Package 'tinyoperations'

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```
Title Functions and infix operators to help in your programming etiquette
Version 0.0.0.9
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Description The 'tinyoperations' R-
      package adds some functions and infix operators to help in your programming etiquette.
      It primarily focuses on 4 things.
      1) Safer decimal (in)equality testing, safer atomic conversions, and other functions for safer cod-
      ing.
      2) A new package import system,
      that combines the benefits of aliasing a package with the benefits of attaching a package.
      3) Extending the string manipulation capabilities of the `stringi` R package.
      4) Reducing repetitive code.
      The 'tinyoperations' R-package has only one dependency, namely 'stringi'.
      Most functions in this R-package are fully vectorized and optimized,
      and have been well documented.
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atomic_conversions

Safer atomic type casting

Description

Index

Atomic type casting in R is generally performed using the functions as.logical, as.integer, as.double, as.character.

Converting an object between atomic types using these functions strips the object of its attributes, including attributes such as names and dimensions.

The functions provided here by the tinyoperations package do not strip away attributes - except the "class" attribute.

The functions are as follows:

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- as_bool(): converts object to class logical (TRUE, FALSE, NA).
- as_int(): converts object to class integer.
- as_dbl(): converts object to class double (AKA decimal numbers).
- as_chr(): converts object to class character.

Moreover, the function is_wholenumber() is added, to safely test for whole numbers.

Usage

```
as_bool(x, ...)
as_int(x, ...)
as_dbl(x, ...)
as_chr(x, ...)
is_wholenumber(x, tol = .Machine$double.eps^0.5)
```

Arguments

```
    vector, matrix, array (or similar object where all elements share the same atomic class), to be converted to some other atomic class.
    further arguments passed to or from other methods.
    tol the tolerance.
```

Value

The converted object.

See Also

```
tinyoperations_safer()
```

```
x <- c(rep(0, 2), seq(0, 2.5, by=0.5)) |> matrix(ncol=2)
colnames(x) <- c("one", "two")
attr(x, "test") <- "test"
print(x)

# notice that in all following, attributes are conserved:
as_bool(x)
as_int(x)
as_dbl(x)
as_chr(x)</pre>
```

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decimal_truth

Safer decimal number (in)equality testing operators

Description

The %d==%, %d!=% %d<%, %d>%, %d<=%, %d>=% (in)equality operator perform decimal (class "double") number truth testing.

They are virtually equivalent to the regular (in)equality operators,

```
==, !=, <, >, <=, >=,
```

except for two aspects.

First: The decimal number (in)equality operators assume that if the absolute difference between any two numbers x and y is smaller than the Machine tolerance, sqrt(.Machine\$double.eps), then x and y should be consider to be equal.

Second: The decimal number (in)equality operators do NOT allow vector recycling. When comparing any two numbers x and y, they must be of equal length, or one of them must be a scalar (i.e. length of 1). Otherwise, an error is returned.

Thus these operators provide safer decimal number (in)equality tests.

For example: 0.1*7 == 0.7 returns FALSE, even though they are equal, due to the way decimal numbers are stored in programming languages (like R).

But 0.1*7 %d==% 0.7 returns TRUE.

Another example: The code 1:10 == c(4, 6) is silently transformed to 1:10 == rep(c(4, 6), each = 5), and thus gives c(rep(FALSE, 5), TRUE, rep(FALSE, 4)).

This may not be what the user wants. But 1:10 %d==% c(4, 6) prevents this by returning an error instead.

There are also the $x %d{}\%$ bnd and $x %d!{}\%$ 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 %d{}% bnd operator checks if x is within the closed interval with bounds defined by bnd.

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

Usage

```
x %d==% y
```

x %d!=% y

x %d<% y

x %d>% y

x %d<=% y

x %d>=% y

x %d{}% bnd

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```
x %d!{}% bnd
```

Arguments

x, y numeric vectors, matrices, or arrays.

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

Value

A logical vector with the same dimensions as x, indicating the result of the element by element comparison.

See Also

```
tinyoperations_safer()
```

```
x <- c(0.3, 0.6, 0.7)
y \leftarrow 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 \# gives TRUE, should be FALSE
x %d==% y # here it's done correctly
x %d!=% y # correct
x %d<% y # correct
x %d>% y # correct
x %d<=% y # correct
x %d>=% y # correct
x < -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 %d{}% bnd
x %d!{}% bnd
# These operators still work for non-decimal number numerics also:
x <- 1:5
y <- 1:5
x %d==% y
x %d!=% y
x %d<% y
x %d>% y
x %d<=% y
x %d>=% y
x <- 1:5
y <- x+1
x %d==% y
x %d!=% y
```

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```
x %d<% y
x %d>% y
x %d<=% y
x %d>=% y
x <- 1:5
y <- x-1
x %d==% y
x %d!=% y
x %d<% y
x %d<=% y
x %d<=% y
```

import_as

Load main package + its re-exports + its dependencies + its enhances + its extensions under one alias

Description

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

Usage

```
import_as(
  alias,
  main_package,
  re_exports = TRUE,
  dependencies = NULL,
  enhances = NULL,
  extensions = NULL,
  lib.loc = .libPaths(),
  loadorder = c("dependencies", "main_package", "enhances", "extensions")
)
```

Arguments

alias

a syntactically valid non-hidden name giving the alias object where the package(s) are to be loaded into.

