Package 'tinyoperators'

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Title Infix operators and some functions to help in your coding etiquette
Version 0.0.9
Description The 'tinyoperators' R-package adds some infix operators, and a few functions. It primarily focuses on 4 things. (1) Float truth testing. (2) Reducing repetitive code. (3) Extending the string manipulation capabilities of the 'stringi' R package. (4) A new package and module import system, that combines the benefits of aliasing a package with the benefits of attaching a package. The 'tinyoperators' R-package has only one dependency, namely 'stringi'. However, it does allows multi-threading of some of the string-related functions (when appropriate) via the suggested 'stringfish' R-package.
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float_truth_testing Safer float (in)equality operators

Description

The %f==%, %f!=% %f<%, %f>%, %f<=%, %f>=% (in)equality operator perform float truth testing. They are virtually equivalent to the regular (in)equality operators,

except for one aspect. The float truth testing operators assume that if the absolute difference between x and y is smaller than the Machine tolerance, sqrt(.Machine\$double.eps), then x and y ought to be consider to be equal.

Thus these provide safer float truth testing.

For example: 0.1*7 == 0.7 returns FALSE, even though they are equal, due to the way floating numbers are stored in programming languages like R. But 0.1*7 %f==% 0.7 returns TRUE.

There are also the $x \%f{}\%$ bnd and $x \%f{}\%$ bnd operators, where bnd is a vector of length 2, or a 2-column matrix (nrow(bnd)==length(x) or nrow(bnd)==1).

The $x \%f{}$ % bnd operator checks if x is within the closed interval with bounds defined by bnd.

The x %f!{}% bnd operator checks if x is outside the closed interval with bounds defined by bnd.

Usage

x %f==% y

x %f!=% y

x %f<% y

x %f>% y

x %f<=% y

x %f>=% y

x %f{}% bnd

x %f!{}% bnd

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Arguments

bnd

x, y numeric vectors, matrices, or arrays, though these operators were specifically designed for floats (class "double").

either a vector of length 2, or a matrix with 2 columns and 1 row, or else a matrix

with 2 columns where nrow(bnd)==length(x).

The first element/column of bnd gives the lower bound of the closed interval:

The first element/column of bnd gives the lower bound of the closed interval; The second element/column of bnd gives the upper bound of the closed interval;

```
x < -c(0.3, 0.6, 0.7)
y < -c(0.1*3, 0.1*6, 0.1*7)
print(x); print(y)
x == y \# gives FALSE, but should be TRUE
x!= y \# gives TRUE, should be FALSE
x > y \# not wrong
x < y \text{ \# gives TRUE, should be FALSE}
x %f==% y # here it's done correctly
x %f!=% y # correct
x %f<% y # correct
x %f>% y # correct
x %f<=% y # correct
x %f>=% y # correct
x \leftarrow c(0.3, 0.6, 0.7)
bnd <- matrix(c(0.29, 0.59, 0.69, 0.31, 0.61, 0.71), ncol=2)
x %f{}% bnd
x %f!{}% bnd
# These operators still work for non-float numerics also:
x <- 1:5
y < -1:5
x %f==% y
x %f!=% y
x %f<% y
x %f>% y
x %f<=% y
x %f>=% y
x <- 1:5
y <- x+1
x %f==% y
x %f!=% y
x %f<% y
x %f>% y
x %f<=% y
x %f>=% y
x <- 1:5
y <- x-1
x %f==% y
x %f!=% y
x %f<% y
x %f>% y
x %f<=% y
```

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```
x %f>=% y
```

import

Additional package import management

Description

These functions implement a new package import system, that attempts to combine the benefits of aliasing a package with the benefits of attaching a package.

import_as:

The import_as() function imports the namespace of an R package, and optionally also its dependencies, enhances, and extensions, under the same alias. The specified alias will be placed in the current environment (like the global environment, or the environment within a function).

import_inops:

The import_inops() function exposes the infix operators of the specified packages to the current environment (like the global environment, or the environment within a function).

To ensure the user can still verify which operator function came from which package, a "package" attribute is added to each exposed operator.

Naturally, the namespaces of the operators remain intact.

import_data:

The import_data() function gets a specified data set from a package.

Unlike utils::data(), the import_data() function returns the data set directly, and allows assigning the data set like so:

```
mydata <- import_data(...).</pre>
```

import_lsf:

The import_lsf(package, ...) function gets a list of exported functions/operators from a package.

Usage

