet the criterion. A single value of 0 indicates no conditions meet the criterion.

**Examples**

```r
x <- c(1, 4, 0, 0, 0, 2, 0, 10)
fx <- function(x) x == 0
maxRuns(x, fx)

fx <- function(x) x > 0
maxRuns(x, fx)

fx <- function(x) x > 0 & x < 5
max Runs(x, fx)

x <- c(1, 4, 0, 0, 0, 2, 0, 10)
fx <- function(x, th) x == th
maxRuns(x, fx, args=list(th=0))

# "count" NA as an observation
x <- c(1, 4, 0, 0, 0, NA, 0, 10)
fx <- function(x, th) ifelse(is.na(x), FALSE, x == th)
maxRuns(x, fx, args=list(th=0))

# include NAs as part of a run
x <- c(1, 4, 0, 0, 0, NA, 0, 10)
fx <- function(x, th) ifelse(is.na(x), TRUE, x == th)
maxRuns(x, fx, args=list(th=0))
```

---

**memUse**

Size of objects taking most memory use

**Description**

Displays the largest objects in `memUse`.

**Usage**

```r
memUse(
  n = 10,
  orderBy = "size",
  decreasing = TRUE,
  pos = 1,
  display = TRUE,
  ...
)
```
Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>n</td>
<td>Positive integer, maximum number of objects to display.</td>
</tr>
<tr>
<td>orderBy</td>
<td>Either 'size' (default) or 'name'.</td>
</tr>
<tr>
<td>decreasing</td>
<td>Logical, if TRUE (default), objects are displayed from largest to smallest.</td>
</tr>
<tr>
<td>pos</td>
<td>Environment from which to obtain size of objects. Default is 1. See ls.</td>
</tr>
<tr>
<td>display</td>
<td>If TRUE (default), print a table with memUse used.</td>
</tr>
</tbody>
</table>

Value

Data frame (invisible).

Examples

```r
memUse()
memUse(3)
```

mergeLists

Merge two lists with precedence

Description

This function merges two lists to create a single, combined list. If there is a conflict (e.g., two elements have the same name), items in the second list gain preference. Adapted from Stack Overflow (http://stackoverflow.com/questions/13811501/r-merge-lists-with-overwrite-and-recursion).

Usage

mergeLists(list1, list2)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>list1</td>
<td>List object.</td>
</tr>
<tr>
<td>list2</td>
<td>List object.</td>
</tr>
</tbody>
</table>

Value

List object.

Examples

```r
list1 <- list(a=1:3, b='Hello world', c=LETTERS[1:3])
list2 <- list(x=4, b='Goodbye world', z=letters[1:2])
mergeLists(list1, list2)
mergeLists(list2, list1)
list3 <- list(m=list(n=4:7, o=pi), a=1:5)
mergeLists(list1, list3)
```
Description

This function creates a "mirror" image of a character string, a number, a matrix, or a data frame. For example "Shots were exchanged at the hospital" becomes "latipsoh eht ta degnahcxe ewr stohS" and 3.14159 becomes 95141.3. Data frames and matrices will be returned with the order of columns or order of rows reversed.

Usage

mirror(x, direction = "lr")

Arguments

x Numeric or character, a vector of numeric or character values, or a matrix or data frame.
direction Only used if x is a matrix or data frame. Accepted values are 'lr' (left-right mirror) or 'ud' (up-down mirror).

Value

Object with same class as x.

Examples

x <- 'Shots were exchanged at the hospital'
mirror(x)

x <- c('Water', 'water', 'everywhere')
mirror(x)

# last value will return NA because the exponentiation does not
# make sense when written backwards
x <- c(3.14159, 2.71828, 6.02214076e+23)
mirror(x)
x <- data.frame(x=1:5, y=6:10)
mirror(x)

x <- matrix(1:10, nrow=2)
mirror(x)
### mmode

<table>
<thead>
<tr>
<th>Modal value(s)</th>
</tr>
</thead>
</table>

**Description**

Modal value. If there is more than one unique mode, all modal values are returned.

**Usage**

```
mmode(x)
```

**Arguments**

- `x`: Numeric or character vector.