This name can be given either as a single string (i.e. "alias."), or as a one-sided formula with a single term (i.e. ~alias.).

NOTE: To keep aliases easily distinguishable from other objects that can also be subset with the \$ operator, I recommend ending (not starting!) all alias names with a dot (.) or underscore (_).

main_package

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

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re_exports

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

This argument determines what the import_as function will do with the reexports of the main_package:

- If TRUE the re-exports from the main_package are added to the alias, even if dependencies = NULL. This is the default, as it is analogous to the behaviour of base R's :: operator.
- If FALSE, these re-exports are not added, and the user must specify the appropriate packages in argument dependencies.

dependencies

an optional character vector, giving the names of the dependencies of the main_package

to be loaded also under the alias.

Defaults to NULL, which means no dependencies are loaded.

See pkg_get_deps to quickly get dependencies from a package.

enhances

an optional 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.

extensions

an optional character vector, giving the names of the extensions of the main_package

to be loaded also under the alias.

Defaults to NULL, which means no extensions are loaded.

lib.loc

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

The lib.loc argument would usually be .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 re-exports (if re_exports=TRUE).
- 3. The enhances, in the order specified by the enhances argument.
- 4. The extensions, in the order specified by the extensions argument.

Details

On the dependencies, enhances and extensions arguments

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

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• enhances: Enhances are defined as packages appearing in the "Enhances" field of the Description file of the main_package.

• extensions: "Extensions" here are defined as reverse-depends or reverse-imports. It does not matter if these are CRAN or non-CRAN packages. However, the intended meaning of an extension is not merely being a reverse dependency, but a package that actually extends the functionality of the main_package.

As implied in the description of the loadorder argument, the order of the character vectors given in the dependencies, enhances, and extensions arguments matter:

If multiple packages share objects with the same name, the objects of the package named last will overwrite those of the earlier named packages.

Additional details

The import_as() function does not support loading base/core R under an alias.

Value

A locked environment object, similar to the output of loadNamespace, with the name as specified in the alias argument, will be created in the current environment (like the global environment, or the environment within a function).

```
To use, for example, function "some_function()" from alias "alias.", use: alias.$some_function()
To see the special attributes of this alias object, use attr.import.
To "unload" the package alias object, simply remove it (i.e. rm(list="alias.")).
```

See Also

```
tinyoperations_import()
```

```
## Not run:
import_as( # this creates the 'tdt.' object
   "tdt.", "tidytable", dependencies = "data.table"
)
# same as:
import_as(
   ~ tdt., "tidytable", dependencies = "data.table"
)
# using a function from the alias:
tdt.$mutate
## End(Not run)
```

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import_data

Directly return a data-set from a package

Description

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

Usage

```
import_data(package, dataname, lib.loc = .libPaths())
```

Arguments

package a single string, giving the name of the R-package.

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

lib.loc character vector specifying library search path (the location of R library trees to

search through).

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

Value

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

See Also

```
tinyoperations_import()
```

Examples

```
d <- import_data("gamair", "chicago")
head(d)</pre>
```

import_inops

(un)expose infix operators in the current environment

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Description

import_inops(expose=...) exposes infix operators specified in a package or an alias object to the current environment (like the global environment or the environment within a function).

import_inops(unexpose=...) unexposes (i.e. removes) the infix operators specified in a packages or an alias object from the current environment (like the global environment or the environment within a function).

Note that in this case only infix operators exposed by the tinyoperations import system will be removed from the current environment;

"regular" infix operators (i.e. user-specified) will not be touched.

Usage

```
import_inops(expose = NULL, unexpose = NULL, lib.loc = .libPaths(), ...)
```

Arguments

expose

either one of the following:

- a package alias as produced by the import_as function.
- a string giving the package name from which to load infix operators, and place them in the current environment.

unexpose

either one of the following:

- a package alias as produced by the import_as function.
- a string giving the package name.

lib.loc

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

Only used when supplying a character vector of package names to expose / unexpose, and ignored when supplying an alias object to expose / unexpose (the library is already defined inside the alias object).

The lib.loc argument would usually be .libPaths().

See also loadNamespace.

... additional arguments, only relevant if the expose argument is used.

See import_inops.control.

Details

The import_inops() function does not support overloading base/core R operators.

When using import_inops() to remove infix operators from the current environment, it will use the attributes of those operators to determine if the infix operator came from the tinyoperations import system, or if they were user-defined.

Value

If using argument expose:

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

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If using argument unexpose:

The infix operators specified in the given package or alias, exposed by import_inops(), will be deleted.

If such infix operators could not be found, this function returns NULL.