```
import_as(
   alias,
   main_package,
   foreign_exports = TRUE,
   dependencies = FALSE,
   enhances = NULL,
   extensions = NULL,
   lib.loc = .libPaths(),
   loadorder = c("dependencies", "main_package", "enhances", "extensions")
)
import_inops(pkgs, lib.loc = .libPaths(), exclude, include.only)
import_data(dataname, package, lib.loc = .libPaths())
import_lsf(package, type, lib.loc = .libPaths())
```

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Arguments

alias a variable name (unquoted), giving the (not yet existing) object where the pack-

age(s) are to be assigned to.

Syntactically invalid names are not allowed for the alias name.

main_package

a single string, giving the name of the main package to load under the given

alias.

foreign_exports

logical.

Some R packages export functions that are not defined in their own package, but in their direct dependencies; "foreign exports", if you will.

If foreign_exports = TRUE these foreign exports are added to the namespace of main_package, even if dependencies = FALSE.

If foreign_exports = FALSE, these foreign exports are not added, and the user must specify the appropriate packages in argument dependencies.

Defaults to TRUE, which is similar to the behaviour of base R's :: operator.

dependencies

either logical, or a character vector.

If FALSE (default), no dependencies are loaded under the alias.

If TRUE, ALL direct dependencies of the main_package are loaded under the alias, but **excluding** base/core R, and also **excluding** pre-installed "recommended" R packages.

See also pkgs_get_deps.

If a character vector, then it is taken as the direct dependencies of the package to be loaded also under the alias.

NOTE (1): "Dependencies" here are defined as any package appearing in the "Depends", "Imports", or "LinkingTo" sections of the Description file of the main_package. So no recursive dependencies.

NOTE (2): If dependencies is a character vector: The order of the character vector matters! If multiple packages share objects with the same name, the package named last will overwrite the earlier named packages.

enhances

a character vector, giving the names of the packages enhanced by the main_package to be loaded also under the alias.

Defaults to NULL, which means no enhances are loaded.

NOTE(1): Enhances are defined as packages appearing in the "Enhances" section of the Description file of the main_package.

NOTE (2): The order of the character vector matters! If multiple packages share objects with the same name, the objects of the package named last will overwrite those of the earlier named packages.

extensions

a character vector, giving the names of the extensions / reverse-dependencies of the main_package to be loaded also under the alias.

Defaults to NULL, which means no extensions are loaded.

NOTE (1): "Extensions" here are defined as reverse-depends or reverse-imports. It does not matter if these are CRAN or non-CRAN packages.

NOTE (2): The order of the character vector matters! If multiple packages share objects with the same name, the objects of the package named last will overwrite those of the earlier named packages.

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lib.loc

character vector specifying library search path (the location of R library trees to search through).

This is usually .libPaths(). See also loadNamespace.

loadorder

the character vector

c("dependencies", "main_package", "enhances", "extensions"), or some re-ordering of this character vector, giving the relative load order of the groups of packages.

The default setting (which is highly recommended) is the character vector c("dependencies", "main_package", "enhances", "extensions"), which results in the following load order:

- (1) The dependencies, in the order specified by the dependencies argument.
- (2) The main_package (see argument main_package), including foreigg exports (if foreign_exports=TRUE).
- (3) The enhances, in the order specified by the enhances argument.
- (4) The reverse-dependencies/extensions, in the order specified by the extensions argument.

pkgs

a single string, or character vector, with the package name(s).

NOTE (1): The order of the character vector matters! If multiple packages share objects with the same name, the objects of the package named last will overwrite those of the earlier named packages.

NOTE (2): The import_inops function performs a basic check that the packages are mostly (reverse) dependencies of each other. If not, it will give an error.

exclude

a character vector, giving the infix operators NOT to expose to the current environment.

This can be handy to prevent overwriting any (user defined) infix operators already present in the current environment.

NOTE: You cannot specify both the exclude and include.only arguments. Only one or the other, or neither.

include.only

a character vector, giving the infix operators to expose to the current environment, and the rest of the operators will not be exposed.

This can be handy to prevent overwriting any (user defined) infix operators already present in the current environment.

NOTE: You cannot specify both the exclude and include.only arguments. Only one or the other, or neither.

dataname

a single string, giving the name of the data set.

package

the quoted package name.

type

The type of functions to list. Possibilities:

"inops" or "operators": Only infix operators.

"regfuns": Only regular functions (thus excluding infix operators).
"all": All functions, both regular functions and infix operators.

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Details

The import_as() function does not allow importing base/core R under an alias, so don't try. For a more detailed description of the import system introduced by the tinyoperators R package, please refer to the Read Me file on the GitHub main page:

```
https://github.com/tony-aw/tinyoperators
```

Value

```
For import_as:
```

The variable named in the alias argument will be created (if it did not already exist), in the current environment (like the global environment, or the environment within a function). The alias object will contain the (merged) package environment.

```
For import_inops():
```

The infix operators from the specified packages will be placed in the current environment (like the Global environment, or the environment within a function).

```
For import_data():
```

Returns the data directly. Thus, one can assign the data like so: mydata <- import_data(...).

```
For import_lsf():
```

Returns a character vector of function and/or operator names.

Examples

```
## Not run:
import_as( # this creates the dr object
dr, "dplyr", depends=c("tibble", "tidyselect"), extends = "powerjoin"
)
import_inops("data.table")
d <- import_data("chicago", "gamair")
head(d)
## End(Not run)</pre>
```

inplace

Generalized in-place (mathematical) modifier

Description

Generalized in-place (mathematical) modifier.

The x %:=% f operator allows performing in-place modification of some object x with a function f.