**Value**

Numeric or character vector.

**Examples**

```
x <- c(1, 2, 3, 3, 4, 5, 3, 1, 2)
mmode(x)
```

```
x <- c(1, 2, 3)
mmode(x)
```

---

### naCompare

**Compare values using <, <=, >, >=, !=, and == (robust to NAs)**

**Description**

This function and set of operators perform simple (vectorized) comparisons using `<`, `<=`, `>`, `>=`, `!=`, or `==` between values and always returns TRUE or FALSE. TRUE only occurs if the condition can be evaluated and it is TRUE. FALSE is returned if the condition is FALSE or it cannot be evaluated.

**Usage**

```
naCompare(op, x, y)
```

```
x %<na% y
```

```
x %<=na% y
```

```
x %==na% y
```
naCompare

x %!:=na% y
x %:=na% y
x %>=na% y

Arguments

op Character, the operation to perform: '>', '<', '<=', '>', '>=', '!', or '=='. Note this must be a character (i.e., put it in quotes).

x, y Vectors of numeric, character, NA, and/or NaN values. This is the first value in the operation x XXX y where XXX is the operator in op. If x is shorter than y then x is recycled.

Value

Vector of logical values.

Examples

naCompare('<', c(1, 2, NA), c(10, 1, 0))
naCompare('<', c(1, 2, NA), 10)
naCompare('<', c(1, 2, NA), NA)

# compare to:
NA < 5
NA < NA

# same operations with operators:
1 %<na% 2
1 %<na% NA
3 %==na% 3
NA %==na% 3
4 %!=na% 4
4 %!=na% NA
5 %>=na% 3
5 %>=na% NA
3 %==na% c(NA, 1, 2, 3, 4)

# compare to:
1 < 2
1 < NA
3 == 3
NA == 3
4 != 4
4 != NA
5 >= 3
5 >= NA
3 == c(NA, 1, 2, 3, 4)
naOmitMulti

*Remove NAs from one or more equal-length vectors*

**Description**

This function removes elements in one or more equal-length vectors in which there is one NA at that position. For example, if there are three vectors A, B, and C, and A has an NA in the first position and C has an NA in the third position, then A, B, and C will each have the elements at positions 1 and 3 removed.

**Usage**

```r
naOmitMulti(...)```

**Arguments**

- `...`: Vectors (numeric or character).

**Value**

List of objects of class `...`.

**See Also**

`na.omit`

**Examples**

```r
a <- c(NA, 'b', 'c', 'd', 'e', NA)
b <- c(1, 2, 3, NA, 5, NA)
c <- c(6, 7, 8, 9, 10, NA)
naOmitMulti(a, b, c)
```

---

naRows

*Index of rows in a data frame or matrix that contain at least one NA*

**Description**

This function returns the row number of any row in a data frame or matrix that has at least one NA. This is the same as `which(!complete.cases(x))`.

**Usage**

```r
naRows(x, inf = FALSE, inverse = FALSE)```
Arguments

- **x**: Data frame or matrix.
- **inf**: Logical, if TRUE then also return row numbers of rows in which at least one element is Inf or -Inf. The default is FALSE.
- **inverse**: Logical, if TRUE then return row numbers of rows that do not have NAs (and possibly Inf or -Inf). The default is FALSE.

Value

Integer vector.

Examples

```r
x <- data.frame(a=1:5, b=c(1, 2, NA, 4, 5), c=c('a', 'b', 'c', 'd', NA))
narows(x)
```

Description

This package contains a set of helper functions.

Details

Create an issue on GitHub.