See Also

```
tinyoperations_import(), import_inops.control()
```

Examples

```
## Not run:
import_as(dt., "data.table")
import_inops(expose = dt.) # expose infix operators from alias
import_inops(unexpose = dt.) # unexposed infix operators from current environment
import_inops(expose = "data.table") # expose infix operators from package
import_inops(unexpose = "data.table") # remove the exposed infix operators from environment
# additional arguments (only used when exposing, not unexposing):
import_as(dt., "data.table")
import_inops(expose = dt., include.only = ":=")
import_inops(unexpose = dt.)
import_inops(expose = "data.table", exclude = ":=")
import_inops(unexpose = "data.table")
import_inops(expose = dt., overwrite = FALSE)
import_inops(unexpose = dt.)
import_inops(expose = "data.table", overwrite = FALSE)
import_inops(unexpose = "data.table")
## End(Not run)
```

```
import_inops.control import_inops.control
```

Description

Additional arguments to control exposing infix operators in the import_inops function.

Usage

```
import_inops.control(
  exclude = NULL,
  include.only = NULL,
  overwrite = TRUE,
  inherits = FALSE
)
```

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Arguments

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.

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.

overwrite

logical, indicating if it is allowed to overwrite existing infix operators.

- If TRUE (default), a warning is given when operators existing in the current environment are being overwritten, but the function continuous nonetheless.
- If FALSE, an error is produced when the to be exposed operators already exist in the current environment, and the function is halted.

inherits

logical; when overwrite=FALSE, should enclosed environments, especially package namespaces, also be taken into account?

Defaults to FALSE. See also exists.

Details

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

Value

This function is used internally in the import_inops function.

See Also

```
import_inops(), tinyoperations_import()
```

```
## Not run:
# additional arguments (only used when exposing, not unexposing):
import_as(dt., "data.table")
import_inops(expose = dt., include.only = ":=")
import_inops(unexpose = dt.)
import_inops(expose = "data.table", exclude = ":=")
import_inops(unexpose = "data.table")
import_inops(expose = dt., overwrite = FALSE)
import_inops(unexpose = dt.)
import_inops(expose = "data.table", overwrite = FALSE)
import_inops(unexpose = "data.table", overwrite = FALSE)
import_inops(unexpose = "data.table")
```

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

import_LL

Miscellaneous import_ - functions

Description

The import_LL() function places specific functions from a package in the current environment, and also locks (see lockBinding) the specified functions to prevent modification.

The primary use-case for this function is for loading functions inside a local environment, like the environment within a function.

The import_int() function directly returns an internal function from a package.

It is similar to the ::: operator, but with 2 key differences:

- 1. It allows the user to explicitly set a library location through the lib.loc argument.
- 2. It only searches internal functions, not exported ones. This makes it clearer in your code that you're using an internal function, instead of making it ambiguous.

Usage

```
import_LL(package, selection, lib.loc = .libPaths())
import_int(form, lib.loc = .libPaths())
```

Arguments

package	a single string, giving the name of the package to take functions from.
selection	a character vector of function names (both regular functions and infix operators). Internal functions or re-exported functions are not supported.
lib.loc	character vector specifying library search path (the location of R library trees to search through). The lib.loc argument would usually be .libPaths(). See also loadNamespace.
form	a two-sided formula, with one term on each side.

The term on the left hand side should give a single package name. The term on the right hand side should give a single internal function.

Example: package_name ~ function_name

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Details

Regarding the Locks in import_LL()

The import_as function returns a locked environment, just like loadNamespace, thus protecting the functions from accidental modification or re-assignment.

The import_inops function returns infix operators, and though these are not locked, one needs to surround infix operators by back ticks to re-assign or modify them, which is unlikely to happen on accident.

The import_LL() function, however, returns "loose" functions. And these functions (unless they are infix operators) do not have the protection due to a locked environment or due to the syntax. Therefore, to ensure safety from (accidental) modification or re-assignment, the import_LL()

function locks these functions (see lockBinding). For consistency, infix operators exposed by import_LL() are also locked.

Other details

The import_LL() and import_int() functions do not support importing from base/core R.

Value

```
For import_LL():
```

The specified functions will be placed in the current environment (like the global environment, or the environment within a function), and locked.

```
To "unload" or overwrite the functions, simply remove them; i.e.: rm(list=c("some_function1", "some_function2")).
```

```
For import_int():
```

The function itself is returned directly.

So one can assign the function directly to some variable, like so:

```
myfun <- import_int(...)</pre>
```

or use it directly without re-assignment like so:

```
import_int(...)(...)
```

See Also

```
tinyoperations_import()
```

```
## Not run:
# Using import_LL ====
import_LL(
    "tidytable", "across"
)
across # <- this function cannot be modified, only used or removed, because it's locked
# Using internal function ====
# Through re-assignment:
fun <- import_int(tinyoperations ~ .internal_paste, .libPaths())
fun("hello", "world")
# Or using directly:
import_int(
    tinyoperations ~ .internal_paste, .libPaths()
)("hello", "world")</pre>
```

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

inplace

Generalized in-place modifier

Description

Generalized in-place modifier.

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

For example this:

Usage

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

Arguments

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

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

like') vector, matrix, or array.

f a (possibly anonymous) function to be applied in-place on x. The function must

take one argument only.

Value

This operator does not return any value:

It is an in-place modifier, and thus modifies the object directly.