```
For example this:
```

```
object[object > 0] %:=% \(x) x + 1
```

```
Is the same as:
object[object > 0] <- object[object > 0] + 1
```

This function-based method is used instead of the more traditional in-place mathematical modification like += to prevent precedence issues (functions come before mathematical arithmetic in R).

Usage

```
x %:=% f
```

Arguments

x an object, with properties such that function f can be use on it.

For example, when function f is mathematical, x should be a number or numeric

(or 'number-like') vector, matrix, or array.

f a function to be applied in-place on x.

Value

This operator does not return any value: it is an in-place modifiers, and thus modifies x directly.

Examples

```
set.seed(1)
object <- matrix(rpois(10, 10), ncol=2)
print(object)
object %:=% \(x) x+3 # same as object <- object + 3
print(object)</pre>
```

inplace_str_arithmetic

In place modifying string arithmetic

Description

In-place modifier versions of string arithmetic:

```
x %s+ =% y is the same as x <- x %s+% y
x %s- =% p is the same as x <- x %s-% p
x %s* =% n is the same as x <- x %s*% n
x %s/ =% p is the same as x <- x %s/% p</pre>
```

See also the documentation on string arithmetic: string arithmetic.

Some of the internal code of these operators was inspired by the roperators R package.

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Usage

```
x %s+ =% y
x %s- =% p
x %s* =% n
x %s/ =% p
```

Arguments

```
x, y, p, n see string arithmetic and s_pattern.
```

pattern with ignore.case=TRUE:

p <- s_pattern(regex = "A|E|I|0|U", ignore.case=TRUE)</pre>

Value

These operators do not return any value: they are in-place modifiers, and thus modify x directly.

References

Wiseman B, Nydick S, Jones J (2022). roperators: Additional Operators to Help you Write Cleaner R Code. https://CRAN.R-project.org/package=roperators

```
y <- "a"
p \leftarrow "a|e|i|o|u"
n < -c(2, 3)
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
x %s+ =% y # same as x <- x %s+% y
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x \%s- = % p \# same as <math>x <- x \%s- \% p
print(x)
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
x %s* =% n # same as <math>x <- x %s\\*% n
print(x)
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
x \%s/ = % p # same as <math>x < - x \%s/\% p
print(x)
y <- "a"
```

inplace_str_subset

```
n < -c(3, 2)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x %s+ =% y # same as x <- x %s+% y
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x %s- =% p # same as x <- x %s-% p
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x %s* =% n # same as <math>x <- x %s\*% n
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x \%s/ = \% p \# same as x <- x \%s/\% p
print(x)
```

inplace_str_subset

In place modifying string subsetting

Description

In-place modifier versions of string subsetting:

```
x %sget =% ss is the same as x <- x %sget% ss
x %strim =% ss is the same as x <- x %strim% ss</pre>
```

See also the documentation on string subsetting (string subset). Note that there is no in-place modifier versions of %ss%.

Some of the internal code of these operators was inspired by the roperators R package.

Usage

```
x %sget =% ss
x %strim =% ss
```

Arguments

```
x, ss see string subset.
```

Value

These operators do not return any value: they are in-place modifiers, and thus modify x directly.

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References

Wiseman B, Nydick S, Jones J (2022). roperators: Additional Operators to Help you Write Cleaner R Code. https://CRAN.R-project.org/package=roperators

Examples

```
ss <- c(2,2)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
x %sget =% ss # same as x <- x %sget% ss
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
x %strim =% ss # same as x <- x %strim% ss
print(x)
ss <- c(2,2)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
x %sget =% ss # same as x <- x %sget% ss
print(x)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
x %strim =% ss # same as x <- x %strim% ss
print(x)
```

logic_ops

Logic operators

Description

Additional logic operators:

The x %xor% y operator is the "exclusive-or" operator, the same as xor(x, y).

The x n&% operator is the "not-and" operator, the same as (!x) & (!y).

The x %out% y operator is the same as !x %in% y.

The x %?=% y operator checks if x and y are **both** unreal or unknown (i.e. NA, NaN, Inf, -Inf).

The n %=numtype% numtype operator is a vectorized operator that checks for every value of numeric vector n if it can be considered a number belonging to type numtype. See arguments for details.

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The s %=strtype% strtype operator is a vectorized operator that checks for every value of character vector s

if it can seen as a certain strtype. See arguments for details.

The s %sgrep% p operator is a vectorized operator that checks for every value of character vector s if it has pattern p.

Usage

```
x %xor% y
x %n&% y
x %out% y
x %?=% y
s %sgrep% p
n %=numtype% numtype
s %=strtype% strtype
```

Arguments

x, y see Logic.

s a character vector.

p the result from s_pattern, or else a character vector of the same length as s with regular expressions.

n a numeric vector.

numtype a single string giving the type if numeric to be checked. The following options are supported:

- "~0": zero, or else a number whose absolute value is smaller than the Machine tolerance (sqrt(.Machine\$double.eps)).
- "B": binary numbers (exactly 0 or exactly 1);
- "prop": proportions numbers between 0 and 1 (exactly 0 or 1 is also allowed);
- "N": Natural numbers (non-negative integers including zero);
- "I": Integers;
- "odd": odd integers;
- "even": even integers;

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- "R": Real numbers;
- "unreal": infinity, NA, or NaN;

strtype

a single string giving the type of string to be checked. The following options are supported:

- "empty": checks if the string only consists of empty spaces.
- "unreal": checks if the string is NA, or if it has literal string "NA", "NaN" or "Inf", regardless if it has leading or trailing spaces.
- "numeric": checks if the string can be converted to a number, disregarding leading and trailing spaces. I.e. the string "5.0" can be converted to the the actual number 5.0.
- "special": checks if the string consists of only special characters.