Data manipulation

- **bracket**: Find values that bracket a given number.
- **combineDf**: Combine data frames with different schema.
- **corner**: Corner of a matrix or data frame.
- **cull**: Force vectors or matrices/data frames to have the same length or number of rows.
- **insertCol** and **insertRow**: Insert column(s)/row(s) in a matrix or data frame.
- **maxRuns**: Maximum number of continuous "runs" of values meeting a particular condition.
- **mirror**: Flip an object left-right (or up-down).
- **mmode**: Modal value(s).
- **mergeLists**: Merge lists.
- **rotateMatrix**: Rotate a matrix.
- **roundTo**: Rounds a value to the nearest target value.
- **rowColIndexing**: Convert between row and column indexing of a matrix.
- **stretchMinMax**: Rescale values to a given range.
Dates

- `domLeap`: Data frame of days of each month in a leap year.
- `domNonLeap`: Data frame of days of each month in a non-leap year.
- `doyLeap`: Data frame of days of year in a leap year.
- `doyNonLeap`: Data frame of days of year in a non-leap year.
- `isLeapYear`: Is a year a leap year?
- `yearFromDate`: Attempt to find the year across dates with non-standard formats.

Geometry

- `pairDist`: Pairwise Euclidean distance between two sets of points.
- `quadArea`: Area of a quadrilateral.

Handling NAs

- `%<=na%`, `%==na%`, `%!=na%`, and `%>=na%`: Comparative operations (`>`, `>=`, `==`, `!=`, `<`, `<=`) but returns FALSE for NA cases (versus NA).
- `isTRUENA` and `isFALSENA`: Logical operators robust to NA.
- `naCompare`: Comparative operations (`>`, `>=`, `==`, `!=`, `<`, `<=`) but returns FALSE for NA cases (versus NA).
- `naOmitMulti`: Remove elements of multiple vectors if at least one element is NA or rows of matrices/data frames if at least one row has an NA.
- `naRows`: Indices of rows with at least one NA. Same as `which(!complete.cases(x))`.

Data properties

- `countDecDigits`: Count number of digits after a decimal.
- `longRun`: Longest run of a given sequence in a vector.
- `roundedSigDigits`: Infers the number of significant digits represented by a decimal representation of a division operation or digits place to which an integer was rounded.
- `which.pmax` and `which.pmin`: Combine `which.max` with `pmax`, and `which.min` with `pmin` (vectorized `which.max` and `which.min`).

System

- `dirCreate`: Nicer version of `dir.create`.
- `eps`: Smallest floating point value your computer can think of.
- `listFiles`: Nicer version of `list.files`.
- `memUse`: Display largest objects in memory.

Text

- `capIt`: Capitalize first letter of a string.
- `prefix`: Add repeating character to a string to ensure it has a user-defined length (e.g., 7 → `007`).
**pairDist**

say: Replacement for print('abc'); flush.console.

**Author(s)**

Adam B. Smith

---

**pairDist**

*Calculate pairwise distances between two matrices or data frames.*

**Description**

This function takes two data frames or matrices and returns a matrix of pairwise Euclidean distances between the two.

**Usage**

`pairDist(x1, x2, na.rm = FALSE)`

**Arguments**

- `x1`: Data frame or matrix one or more columns wide.
- `x2`: Data frame or matrix one or more columns wide.
- `na.rm`: Logical, if TRUE then any rows in `x1` or `x2` with at least one NA are removed first.

**Value**

Matrix with `nrow(x1)` rows and `nrow(x2)` columns. Values are the distance between each row of `x1` and row of `x2`.

**See Also**

`dist`

**Examples**

```r
x1 <- data.frame(x1=1:20, x2=round(100 * rnorm(20)))
x2 <- data.frame(x1=sample(1:30, 30), x2=sort(round(100 * rnorm(30))))
pairDist(x1, x2)
```
prefix  

*Add leading characters to a string*

**Description**

Add leading characters to a string. This function is useful for ensuring, say, files get sorted in a particular order. For example, on some operating systems a file name "file 1" would come first, then "file 10", then "file 11", "file 12", etc., then "file 2", "file 21", and so on. Using `prefix`, you can add one or more leading zeros so that file names are as "file 01", "file 02", "file 03", and so on... and they will sort that way.

**Usage**

`prefix(x, len, pad = "0")`

**Arguments**

- **x** Character or character list to which to add a prefix.
- **len** The total number of characters desired for each string. If a string is already this length or longer then nothing will be prefixed to that string.
- **pad** Character. Symbol to prefix to each string.

**Value**

Character or character vector.