See Also

```
tinyoperations_dry()
```

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

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lock

Lock T. lock F. or create locked constants

Description

One can re-assign the values T and F. One can even run things like $T \leftarrow FALSE$ and $F \leftarrow TRUE$! The lock_TF() function locks the T and F values and sets them to TRUE and FALSE, respectively, to prevent the user from re-assigning them.

Removing the created T and F objects allows re-assignment again.

The X %<-c% A operator creates a constant X and assigns A to it.

Constants cannot be changed, only accessed or removed. So if you have a piece of code that requires some unchangeable constant, use this operator to create said constant.

Removing constant X also removes its binding lock. Thus to change a constant, simply remove it and re-create it.

Usage

```
lock_TF()
X %<-c% A</pre>
```

Arguments

X a syntactically valid unquoted name of the object to be created.

A any kind of object to be assigned to X.

Value

For lock_TF():

Two constants, namely T and F, set to TRUE and FALSE respectively, are created in the current environment, and locked. Removing the created T and F objects allows re-assignment again.

```
For X %<-c% A:
```

The object X containing A is created in the current environment, and this object cannot be changed. It can only be accessed or removed.

See Also

```
tinyoperations_safer()
```

```
lock_TF() 
 X %<-c% data.frame(x=3, y=2) # this data.frame cannot be changed. Only accessed or removed. 
 X[1, drop=FALSE]
```

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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.

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.

Usage

- x %xor% y
- x %n&% y
- x %out% y
- x %?=% y
- n %=numtype% numtype
- s %=strtype% strtype

Arguments

x, y see Logic.

n a numeric vector.

numtype a single string giving the numeric type 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)).

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- "B": binary numbers (exactly 0 or exactly 1);
- "prop": proportions numbers between 0 and 1 (exactly 0 or 1 is also allowed);
- "I": Integers;
- "odd": odd integers;
- "even": even integers;
- "R": Real numbers;
- "unreal": infinity, NA, or NaN;

s a character vector.

strtype

a single string giving the string type 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.

Value

A logical vector.

```
x <- c(TRUE, FALSE, TRUE, FALSE, NA, NaN, Inf, -Inf, TRUE, FALSE)
y <- c(FALSE, TRUE, TRUE, FALSE, rep(NA, 6))
outcome <- data.frame(
    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 <- c(0:5, 0:-5, 0.1, -0.1, 0, 1, Inf, -Inf, NA, NaN)
1e-20 %=numtype% "~0"</pre>
```

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```
n[n %=numtype% "B"]
n[n %=numtype% "prop"]
n[n %=numtype% "odd"]
n[n %=numtype% "even"]
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"]</pre>
```

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, each row ordered independently from the other rows, of matrix x, according to the ordering ranks given in matrix mat.

The x %col~% mat operator re-orders the elements of every column, each column ordered independently from the other columns, 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 element 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 %row% x and x %rol% x is still possible, but probably not the best option. In the non-numeric case, providing a matrix of ordering ranks

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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.

mat <- sample(1:length(x)) |> matrix(ncol=ncol(x)))

then the code

x %row~% mat

will randomly shuffle the elements of every row, where the shuffling order in each row is independent from the shuffling order in the other rows.

Similarly,

x %col~% mat
```

will randomly shuffle the elements of every column, where the shuffling order in each column is independent from the shuffling order in the other columns.

Re-ordering/sorting every row/column of a matrix with these operators is generally faster than doing so through loops or apply-like functions.

Value

A modified matrix.

See Also

```
tinyoperations_misc()
```

```
# numeric matrix ====
x <- matrix(sample(1:25), nrow=5)</pre>
print(x)
x %row~% x # sort elements of every row independently
x %row~% -x # reverse-sort elements of every row independently
x %col~% x # sort elements of every column independently
x %col~% -x # reverse-sort elements of every column independently
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>
mat <- stringi::stri_rank(as.vector(x)) |> matrix(ncol=ncol(x))
x %row~% mat # sort elements of every row independently
x %row~% -mat # reverse-sort elements of every row independently
x col^{\infty} mat # sort elements of every column independently
x %col~% -mat # reverse-sort elements of every column independently
x <- matrix(sample(letters, 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
```

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```
x %col~% mat # randomize shuffle every column independently
```

pkgs

Miscellaneous package related functions

Description

The pkgs %installed in% lib.loc operator checks if one or more package(s) pkgs exist(s) in library location lib.loc, without loading the package(s).

The syntax of this operator forces the user to make it syntactically explicit where to look for installed R package(s).

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

The pkg_lsf(package, ...) function gets a list of exported functions/operators from a package. One handy use for this function is to, for example, globally attach all infix operators from a function using library, like so:

```
library(packagename, include.only = pkg_lsf("packagename", type="inops"))
```

Usage