```
x <- c(TRUE, FALSE, TRUE, FALSE, NA, NaN, Inf, -Inf, TRUE, FALSE)
y <- c(FALSE, TRUE, TRUE, FALSE, rep(NA, 6))
outcome <- data.frame(</pre>
  x=x, y=y,
  "x %xor% y"=x %xor% y, "x %n&% y" = x %n&% y, "x %?=% y" = x %?=% y,
  check.names = FALSE
print(outcome)
1:3 %out% 1:10
1:10 %out% 1:3
n \leftarrow c(0:5, 0:-5, 0.1, -0.1, 0, 1, Inf, -Inf, NA, NaN)
1e-20 %=numtype% "~0"
n[n %=numtype% "B"]
n[n %=numtype% "prop"]
n[n %=numtype% "N"]
n[n %=numtype% "I"]
n[n %=numtype% "odd"]
n[n %=numtype% "even"]
n[n %=numtype% "R"]
n[n %=numtype% "unreal"]
s <- c(" AbcZ123 ", " abc ", " 1.3 ", " !#$%^&*() ", " ", " NA ", " NaN ", " Inf ")
s[s %=strtype% "empty"]
s[s %=strtype% "unreal"]
s[s %=strtype% "numeric"]
s[s %=strtype% "special"]
s <- c("Hello world", "Goodbye world")</pre>
p <- s_pattern(regex = c("Hello", "Hello"))</pre>
```

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s %sgrep% p

matrix_ops

Infix operators for row- and column-wise re-ordering of matrices

Description

Infix operators for custom row- and column-wise re-ordering of matrices

The x %row~% mat operator re-orders the elements of every row of matrix x according to the ordering ranks given in matrix mat.

The $x \col^{\infty}$ mat operator re-orders the elements of every column of matrix x according to the ordering ranks given in matrix mat.

Usage

x %row~% mat

x %col~% mat

Arguments

x a matrix

mat a matrix with the same dimensions as x, giving the ordering ranks of every ele-

ment of matrix x.

Details

If matrix x is a numeric matrix, and one wants to sort the elements of every row or column numerically, x %row~% x or x %col~% x would suffice, respectively.

If matrix x is not numeric, sorting the elements using $x \text{ x-ow} x \text{ and } x \text{ col} x \text{ are still possible, but probably not the best option. In the non-numeric case, providing a matrix of ordering ranks for mat would be faster and give more accurate ordering. See the examples section.$

If mat is a matrix of non-repeating random integers (i.e. sample(1:length(x)), replace=FALSE) sample(1:length(x)) |> matrix(ncol=ncol(x))), x %row~% mat will randomly shuffle the elements of every row, where the shuffling order of every row is independent of the other rows. Similarly, x %col~% mat will randomly shuffle the elements of every column, where the shuffling order of every column is independent of the other columns.

These operators internally only use vectorized operations (no loops or apply-like functions), and are faster than re-ordering matrices using loops or apply-like functions.

Value

A modified matrix.

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Examples

```
# numeric matrix ====
x <- matrix(sample(1:25), nrow=5)</pre>
print(x)
x %row~% x # sort elements of every row
x %row~% -x # reverse-sort elements of every row
x %col~% x # sort elements of every column
x \ \%col^{\sim}\% - x \ \# \ reverse-sort \ elements \ of \ every \ column
x <- matrix(sample(1:25), nrow=5)</pre>
print(x)
mat <- sample(1:length(x)) |> matrix(ncol=ncol(x)) # matrix of non-repeating random integers
x %row~% mat # randomly shuffle every row independently
x %col~% mat # randomize shuffle every column independently
# character matrix ====
x <- matrix(sample(letters, 25), nrow=5)</pre>
print(x)
mat <- stringi::stri_rank(as.vector(x)) |> matrix(ncol=ncol(x))
x %row~% mat # sort elements of every row
x %row~% -mat # reverse-sort elements of every row
x %col~% mat # sort elements of every column
x %col~% -mat # reverse-sort elements of every column
x <- matrix(sample(letters, 25), nrow=5)</pre>
mat <- sample(1:length(x)) \mid > matrix(ncol=ncol(x)) \# matrix of non-repeating random integers
x %row~% mat # randomly shuffle every row independently
x %col~% mat # randomize shuffle every column independently
```

pkgs_get_deps

Miscellaneous package functions

Description

The pkgs %installed in% lib.loc operator checks if one or more package(s) pkgs exist(s) in library location lib.loc.

Now you no longer have to attach a package with require() simply to check if it exists.

Moreover, this operator makes it syntactically explicit in your code where you are looking for your R package(s).

The pkgs_get_deps() function gets the dependencies of a package from the Description file. It works on non-CRAN packages also.

Usage

```
pkgs_get_deps(
```

pkgs_get_deps

```
package,
lib.loc = .libPaths(),
deps_type = c("Depends", "Imports", "LinkingTo"),
base = FALSE,
recom = FALSE
)
pkgs %installed in% lib.loc
```

Arguments

package a single string giving the package name lib.loc character vector specifying library search path (the location of R library trees to search through). This is usually .libPaths(). See also loadNamespace. a character vector, giving the dependency types to be used. deps_type Defaults to c("Depends", "Imports", "LinkingTo"). base logical, indicating whether base/core R packages should be included (TRUE), or not included (FALSE; the default). recom logical, indicating whether the pre-installed "recommended" R packages should be included (TRUE), or not included (FALSE; the default). Note that only the recommended R packages actually installed in your system are taken into consideration.

Value

pkgs

For pkgs %installed in% lib.loc: Returns a logical vector, where TRUE indicates a package is installed, and FALSE indicates a package is not installed.

a single string, or character vector, with the package name(s).

For pkgs_get_deps(): A character vector of dependencies.

References

https://stackoverflow.com/questions/30223957/elegantly-extract-r-package-dependencies-of-a-package-not-listed-on-cran