**Examples**

```r
prefix(1:5, len=2)
prefix(1:5, len=5)
prefix(1:5, len=3, pad="!")
```

---

**quadArea**

*Area of a quadrilateral*

**Description**

Calculates the area of a quadrilateral by dividing it into two triangles and applying Heron's formula.

**Usage**

`quadArea(x, y)`
rotateMatrix

Arguments

x Numeric list. x coordinates of quadrilateral.
y Numeric list. y coordinates of quadrilateral.

Value

Numeric (area of a quadrilateral in same units as x and y).

Examples

x <- c(0, 6, 4, 1)
y <- c(0, 1, 7, 4)
quadArea(x, y)
plot(1, type="n", xlab="x", ylab="y")
polygon(x, y)
text(x, y, LETTERS[1:4], pos=4)
lines(x[1:3], y[1:3], lty="dashed", col="red")

rotateMatrix

Rotate values in a matrix

Description

This function rotates the values in a matrix by a user-specified number of degrees. In almost all cases some values will fall outside the matrix so they will be discarded. Cells that have no rotated values will become NA. Only square matrices can be accommodated. In some cases a rotation will cause cells to have no assigned value because no original values fall within them. In these instances the mean value of surrounding cells is assigned to the cells with missing values. If the angle of rotation is too small then no rotation will occur.

Usage

rotateMatrix(x, rot)

Arguments

x Object of class matrix.

rot Numeric. Number of degrees to rotate matrix. Values represent difference in degrees between "north" (up) and the clockwise direction.

Value

A matrix.

See Also

[base::t()]
Examples

```r
x <- matrix(1:100, nrow=10)
x
rotateMatrix(x, 90) # 90 degrees to the right
rotateMatrix(x, 180) # 180 degrees to the right
rotateMatrix(x, 45) # 45 degrees to the right
rotateMatrix(x, 7) # slight rotation
rotateMatrix(x, 5) # no rotation because angle is too small
```

roundedSigDigits

### Description

This function "examines" a numeric value (typically with numbers after the decimal place) and estimates either:

- The number of significant digits of the numerator and denominator of a fraction that would (approximately) result in the given value.
- The number of digits to which an integer may have been rounded, depending on whether the input has values after the decimal place or is an integer. Negative values are treated as positive values so the negative of a number will returns the same value as its positive version. See Details for more details. *Obviously, values can appear to be rounded or repeating even when they are not!*

### Usage

```r
roundedSigDigits(x, minReps = 3)
```

### Arguments

- **x**: Numeric or numeric vector.
- **minReps**: Integer. Number of times a digit or sequence of digits that occur after a decimal place needs to be repeated to assume it represents a repeating series and thus is assumed to arise from using decimal places to represent a fraction. Default is 3. For example, if `minReps` is 3 then 0.111 would be assumed to represent a repeating value because 1 occurs three times, so -1 would be returned. However, if `minReps` is 4 then the function would assume that if the value had had four digits, the next digit would not have been a 1, so returns -3 because there are three values after the decimal place. When the penultimate digit is >5 and the last digit is equal to the penultimate digit plus 1, then the last digit counts as a repeat of the penultimate digit. So 0.067 is assumed to have two repeating 6s. If `minReps` is 0 or 1 then the function will (usually) return the negative of the total number of decimal places in the value.
**roundedSigDigits**

**Details**

For values with at least one non-zero digit after a decimal place with no repeated series of digits detected, the function simply returns the total number of digits (ignoring trailing zeros) times -1. For example:

- 0.3 returns -1 because there is just one value after the decimal.
- 0.34567 returns -5 because there are no repeats up to the 5th decimal place.
- 0.1212125 returns -7 because there are no repeats (starting from the right) up to the 7th decimal place.
- 0.111117 returns -6 because there are no repeats (starting from the right) up to the 7th decimal place.

The function takes account of rounding up:

- 0.666 might be a truncated version of 2/3. Two and three each have 1 significant digit, so the function returns -1 (1 value after the decimal).
- 0.667 also returns -1 because this might represent a rounding of 2/3 and it is customary to round digits up if the next digit would have been >5.
- 0.3334 returns -4 because it is inappropriate to round 3 up to 4 if the next digit would have been 5 or less.