```
pkgs %installed in% lib.loc

pkg_get_deps(
   package,
   lib.loc = .libPaths(),
   deps_type = c("LinkingTo", "Depends", "Imports"),
   base = FALSE,
   recom = FALSE,
   rstudioapi = FALSE
)

pkg_lsf(package, type, lib.loc = .libPaths())
```

Arguments

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

lib.loc character vector specifying library search path (the location of R library trees to

search through).

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

package a single string giving the package name.

deps_type a character vector, giving the dependency types to be used.

Defaults to c("LinkingTo", "Depends", "Imports").

The order of the character vector given in deps_type affects the order of the

returned character vector; see Details sections.

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base logical, indicating whether base/core R should be included (TRUE), or not in-

cluded (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 con-

sideration.

rstudioapi logical, indicating whether the rstudioapi R package should be included (TRUE),

or not included (FALSE; the default).

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.

Details

For pkg_get_deps():

If using the pkgs_get_deps() function to fill in the dependencies argument of the import_as function, one may want to know the how character vector returned by pkgs_get_deps() is ordered. The order is determined as follows.

For each string in argument deps_type, the package names in the corresponding field of the Description file are extracted, in the order as they appear in that field.

The order given in argument deps_type also affects the order of the returned character vector: The default,

c("LinkingTo", "Depends", "Imports"),

means the package names are extracted from the fields in the following order:

- 1. "LinkingTo";
- 2. "Depends";
- 3. "Imports".

The unique (thus non-repeating) package names are then returned to the user.

Value

For pkgs %installed in% lib.loc:

Returns a named logical vector, with the names giving the package names, and where the value TRUE indicates a package is installed, and the value FALSE indicates a package is not installed.

For pkg_get_deps():

A character vector of unique dependencies.

For pkg_lsf():

Returns a character vector of exported function names in the specified package.

References

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

See Also

```
tinyoperations_import()
```

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Examples

```
## Not run:
pkgs <- pkg_get_deps("devtools")
pkgs %installed in% .libPaths()
pkg_lsf("devtools", "all")
## End(Not run)</pre>
```

report_inops

Report infix operators

Description

The report_inops() function returns a data.frame listing the infix operators defined in the current environment (like the global environment, or the environment within a function), or a user specified environment. It also reports from which packages the infix operators came from.

Usage

```
report_inops(env)
```

Arguments

env

an optional environment to give, where the function should look for infix oper-

ators.

When not specified, the current environment (like the global environment, or the

environment within a function) is used.

Value

A data.frame.

See Also

```
tinyoperations_misc()
```

```
## Not run:
report_inops()
## End(Not run)
```

24 source_selection

n	
---	--

Description

The source_selection() function is the same as base R's source function, except that it allows only placing the selected objects and functions into the current environment, instead of all objects.

The objects to be selected can be specified using any combination of the following:

- by supplying a character vector of exact object names to the select argument.
- by supplying a character vector of regex patterns to the regex argument.
- by supplying a character vector of fixed patterns to the fixed argument.

Note that the source_selection() function does NOT suppress output (i.e. plots, prints, messages) from the sourced module file.

Usage

```
source_selection(lst, select = NULL, regex = NULL, fixed = NULL)
```

Arguments

lst	named list, giving the arguments to be passed to the source function.
100	named hist, giving the digaments to be pussed to the source function.

The local argument should not be included in the list.

select a character vector, giving the exact names of the functions or objects appearing

in the script, to expose to the current environment.

regex a character vector of regex patterns (see about_search_regex).

These should give regular expressions that match to the names of the functions

or objects appearing in the script, to expose to the current environment. For example, to expose the following methods to the current environment, mymethod.numeric() and mymethod.character() from generic mymethod(),

one could specify regex = "^mymethod".

fixed a character vector of fixed patterns (see about_search_fixed).

These should give fixed expressions that match to the names of the functions or

objects appearing in the script, to expose to the current environment. For example, to expose the following methods to the current environment, mymethod.numeric() and mymethod.character() from generic mymethod(),

one could specify fixed= "mymethod".

Details

One can specify which objects to expose using arguments select, regex, or fixed. The user can specify all 3 of them, but at least one of the 3 must be specified. It is not a problem if the specifications overlap.

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Value

Any specified objects will be placed in the current environment (like the Global environment, or the environment within a function).

See Also

```
tinyoperations_misc, base::source()
```

Examples

```
exprs <- expression({</pre>
helloworld = function()print("helloworld")
goodbyeworld <- function() print("goodbye world")</pre>
`%s+test%` <- function(x,y) stringi::`%s+%`(x,y)
`%s*test%` <- function(x,y) stringi::`%s*%`(x,y)
mymethod <- function(x) UseMethod("mymethod", x)</pre>
mymethod.numeric <- function(x)x * 2
mymethod.character \leftarrow function(x)chartr(x, old = "a-zA-Z", new = "A-Za-z")
})
source_selection(list(exprs=exprs), regex = "^mymethod")
mymethod(1)
mymethod("a")
temp.fun <- function(){</pre>
  source_selection(list(exprs=exprs), regex = "^mymethod", fixed = c("%", ":="))
  ls() \mbox{\tt\#} list all objects residing within the function definition
}
temp.fun()
temp.fun <- function(){</pre>
  source_selection(list(exprs=exprs), select = c("helloworld", "goodbyeworld"))
  ls() # list all objects residing within the function definition
temp.fun()
```

strcut_loc

Cut strings

Description

The strcut_loc() function cuts every string in a character vector around a location range loc, such that every string is cut into the following parts:

- the sub-string **before** loc;
- the sub-string at loc itself;
- the sub-string after loc.

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The location range loc would usually be matrix with 2 columns, giving the start and end points of some pattern match.

The strcut_brk() function (a wrapper around stri_split_boundaries) cuts every string into individual text breaks (like character, word, line, or sentence boundaries).

The main difference between the strcut_ - functions and stri_split / strsplit, is that the latter generally removes the delimiter patterns in a string when cutting, while the strcut_-functions do not attempt to remove parts of the string by default, they only attempt to cut the strings into separate pieces. Moreover, the strcut_ - functions always return a matrix, not a list.

Usage