```
## Not run:
pkgs <- c(unlist(tools::package_dependencies("devtools")), "devtools")
pkgs %installed in% .libPaths()
## End(Not run)</pre>
```

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source_module	Additional module import management
---------------	-------------------------------------

Description

The alias %@source% list(file=...) operator imports all objects from a source-able script file under an alias.

The source_inops() function exposes the infix operators defined in a source-able script file to the current environment (like the global environment, or the environment within a function).

Note that the alias %@source% list(file=...) operator and the source_inops() function do NOT suppress output (i.e. plots, prints, messages) from the sourced module file.

Usage

```
alias %@source% lst
source_inops(...)
```

Arguments

alias	a variable name (unquoted), giving the (not yet existing) object where the sourced
	objects from the module are to be assigned to.
	Syntactically invalid names are not allowed for the alias name.
lst	a named list, giving the arguments to be passed to the source function.
	For example: alias %@source% list(file="mydir/myscript.R")
	arguments to be passed to the source function, such as the file argument.

Value

For the alias %@source% list(file=...) operator:

The variable named as the alias will be created (if it did not already exist) in the current environment, and will contain all objects from the sourced script.

```
For source_inops():
```

The infix operators from the specified module will be placed in the current environment (like the Global environment, or the environment within a function).

```
## Not run:
alias %@source% list(file="mydir/mymodule.R")
source_inops(file="mydir/mymodule.R")
## End(Not run)
```

18 stri_join_mat

stri_join_mat

Concatenate Character Matrix Row-wise or Column-wise

Description

The stri_join_mat() function (and their aliases stri_c_mat and stri_paste_mat) perform rowwise (margin=1; the default) or column-wise (margin=2) joining of a matrix of strings, thereby transforming a matrix of strings into a vector of strings.

Usage

```
stri_join_mat(mat, margin = 1, sep = "", collapse = NULL)
stri_c_mat(mat, margin = 1, sep = "", collapse = NULL)
stri_paste_mat(mat, margin = 1, sep = "", collapse = NULL)
```

Arguments

mat a matrix of strings

margin the margin over which the strings must be joined.

If margin=1, the elements in each row of matrix mat are joined into a single

string. Thus if the matrix has 10 rows, it returns a vector of 10 strings.

If margin=2, the elements in each column of matrix mat are joined into a single string. Thus if the matrix has 10 columns, it returns a vector of 10 strings.

sep, collapse as in stri_join.

Details

The examples section show the uses of the stri_join_mat() function.

Value

The stri_join_mat() function, and its aliases, return a vector of strings.

stri_locate_ith

```
x <- c("Hello everyone", "Goodbye everyone")</pre>
mat <- stringi::stri_split_boundaries(x, simplify = TRUE, type="word")</pre>
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank</pre>
print(sorted)
stri_c_mat(sorted, margin=1) # <- alias for stri_join_mat</pre>
stri_c_mat(sorted, margin=2)
# re-ordering sentences ====
x <- c("Hello, who are you? Oh, really?! Cool!", "I don't care. But I really don't.")
print(x)
mat <- stringi::stri_split_boundaries(x, simplify = TRUE, type="sentence")</pre>
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank
print(sorted)
stri_paste_mat(sorted, margin=1) # <- another alias for stri_join_mat</pre>
stri_paste_mat(sorted, margin=2)
```

stri_locate_ith

Locate i^th Pattern Occurrence

Description

The stri_locate_ith function locates the i^{th} occurrence of a pattern in each string of some character vector.

Usage

```
stri_locate_ith(str, i, ..., regex, fixed, coll, charclass)
```

Arguments

str

a string or character vector.

i

a number, or a numeric vector of the same length as str. This gives the i^{th} instance to be replaced.

Positive numbers are counting from the left. Negative numbers are counting from the right. I.e.:

 $stri_locate_ith(str, i=1, ...)$ gives the position (range) of the first occurrence of a pattern.

stri_locate_ith(str, i=-1, ...) gives the position (range) of the last occurrence of a pattern.

 $stri_locate_ith(str, i=2, ...)$ gives the position (range) of the second occurrence of a pattern.

 $stri_locate_ith(str, i=-2, ...)$ gives the position (range) of the second-last occurrence of a pattern.

If abs(i) is larger than the number of instances, the first (if i < 0) or last (if i > 0) instance will be given.

20 stri_locate_ith

```
For example: suppose a string has 3 instances of some pattern; then if i >= 4 the third instance will be located, and if i <= -3 the first instance will be located.

... more arguments to be supplied to stri_locate and stri_count.

regex, fixed, coll, charclass
a character vector of search patterns, as in stri_locate.
```

Value

The stri_locate_ith() function returns an integer matrix with two columns, giving the start and end positions of the i^{th} matches, and two NAs if no matches are found.