Repeating series are accounted for. For example:

- 0.121212 returns -2 because "12" starts repeating after the second decimal place.
- 0.000678678678 returns -6 because "678" starts repeating after the 6th place.
- 0.678678678 returns -3.
- 0.678678679 also returns -3 because 678 could be rounded to 679 if the next digit were 6.

Note that you can set the minimum number of times a digit or series needs to be repeated to count as being repeated using the argument `minReps`. The default is 3, so digits or series of digits need to be repeated at least 3 times to count a repetition, but this can be changed:

- 0.1111 returns -1 using the default requirement for 3 repetitions but -4 if the number of minimum repetitions is 5 or more.
- 0.121212 returns -2 using the default requirement for 3 repetitions but -6 if the number of minimum repetitions is 4 or more.

Trailing zeros are ignored, so 0.12300 returns -3. When values do not have digits after a decimal place the location of the first non-zero digit from the right is returned as a positive integer. For example:

- 234 returns 1 because the first non-zero digit from the right is in the 1s place.
- 100 returns 3 because the first non-zero digit from the right is in the 100s place.
- 70001 returns 1 because the first non-zero digit from the right is in the 1s place.

However, note a few oddities:

- 4E5 returns 6 but 4E50 probably will not return 51 because many computers have a hard time internally representing numbers that large.
• 4E-5 returns -5 but probably will not return -50 because many computers have a hard time internally representing numbers that small.
• -100 and 100 return 3 and -0.12 and 0.12 return -2 because the negative sign is ignored.
• 0 returns 0.
• NA and NaN returns NA.

Value

Integer (number of digits) or NA (does not appear to be rounded).

Examples

roundedSigDigits(0.3)
roundedSigDigits(0.34567)
roundedSigDigits(0.1212125)
roundedSigDigits(0.111117)
roundedSigDigits(0.666)
roundedSigDigits(0.667)
roundedSigDigits(0.3334)
roundedSigDigits(0.121212)
roundedSigDigits(0.000678678678)
roundedSigDigits(0.678678678)
roundedSigDigits(0.678678679)
roundedSigDigits(0.1111)
roundedSigDigits(0.1111, minReps=5)
roundedSigDigits(0.121212)
roundedSigDigits(0.121212, minReps=4)
roundedSigDigits(234)
roundedSigDigits(100)
roundedSigDigits(70001)
roundedSigDigits(4E5)
roundedSigDigits(4E50)
roundedSigDigits(4E-5)
roundedSigDigits(4E-50)
roundedSigDigits(0)
roundedSigDigits(NA)

x <- c(0.0667, 0.0667, 0.067)
roundedSigDigits(x)

roundTo  

\textit{Round to nearest target value}

Description

This function rounds a value to a nearest "target" value (e.g., you could round 0.72 to the nearest 0.25, or 0.75).
rowColIndexing

Usage

roundTo(x, target, roundFx = round)

Arguments

x Numeric.
target Numeric.
roundFx Any of round, floor, or ceiling.

Value

Numeric.

Examples

roundTo(0.73, 0.05)
roundTo(0.73, 0.1)
roundTo(0.73, 0.25)
roundTo(0.73, 0.25, floor)
roundTo(0.73, 1)
roundTo(0.73, 10)
roundTo(0.73, 10, ceiling)

rowColIndexing Convert between row- and column-style indexing of matrices

Description

These functions converts index values of cells between row- and column-style indexing of cells in matrices. Column indexing (the default for matrices) has the cell "1" in the upper left corner of the matrix. The cell "2" is below it, and so on. The numbering then wraps around to the top of the next column. Row indexing (the default for rasters, for example), also has cell "1" in the upper left, but cell "2" is to its right, and so on. Numbering then wraps around to the next row.

Usage

rowColIndexing(x, cell, dir)

Arguments

x Either a matrix or a vector with two values, one for the number of rows and one for the number of columns in a matrix.
cell One or more cell indices (positive integers).
dir The "direction" in which to convert. If 'row', it is assumed that cell is a column-style index and so should be converted to a row-style index. If 'col', it is assumed that cell is a row-style index and so should be converted to a column-style index.
Value

One or more positive integers.