```
strcut_loc(str, loc, fill_loc = TRUE)
strcut_brk(str, brk = "chr", ...)
```

Arguments

str

a string or character vector.

loc

Either one of the following:

- the result from the stri_locate_ith function.
- a matrix of 2 integer columns, with nrow(loc)==length(str), giving the location range of the middle part.
- a vector of length 2, giving the location range of the middle part.

fill_loc

logical, indicating what should be done if for some row i, loc[i,] is c(NA, NA).

- If TRUE, c(NA, NA) in loc[i,] is translated to c(1, nc[i]), where nc[i] is the number of characters of str[i]
- If FALSE, strcut_loc() will return c(NA, NA, NA) for when loc[i,] is c(NA, NA).

brk

a single string, giving one of the following:

- "chr": attempts to split string into individual characters.
- "line": attempts to split string into individual lines (NOTE: this is somewhat locale dependent).
- "word": attempts to split string into individual words (NOTE: this is highly locale dependent!).
- "sentence": attempts to split string into individual sentences (NOTE: this is highly locale dependent!).

For information on the boundary rules and definitions, please see:

The ICU User Guide on Boundary Analysis

(https://unicode-org.github.io/icu/userguide/boundaryanalysis/)

additional settings for stri_opts_brkiter

Value

For the strcut_loc() function:

A character matrix with length(str) rows and 3 columns:

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- the first column contains the sub-strings **before** loc;
- the second column contains the sub_strings at loc;
- the third and last column contains the sub-strings after loc.

For the strcut_brk() function:

A character matrix with length(str) rows and a number of columns equal to the maximum number of pieces str was cut in.

See Also

```
tinyoperations_strings()
```

Examples

```
x <- rep(paste0(1:10, collapse=""), 10)
print(x)
loc <- stri_locate_ith(x, 1:10, fixed = as.character(1:10))
strcut_loc(x, loc)
strcut_loc(x, c(5,5))

test <- "The\u00a0above-mentioned features are very useful. " %s+%
"Spam, spam, eggs, bacon, and spam. 123 456 789"
strcut_brk(test, "line")
strcut_brk(test, "word")
strcut_brk(test, "sentence")
strcut_brk(test, "chr")</pre>
```

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)
```

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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.

See Also

```
tinyoperations_strings()
```

```
# re-ordering characters in strings ====
x <- c("Hello world", "Goodbye world")</pre>
print(x)
mat <- strcut_brk(x, "chr")</pre>
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank</pre>
print(sorted)
stri_join_mat(sorted, margin=1)
stri_join_mat(sorted, margin=2)
# re-ordering words ====
x <- c("Hello everyone", "Goodbye everyone")
mat <- strcut_brk(x, "word")</pre>
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank
print(sorted)
\label{lem:stri_c_mat} stri\_c\_mat(sorted, margin=1) \ \# <- \ alias \ for \ stri\_join\_mat
stri_c_mat(sorted, margin=2)
# re-ordering sentences ====
```

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```
x <- c("Hello, who are you? Oh, really?! Cool!", "I don't care. But I really don't.")
print(x)
mat <- strcut_brk(x, "sentence")
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
stri_paste_mat(sorted, margin=2)</pre>
```

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.

For example: suppose a string has 3 instances of some pattern; then if $i \ge 3$ the third instance will be located, and if $i \le -3$ the first instance will be located.

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

Do not supply the arguments omit_no_match, get_length, or pattern, as they are already specified internally. Supplying these arguments anyway will result in an error.

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, two NAs if no matches are found, and also two NAs if str is NA.

See Also

```
tinyoperations_strings()
```

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

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```
p <- rep("AB", 2)
out <- stri_locate_ith(x, c(1, -1), regex=p, case_insensitive=TRUE)
print(out)
substr(x, out[,1], out[,2])</pre>
```

```
# 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>
```

stri_rgx

Pattern specifications for string related infix operators.

Description

The %s-% and %s/% operators, as well as the string detection operators (%s{}%, %s!{}%), perform pattern matching for some purpose, where the pattern is given on the right hand side.

When a character vector or string is given on the right hand side, this is interpreted as case-sensitive regex patterns from stringi.

Instead of giving a string or character vector of regex patterns, one can also supply a list to specify exactly how the pattern should be interpreted. The list should use the exact same argument convention as stringi.

For example:

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

All arguments in the list are simply passed to the appropriate functions in stringi.