```
# simple pattern ====
x <- rep(paste0(1:10, collapse=""), 10)</pre>
print(x)
out <- stri_locate_ith(x, 1:10, regex = as.character(1:10))</pre>
cbind(1:10, out)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
p <- rep("a|e|i|o|u",2)</pre>
out <- stri_locate_ith(x, c(-1, 1), regex=p)</pre>
print(out)
substr(x, out[,1], out[,2])
# ignore case pattern ====
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- rep("A|E|I|0|U", 2)</pre>
out <- stri_locate_ith(x, c(1, -1), regex=p, case_insensitive=TRUE)</pre>
substr(x, out[,1], out[,2])
# multi-character pattern ====
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
# multi-character pattern:
p \leftarrow rep("AB", 2)
out <- stri_locate_ith(x, c(1, -1), regex=p, case_insensitive=TRUE)
print(out)
substr(x, out[,1], out[,2])
```

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```
# Replacement transformation using stringi ====
x <- c("hello world", "goodbye world")
loc <- stri_locate_ith(x, c(1, -1), regex="a|e|i|o|u")
extr <- stringi::stri_sub(x, from=loc)
repl <- chartr(extr, old = "a-zA-Z", new = "A-Za-z")
stringi::stri_sub_replace(x, loc, replacement=repl)</pre>
```

str_arithmetic

String arithmetic

Description

String arithmetic operators.

The x %s+%y operator is equivalent to stringi::stri_c(x,y).

The x %s-% p operator removes character/pattern defined in p from x.

The x %s*% n operator repeats every element of x for n times, and glues them together.

The x %s/% p operator counts how often regular expression or character pattern p occurs in each element of x.

Usage

x %s+% y

x %s-% p

x %s*% n

x %s/% p

Arguments

Χ	a string or character vector.

y a string, or a character vector of the same length as x.

p the result from s_pattern, or else a character vector of the same length as x with

regular expressions.

n a number, or a numeric vector of the same length as x.

22 str_subset_ops

Details

These operators and functions serve as a way to provide straight-forward string arithmetic, missing from base R.

Value

The %s+%, %s-%, and %s*% operators return a character vector of the same length as x. The %s/% returns a integer vector of the same length as x.

Examples

```
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
y <- c("a", "b")
p <- rep("a|e|i|o|u", 2) \ \# \ same \ as \ p <- \ s\_pattern(regex=rep("a|e|i|o|u", 2))
n < -c(3, 2)
x %s+% y # =paste0(x,y)
x %s-% p # remove all vowels from x
x %s*% n
x %s/% p # count how often vowels appear in each string of vector x.
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
y <- "a"
# pattern that ignores case:
p <- s_pattern(regex=rep("A|E|I|0|U", 2), ignore.case=TRUE)</pre>
n < -c(2, 3)
x %s+% y # =paste0(x,y)
x %s-% p # remove all vowels from x
x %s*% n
x %s/% p # count how often vowels appears in each string of vector x.
```

str_subset_ops

String subsetting operators

Description

String subsetting operators.

The x %ss% s operator allows indexing a single string as-if it is an iterable object.

The x %sget% ss operator gives a certain number of the first and last characters of x.

The x %strim% ss operator removes a certain number of the first and last characters of x.

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Usage

```
x %ss% s
x %sget% ss
x %strim% ss
```

Arguments

X	a string or character vector.
S	a numeric vector giving the subset indices.
SS	a vector of length 2, or a matrix with 2 columns with nrow(ss)==length(x). The object ss should consist entirely of non-negative integers (thus 0, 1, 2, etc. are valid, but -1, -2, -3 etc are not valid). The first element/column of ss gives the number of characters counting from the left side to be extracted/removed from x. The second element/column of ss gives the number of characters counting

from the right side to be extracted/removed from x.

Details

These operators serve as a way to provide straight-forward string sub-setting.

Value

The %ss% operator always returns a vector or matrix, where each element is a single character.

```
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
ss <- c(2,3)
x %sget% ss
x <- c(paste0(letters[1:13], \ collapse=""), \ paste0(letters[14:26], \ collapse=""))
print(x)
ss <- c(1,0)
x %sget% ss
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
ss <- c(2,3)
x %strim% ss
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
ss <- c(1,0)
x %strim% ss
```

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substr_repl

Substr - functions

Description

Fully vectorized sub-string functions.

These functions extract, replace, add-in, transform, or re-arrange, the i^{th} pattern occurrence or position range.

The $substr_repl(x, rp, ...)$ function replaces a position (range) with string rp.

The substr_chartr(x, old, new, ...) function transforms the sub-string at a position (range) using chartr(old, new).

The $substr_addin(x, addition, side, ...)$ function adds the additional string addition at the side (specified by argument side) of a position.

The $substr_extract(x, type, ...)$ function extracts the string at, before, or after some position.

The substr_arrange(x, arr, ...) function sorts (alphabetically or reverse-alphabetically) or reverses the sub-string at a position (range).

Usage

```
substr_repl(x, rp, ..., loc = NULL, start = NULL, end = NULL, fish = FALSE)
substr_chartr(
    x,
    old = "a-zA-Z",
    new = "A-Za-z",
    ...,
    loc = NULL,
    start = NULL,
    end = NULL,
    fish = FALSE
)
substr_addin(
```