Examples

# column versus row indexing
colIndex <- matrix(1:40, nrow=5, ncol=8)
rowIndex <- matrix(1:40, nrow=5, ncol=8, byrow=TRUE)
print(colIndex)
print(rowIndex)

# examples
x <- matrix('a', nrow=5, ncol=8, byrow=TRUE)
rowColIndexing(x, cell=c(1, 6, 20), 'row')
rowColIndexing(x, cell=c(1, 6, 20), 'col')

rowColIndexing(c(5, 8), cell=c(1, 6, 20), 'row')
rowColIndexing(c(5, 8), cell=c(1, 6, 20), 'col')

say

Nicer version of print() or cat() function

Description

This function is a nicer version of print() or cat(), especially when used inline for functions because it displays immediately and pastes all strings together. It also does some rudimentary but optional word wrapping.

Usage

say(
  ..., 
  pre = 0, 
  post = 1, 
  breaks = NULL, 
  wiggle = 10, 
  preBreak = 1, 
  level = NULL, 
  deco = "#"
)

Arguments

... character strings to print
pre Integer >= 0. Number of blank lines to print before strings
post Integer >= 0. Number of blank lines to print after strings
breaks

Either NULL, which causes all strings to be printed on the same line (no wrap overflow) or a positive integer which wraps lines at this character length (e.g., breaks=80 inserts line breaks every 80 characters).

wiggle

Integer \( \geq 0 \). Allows line to overrun breaks length in characters before inserting line breaks.

preBreak

If wrapping long lines indicates how subsequent lines are indented. NULL causes lines to be printed starting at column 1 on the display device. A positive integer inserts preBreak number of spaces before printing each line. A string causes each line to start with this string.

level

Integer or NULL. If NULL, then the items in \( \ldots \) are displayed as-is. Otherwise, a value of 1, 2, or 3 indicates the heading level, with lower numbers causing more decoration and spacing to be used.

deco

Character. Character to decorate text with if level is not NULL.

Value

Nothing (side effect is output on the display device).

Examples

```r
say('The quick brown fox ', 'jumps over the lazy ', 'Susan.')
say('The quick brown fox ', 'jumps over the lazy ', 'Susan.', breaks=10)
say('The quick brown fox ', 'jumps over the lazy ', 'Susan.', level=1)
say('The quick brown fox ', 'jumps over the lazy ', 'Susan.', level=2)
say('The quick brown fox ', 'jumps over the lazy ', 'Susan.', level=3)
```

<table>
<thead>
<tr>
<th>stretchMinMax</th>
<th>Rescale values</th>
</tr>
</thead>
</table>

Description

This function rescales a vector of numeric values to an arbitrary range. Optionally, after the stretch values equal to the lowest value can be "nudged" slightly higher to half the minimum value across the rescaled vector of values \( > 0 \).

Usage

```r
stretchMinMax(
  x,
  lower = 0,
  upper = 1,
  nudgeUp = FALSE,
  nudgeDown = FALSE,
  na.rm = FALSE
)
```
Arguments

- **x**: Numeric list.
- **lower**: Numeric, low end of range to which to stretch.
- **upper**: Numeric, high end of range to which to stretch.
- **nudgeUp**, **nudgeDown**: Logical, if FALSE (default) then do nothing. If TRUE then *after* rescaling to [0, 1], a small value will be added to all values of x equal to 0. This value is equal to $0.5 \times \min(x[x > 0])$.
- **na.rm**: Logical, if FALSE (default) then if any values of x are NA then the returned value will be NA. If TRUE then NA's are ignored in calculation.

Value

Numeric value.