For example:

```
x %s/% p
```

counts how often regular expression specified in character vector p occurs in x, whereas the following,

```
x %s/% list(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.

Related to the above, tinyoperations adds some convenience functions based on the stri_opts_ - functions in stringi (convenient in the sense they already have argument names, thus allowing

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for auto code completion):

```
stri_rgx(p, ...) is equivalent to c(list(regex = p), ...)
stri_fxd(p, ...) is equivalent to c(list(fixed = p), ...)
stri_cll(p, ...) is equivalent to c(list(coll = p), ...)
stri_chrcls(p, ...) is equivalent to c(list(charclass = p), ...)
```

With the ellipsis (...) being passed to the appropriate $stri_{opts-functions}$ when it matches their arguments.

Usage

```
stri_rgx(
  case_insensitive,
  comments,
  dotall,
  multiline,
  time_limit,
  stack_limit,
)
stri_fxd(p, case_insensitive, overlap, ...)
stri_cll(
  р,
  locale,
  strength,
  alternate_shifted,
  french,
  uppercase_first,
  case_level,
  numeric,
  normalization,
)
stri_chrcls(p, ...)
```

Arguments

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Value

A list.

See Also

```
tinyoperations_strings()
```

Examples

```
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
p \leftarrow rep("a|e|i|o|u", 2) \# same as p \leftarrow list(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=""))</pre>
print(x)
x %s/% list(regex = rep("A|E|I|0|U", 2), case_insensitive = TRUE)
x %s/% stri_rgx(rep("A|E|I|0|U", 2), case_insensitive = TRUE)
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
p <- list(regex = c("A", "A"), case_insensitive=TRUE)</pre>
x %s{}% p
x %s!{}% p
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
p <- list(fixed = c("A", "A"), case_insensitive=TRUE)</pre>
x %s{}% p
x %s!{}% p
```

str_arithmetic

String arithmetic

Description

String arithmetic operators.

The x %s+% y operator is exported from stringi, and concatenates character vectors x and y.

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

The x %s *% n operator is exported from stringi, and duplicates each string in x n times, and concatenates the results.

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

The e1 %s\$% e2 operator is exported from stringi, and provides access to stri_sprintf in the form

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of an infix operator.

Usage

```
x %s-% p
```

Arguments

x a string or character vector.

p either a list with stringi arguments (see stri_rgx), or else a character vector of the same length as x with regular expressions.

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.

The %s\$% operator returns a character vector.

See Also

```
tinyoperations_strings()
```

Examples

```
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
y <- c("a", "b")
p <- \ \text{rep}("a|e|i|o|u",\ 2) \ \# \ \text{same as} \ p <- \ \text{list(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 <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
y <- "a"
# pattern that ignores case:
p <- list(regex=rep("A|E|I|0|U", 2), case_insensitive = 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 35

str_subset_ops Sta	ring subsetting operators
--------------------	---------------------------

Description

String subsetting operators.

The string %ss% ind 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 character vector x.

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

Usage

```
string %ss% ind
x %sget% ss
```

x %strim% ss

Arguments

string	a single string.
ind	a numeric vector giving the subset indices.
X	a character vector.
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 and non-missing integers, or be coerce-able to such integers. (thus negative integers, and missing values are not allowed; decimal numbers will be converted to integers). 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

Details

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

the right side to be extracted/removed from x.

Value

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

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

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

See Also

```
tinyoperations_strings()
```

Examples

```
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
ss <- c(2,3)
x %sget% ss
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
ss <- c(1,0)
x %sget% ss
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
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
"hello" %ss% 5:1
```

 $subset_if$

The subset_if operators and the unreal in place modifier

Description

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

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

subset_if 37

```
The x \%unreal =\% repl operator modifies all unreal (NA, NaN, Inf, -Inf) values of x with replacement value repl. Thus,
```

```
x %unreal =% repl,
is the same as,
x[is.na(x)|is.nan(x)|is.infinite(x)] <- repl</pre>
```

Usage

```
x %[if]% cond
x %[!if]% cond
x %unreal =% repl
```

Arguments

x a vector, matrix, or array.

cond a (possibly anonymous) function that returns a logical vector of the same

length/dimensions as x. For example: $\(x)x>0$.

repl the replacement value.

Value

For the **subset-if** operators:

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

For the x %unreal =% repl operator:

The x %unreal =% repl operator does not return any value:

It is an in-place modifier, and thus modifies x directly. The object x is modified such that all NA, NaN, Inf, and -Inf elements are replaced with repl.