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```
х,
  addition,
  side = "after",
  loc = NULL,
  at = NULL,
  fish = FALSE
substr_extract(
  type = "at",
  ...,
  loc = NULL,
  start = NULL,
  end = NULL,
  fish = FALSE
substr_arrange(
  arr = "incr",
  ...,
  loc = NULL,
  start = NULL,
  end = NULL,
  opts_collator = NULL,
  fish = FALSE
)
```

Arguments

X	a string or character vector.
rp	a string, or a character vector of the same length as \boldsymbol{x} , giving the replacing strings.
	only applicable if fish=TRUE; other arguments to be passed to the stringfish functions.
loc	The result from the stri_locate_ith function. NOTE: you cannot fill in both loc and start, end, or both loc and at. Choose one or the other.
start, end	integers, or integer vectors of the same length as \boldsymbol{x} , giving the start and end position of the range to be modified.
fish	although tinyoperators has no dependencies other than stringi, it does allow the internal functions to use the multi-threadable stringfish functions. To do so, set fish=TRUE; this requires stringfish to be installed.
old, new	see chartr. Defaults to old="a-zA-Z", new="A-Za-z", which means upper case characters will be transformed to lower case characters, and vice-versa.
addition	a string, or a character vector of the same length as \mathbf{x} , giving the string(s) to add-in.
side	which side of the position to add in the string. Either "before" or "after".

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at

an integer, or integer vector of the same length as x, giving the position after or before which the string is to be added.

type

a single string, giving the part of the string to extract. 3 options available:

- type = "at": extracts the string part at the position range;
- type = "before": extracts the string part before the position range;
- type = "after": extracts the string part after the position range.

arr

a single string, giving how the sub-string should be arranged. 3 options available:

```
• arr = "incr": sort the sub-string alphabetically.
```

- arr = "decr": sort the sub-string reverse alphabetically.
- arr = "rev": reverse the sub-string.
- arr = "rand": randomly shuffles the sub-string.

opts_collator as in stri_rank. Only used when arr = "incr" or arr = "decr".

Details

These functions serve as a way to provide straight-forward sub-string modification and/or extraction

All substr_ functions internally only use fully vectorized R functions (no loops or apply-like functions).

Value

A modified character vector. If no match is found in a certain string of character vector x, the unmodified string is returned. The exception is for the substr_extract() function: in this function, non-matches return NA.

```
# numerical substr ====

x <- rep("12345678910", 2)
start=c(1, 2); end=c(3,4)
substr_extract(x, start=start, end=end)
substr_extract(x, type="before", start=start, end=end)
substr_extract(x, type="after", start=start, end=end)
substr_repl(x, c("??", "!!"), start=start, end=end)
substr_chartr(x, start=start, end=end)
substr_addin(x, c(" ", "~"), "after", at=end)
substr_addin(x, c(" ", "~"), "before", at=start)
substr_arrange(x, start=start, end=end)
substr_arrange(x, "decr", start=start, end=end)
substr_arrange(x, "rev", start=start, end=end)
substr_arrange(x, "rand", start=start, end=end)
substr_arrange(x, "rand", start=start, end=end)
substr_arrange(x, "rand", start=start, end=end)</pre>
```

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```
substr_extract(x, start=start, end=end)
substr_extract(x, type="before", start=start, end=end)
substr_extract(x, type="after", start=start, end=end)
substr_repl(x, "??", start=start, end=end)
substr_chartr(x, start=start, end=end)
substr_addin(x, " ", "after", at=end)
substr_addin(x, " ", "before", at=start)
start=5: end=6
substr_extract(x, start=start, end=end)
substr_extract(x, type="before", start=start, end=end)
substr_extract(x, type="after", start=start, end=end)
substr_repl(x, "??", start=start, end=end)
substr_chartr(x, start=start, end=end)
substr_addin(x, " ", "after", at=end)
substr_addin(x, " ", "before", at=start)
# simple pattern ====
x <- c("goodGOODGoodgOOd", "goodGOODGoodgOOd", paste0(letters[1:13], collapse=""))</pre>
print(x)
loc <- stri_locate_ith(</pre>
  # locate second-last occurrence of "good" of each string in x:
  x, -2, regex="good", case_insensitive=TRUE
substr_extract(x, loc=loc) # extract second-last "good"
substr_repl(x, c("??", "!!", " "), loc=loc) # replace second-last "good"
substr_chartr(x, loc=loc) # switch upper/lower case of second-last "good"
substr\_addin(x,\ c("\ ",\ "^{-}",\ "\ "),\ "after",\ loc=loc)\ \#\ add\ white\ space\ after\ second-last\ "good"
substr_addin(x, c(" ", "~", " "), "before", loc=loc) # add white space before second-last "good"
substr_arrange(x, loc=loc) # sort second-last "good"
substr_arrange(x, "decr", loc=loc) # reverse-sort second-last "good"
substr_arrange(x, "rev", loc=loc) # reverse second-last "good"
substr_arrange(x, "rand", loc=loc) # randomly shuffles "good"
```

s_pattern

Pattern attribute assignment

Description

The %s-% and %s/% operators, their in-place equivalents, as well as the %sgrep% operator, all perform pattern matching for some purpose. By default the pattern matching is interpreted as case-sensitive regex patterns from stringi.

The s_pattern function allows the user to specify exactly how the pattern should be interpreted. To use more refined pattern definition, simply replace the right-hand-side expression p in the relevant operators with a call from the s_pattern() function.

The s_pattern() function uses the exact same argument convention as stringi. For example:

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```
s_pattern(regex=p, case_insensitive=FALSE, ...)
s_pattern(fixed=p, ...)
s_pattern(coll=p, ...)
s_pattern(charclass=p, ...)
```

All arguments in s_pattern() are simply passed to the appropriate functions in stringi. For example:

x %s/% p counts how often regular expression p occurs in x,

whereas x %s/% s_pattern(fixed=p, case_insensitive=TRUE) will do the same, except it uses fixed (i.e. literal) expression, and it does not distinguish between upper case and lower case characters.

For consistency with base R, one can also fill in <code>ignore.case=TRUE</code> or <code>ignore_case=TRUE</code> instead of <code>case_insensitive=TRUE</code>, and <code>s_pattern</code> will still understand that.

Usage

```
s_pattern(...)
```

Arguments

pass stringi arguments here. I.e. regex=p, coll=p, charclass=p, case_insensitive=FALSE, etc. See the documentation in the stringi R package.

Details

The s_pattern() function only works in combination with the functions and operators in this package. It does not affect functions from base R or functions from other packages.

Value

The s_pattern(...) call returns a list with arguments that will be passed to the appropriate functions in stringi.