See Also

-scale

Examples

```r
x <- 1:10
stretchMinMax(x)
stretchMinMax(x, lower=2, upper=5)
stretchMinMax(x, nudgeUp=TRUE)
stretchMinMax(x, lower=2, upper=5, nudgeUp=TRUE)
stretchMinMax(x, nudgeDown=TRUE)
stretchMinMax(x, lower=2, upper=5, nudgeUp=TRUE, nudgeDown=TRUE)
x <- c(1:5, NA)
stretchMinMax(x)
stretchMinMax(x, na.rm=TRUE)
```

which.pmax

*Which vector has maximum value at each element*

Description

These functions are vectorized versions of which.max and which.min, which return the index of the value that is maximum or minimum (or the first maximum/minimum value, if there is a tie). In this case, the function is supplied two or more vectors of the same length. For each element at the same position (e.g., the first element in each vector, then the second element, etc.) the function returns an integer indicating which vector has the highest or lowest value (or the index of the first vector with the highest or lowest value in case of ties).
which.pmax

Usage

which.pmax(..., na.rm = TRUE)

which.pmin(..., na.rm = TRUE)

Arguments

...  Two or more vectors. If lengths do not match, the results will likely be unanticipated.

na.rm  Logical, if FALSE and any of the vectors contains an NA or NaN, the function will return an NA. If TRUE (default), then NA will only be returned if all elements at that position are NA or NaN.

Value

Vector the same length as the input, with numeric values indicating which vector has the highest value at that position. In case of ties, the index of the first vector is returned.

Functions

• which.pmin: Which vector has minimum value at each element

See Also

which.max, which.min, pmax, pmin

Examples

set.seed(123)
a <- sample(9, 5)
b <- sample(9, 5)
c <- sample(9, 5)
a[2:3] <- NA
b[3] <- NA
a[6] <- NA
b[6] <- NA
c[6] <- NA
which.pmax(a, b, c)
which.pmin(a, b, c)
which.pmax(a, b, c, na.rm=FALSE)
which.pmin(a, b, c, na.rm=FALSE)
### Description

This function attempts to return the year from characters representing dates formats. The formats can be ambiguous and varied within the same set. For example, it returns "1982" (or 9982 if century is ambiguous) from "11/20/82", "1982-11-20", "Nov. 20, 1982", "20 Nov 1982", "20-Nov-1982", "20/Nov/1982", "20 Nov 82", "20 Nov 82". The function handles ambiguous centuries (e.g., 1813, 1913, 2013) by including a dummy place holder for the century place (i.e., 9913). Note that it may return warnings like "NAs introduced by coercion".

### Usage

```r
yearFromDate(x, yearLast = TRUE)
```

### Arguments

- **x**: Character or character list, one or more dates.
- **yearLast**: Logical, if TRUE assume that dates like "XX/YY/ZZ" list the year last (=ZZ). If FALSE, assume they're first (=XX).

### Value

Numeric.

### Examples

```r
eyearFromDate(1969, yearLast=TRUE)
yearFromDate('10-Jul-71', yearLast=TRUE) # --> 9971
yearFromDate('10-Jul-1971', yearLast=TRUE) # --> 1971
yearFromDate('10-19-71', yearLast=TRUE) # --> 9971
yearFromDate('10-19-1969', yearLast=TRUE) # --> 1969
yearFromDate('10-1-71', yearLast=TRUE) # --> 9971
yearFromDate('3-22-71', yearLast=TRUE) # --> 9971
yearFromDate('3-2-71', yearLast=TRUE) # --> 9971
yearFromFile('10-1-1969', yearLast=TRUE) # --> 1969
yearFromDate('3-22-1969', yearLast=TRUE) # --> 1969
yearFromDate('3-19-1969', yearLast=TRUE) # --> 1969
yearFromFile('10/Jul/71', yearLast=TRUE) # --> 9971
yearFromFile('10/Jul/1971', yearLast=TRUE) # --> 1971
yearFromFile('10/19/71', yearLast=TRUE) # --> 9971
yearFromFile('10/19/1969', yearLast=TRUE) # --> 1969
yearFromFile('10/1/71', yearLast=TRUE) # --> 9971
yearFromFile('3/22/71', yearLast=TRUE) # --> 9971
yearFromFile('3/2/71', yearLast=TRUE) # --> 9971
yearFromFile('10/1/1969', yearLast=TRUE) # --> 1969
yearFromFile('3/22/1969', yearLast=TRUE) # --> 1969
yearFromFile('3/2/1969', yearLast=TRUE) # --> 1969
```
yearFromDate('10 1 1969', yearLast=TRUE) # --> 1969
yearFromDate('10 19 1969', yearLast=TRUE) # --> 1969
yearFromDate('3 22 1969', yearLast=TRUE) # --> 1969
yearFromDate('3 2 1969', yearLast=TRUE) # --> 1969
yearFromDate('Oct. 19, 1969', yearLast=TRUE) # --> 1969
yearFromDate('19 October 1969', yearLast=TRUE) # --> 1969
yearFromDate('How you do dat?', yearLast=TRUE) # --> NA
yearFromDate('2014-07-03', yearLast=TRUE) # --> 2014
yearFromDate('2014-7-03', yearLast=TRUE) # --> 2014
yearFromDate('2014-07-3', yearLast=TRUE) # --> 2014
yearFromDate('2014/07/03', yearLast=TRUE) # --> 2014
yearFromDate('2014/7/03', yearLast=TRUE) # --> 2014
yearFromDate('2014/07/3', yearLast=TRUE) # --> 2014
yearFromDate('2014/7/3', yearLast=TRUE) # --> 2014
yearFromDate('2014 07 03', yearLast=TRUE) # --> 2014
yearFromDate('2014 07 3', yearLast=TRUE) # --> 2014
yearFromDate('2014 7 3', yearLast=TRUE) # --> 2014
yearFromDate('2014 7 3', yearLast=TRUE) # --> 2014