```
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse="")) print(x) p <- rep("a|e|i|o|u", 2) # same as p <- s_pattern(regex=rep("a|e|i|o|u", 2)) x %s/% p # count how often vowels appear in each string of vector x.   
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse="")) print(x) p <- s_pattern(regex=rep("A|E|I|O|U", 2), case_insensitive=TRUE) x %s/% p # count how often vowels appear in each string of vector x.
```

tinyoperators_help 29

Description

Welcome to the tinyoperators help page!

The tinyoperators R-package adds some infix operators, and a few functions. It primarily focuses on 4 things:

- (1) Float truth testing.
- (2) Reducing repetitive code.
- (3) Extending the string manipulation capabilities of the stringi R package.
- (4) A new package and module import system, that combines the benefits of aliasing a package with the benefits of attaching a package.

The tinyoperators R-package has only one dependency, namely stringi, though it does allows multi-threading of some of the string-related functions (when appropriate) via the suggested stringfish R-package.

The tinyoperators R package adds the following functionality:

- Infix logical operators for exclusive-or, not-and, not-in, number-type, and string-type.
- Safer (in)equality operators for floating numbers.
- Infix operators for string arithmetic.
- Infix operators for string sub-setting.
- Several operators for the "Don't Repeat Yourself" coding principle (DRY). This includes the generalized in-place (mathematical) modification operator, infix operators for In-place modifying string arithmetic, and infix operators for In-place modifying string sub-setting.
- Infix operators for row- and column-wise re-ordering of matrices.
- The tinyoperators package adds additional stringi functions, namely stri_locate_ith and stri_join_mat (and aliases). These functions use the same naming and argument convention as the rest of the stringi functions, thus keeping your code consistent.
- The fully vectorized sub-string functions, that extract, replace, add-in, transform, or re-arrange, the i^{th} pattern occurrence or location.
- The s_pattern helper function for string operators.

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- New package import functions, and new module sourcing functions.
- Most stringi pattern expressions options are available for the string-pattern-related functions, when appropriate.
- This R package has only one dependency: stringi. No other dependencies, as to avoid "dependency hell".
- Although this package has no other dependencies, it allows multi-threading of the sub-string functions through the stringfish R package.

Please also have a look at the Read-Me file on the GitHub main page before using this package: https://github.com/tony-aw/tinyoperators

Usage

```
tinyoperators_help()
```

transform_if

The transform_if function and the subset_if operators

Description

Consider the following code:

```
ifelse(cond(x), f(x), g(x))
```

Here a conditional subset of the object x is transformed, where the condition is using a function referring to x itself. Consequently, reference to x is written four times!

The tinyoperators package therefore adds the transform_if() function which will tiny this up.

The tinyoperators package also adds 2 "subset_if" operators:

The $x \leq if \leq cond$ operator selects elements from vector/matrix/array x, for which the result of cond(x) returns TRUE.

The $x \leq [if]$ % cond operator selects elements from vector/matrix/array x, for which the result of cond(x) returns FALSE.

The tinyoperators package also adds the x %unreal =% repl operator: x %unreal = % repl is the same as x[is.na(x)|is.nan(x)|is.infinite(x)] <- repl

Usage

```
transform_if(x, cond, trans_T = function(x) x, trans_F = function(x) x)
x %[if]% cond
x %[!if]% cond
x %unreal =% repl
```

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Arguments

a vector, matrix, or array.

cond

a function that returns a binary logic (TRUE, FALSE) vector of the same length/dimensions as x (for example: is.na).

trans_T

the transformation function to use when cond(x)==TRUE.
For example: log.
If this is not specified, trans_T defaults to function(x)x.

trans_F

the transformation function to use when cond(x)==FALSE.
For example: log.
If this is not specified, trans_F defaults to function(x)x.

repl

the replacement value.

Details

The transform_if(x, cond, trans) function does not rely on any explicit or implicit loops, nor any third-party functions.

Value

```
For transform_if(): Similar to ifelse. However, unlike ifelse(), the transformations are evaluated as trans_T(x[cond(x)]) and trans_F(x[!cond(x)]), ensuring no unnecessary warnings or errors occur.
```

The subset_if - operators all return a vector with the selected elements.

The x %unreal =% repl operator does not return any value: It is an in-place modifiers, and thus modifies x directly. The object x is modified such that all NA, NaN and Inf elements are replaced with repl.

```
object_with_very_long_name <- matrix(-10:9, ncol=2)
print(object_with_very_long_name)
object_with_very_long_name |> transform_if(\(x)x>0, log)
object_with_very_long_name |> transform_if(\(x)x>0, log, \(x)x^2)
object_with_very_long_name %[if]% \(x)x %in% 1:10
object_with_very_long_name %[!if]% \(x)x %in% 1:10

x <- c(1:9, NA, NaN, Inf)
print(x)
x %unreal =% 0 # same as x[is.na(x)|is.nan(x)|is.infinite(x)] <- 0
print(x)</pre>
```

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