yearFromDate(1969, yearLast=FALSE)
yearFromDate('10-Jul-71', yearLast=FALSE) # --> 1971
yearFromDate('10-Jul-1971', yearLast=FALSE) # --> 1971
yearFromDate('10-19-71', yearLast=FALSE) # --> 1971
yearFromDate('10-19-1969', yearLast=FALSE) # --> 1969
yearFromDate('10-1-71', yearLast=FALSE) # --> 1971
yearFromDate('3-22-71', yearLast=FALSE) # --> 1971
yearFromDate('3-2-71', yearLast=FALSE) # --> 1971
yearFromDate('3-2-1969', yearLast=FALSE) # --> 1969
yearFromDate('3-22-1969', yearLast=FALSE) # --> 1969
yearFromDate('10-19/71', yearLast=FALSE) # --> 1971
yearFromDate('10/19/71', yearLast=FALSE) # --> 1971
yearFromDate('10/1/1969', yearLast=FALSE) # --> 1969
yearFromDate('10/1/71', yearLast=FALSE) # --> 1971
yearFromDate('3/22/71', yearLast=FALSE) # --> 1971
yearFromDate('3/2/71', yearLast=FALSE) # --> 1971
yearFromDate('3/2/1969', yearLast=FALSE) # --> 1969
yearFromDate('3/2/1969', yearLast=FALSE) # --> 1969
yearFromDate('3/2 1969', yearLast=FALSE) # --> 1969
yearFromDate('3 2 1969', yearLast=FALSE) # --> 1969
yearFromDate('3 22 1969', yearLast=FALSE) # --> 1969
yearFromDate('3 2 1969', yearLast=FALSE) # --> 1969
yearFromDate('Oct. 19, 1969', yearLast=FALSE) # --> 1969
yearFromDate('19 October 1969', yearLast=FALSE) # --> 1969
yearFromDate('How you do dat?', yearLast=FALSE) # --> NA
yearFromDate('2014-07-03', yearLast=FALSE) # --> 2014
yearFromDate('2014-7-03', yearLast=FALSE) # --> 2014
yearFromDate('2014-07-3', yearLast=FALSE) # --> 2014
yearFromDate('2014-7-3', yearLast=FALSE) # --> 2014
yearFromDate('2014/07/03', yearLast=FALSE) # --> 2014
yearFromDate('2014/7/03', yearLast=FALSE) # --> 2014
yearFromDate('2014/07/3', yearLast=FALSE) # --> 2014
yearFromDate('2014 07 03', yearLast=FALSE) # --> 2014
yearFromDate('2014 7 03', yearLast=FALSE) # --> 2014
yearFromDate('2014 7 3', yearLast=FALSE) # --> 2014
yearFromDate('2014 7 3', yearLast=FALSE) # --> 2014
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