See Also

```
tinyoperations_dry()
```

Examples

```
x <- c(-10:9, NA, NA)
object_with_very_long_name <- matrix(x, ncol=2)
print(object_with_very_long_name)
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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tinyoperations_dry

The tinyoperations "DRY" functionality

Description

"Don't Repeat Yourself", sometimes abbreviated as "DRY", is the coding principle not to write unnecessarily repetitive code. To help you in that effort, the tinyoperations R package introduces a few functions:

- The transform_if function
- The subset_if operators and the in-place unreal modifier operator.
- The generalized in-place (mathematical) modification operator.

Usage

tinyoperations_dry()

See Also

tinyoperations_help()

tinyoperations_help

The tinyoperations help page

Description

Welcome to the tinyoperations introduction help page!

The tinyoperations R-package adds adds some functions and infix operators to help in your programming etiquette.

It primarily focuses on 4 things:

- (1) Safer decimal (in)equality testing, safer atomic conversions, and other functions for safer coding; see tinyoperations_safer.
- (2) A new package import system, that combines the benefits of aliasing a package with the benefits of attaching a package; see tinyoperations_import
- (3) Extending the string manipulation capabilities of the stringi R package; see tinyoperations_strings.
- (4) Reducing repetitive code; see tinyoperations_dry.

And some miscellaneous functionality; see tinyoperations_misc.

The tinyoperations R-package has only one dependency, namely stringi. Most functions in this R-package are fully vectorized and optimized, and have been well documented.

Please also have a look at the GitHub page before using this package:

https://github.com/tony-aw/tinyoperations

Usage

tinyoperations_help()

tinyoperations_import The tinyoperations import system

Description

The tinyoperations R package introduces a new package import system.

One can use a package without attaching the package (i.e. using :: or using a package alias), or one can attach a package (i.e. using library() or require()). The advantages and disadvantages of loading without attaching a package versus attaching a package - at least those relevant here - can be compactly presented in the following table:

```
alias / ::
                                                                                           attaching
prevent masking functions from other packages
                                                                              Yes (+)
                                                                                           No (-)
prevent masking core R functions
                                                                              Yes (+)
                                                                                           No (-)
clarify which function came from which package
                                                                              Yes (+)
                                                                                           No (-)
place/expose functions only in current environment instead of globally
                                                                              Yes (+)
                                                                                           No (-)
prevent namespace pollution
                                                                              Yes (+)
                                                                                           No (-)
minimize typing - especially for infix operators
(i.e. typing package:: '%op%'(x, y) instead of x %op% y is cumbersome)
                                                                                           Yes (+)
use multiple related packages,
without constantly switching between package prefixes
                                                                            No (-)
                                                                                        | Yes (+)
NOTE: + = advantage, - = disadvantage
```

What tinyoperations attempts to do with its import system, is to somewhat find the best of both worlds. It does this by introducing the following functions:

- import_as: Allow a main package + its re-exports + its dependencies + its enhances + its extensions to be loaded under a single alias. This essentially combines the attaching advantage of using multiple related packages, whilst keeping most advantages of aliasing a package.
- import_inops: Expose infix operators from a package or an alias object to the current environment. This gains the attaching advantage of less typing, whilst simultaneously avoiding the disadvantage of attaching functions from a package globally.
- import_data: Directly return a data set from a package, to allow straight-forward assignment.

Furthermore, there are two miscellaneous import_ - functions: import_LL and import_int. And there are also some additional helper functions for the package import system, see x.import and pkgs.

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Usage

```
tinyoperations_import()
```

See Also

tinyoperations_help

Examples

```
## Not run:
# loading "tidytable" + "data.table" under alias "tdt.":
import_as(
    ~ tdt., "tidytable", dependencies = "data.table"
)

# exposing operators from "magrrittr" to current environment:
import_inops("magrittr")

# directly assigning dplyr's "starwars" dataset to object "d":
d <- import_data("dplyr", "starwars")

# see it in action:
d %>% tdt.$filter(species == "Droid") %>%
    tdt.$select(name, tdt.$ends_with("color"))

### End(Not run)
```

tinyoperations_misc

The tinyoperations miscellaneous functionality

Description

Some additional functions provided by the tinyoperations R package:

- Infix logical operators for exclusive-or, not-and, not-in, number-type, and string-type.
- Infix operators for row- and column-wise re-ordering of matrices.
- Report infix operators present in the current environment, or a specified environment.
- source_selection to source only selected objects.

Usage

```
tinyoperations_misc()
```

See Also

```
tinyoperations_help()
```

tinyoperations_safer 41

Description

To help make your code safer, the tinyoperations R package introduces a few functions:

- Safer decimal (in)equality testing.
- Atomic type casting without stripping attributes.
- The lock_TF function to set and lock T and F to TRUE and FALSE.
- The %<-c% operator to assign locked constants.

Usage

tinyoperations_safer()

See Also

tinyoperations_help()

tinyoperations_strings

The tinyoperations expansion of the 'stringi' R package

Description

The tinyoperations R package adds some functions and operators to extend the functionality of the stringi R package:

- Infix operators for string arithmetic.
- Infix operators for string sub-setting.
- Infix operators for row- and column-wise re-ordering of matrices.
- The tinyoperations 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 strcut_-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 the functions are written in R, they have been aggressively optimized to be in the same order of speed as the other stringi functions.

42 transform_if

Usage

```
tinyoperations_strings()
```

See Also

```
tinyoperations_help()
```

transform_if

The transform_if function

Description

The transform_if() function transforms an object x, based on the logical result (TRUE, FALSE, NA) of condition function cond(x) or logical vector cond, such that:

- For every value where cond(x)==TRUE / cond==TRUE, function yes(x) is run or scalar yes is returned.
- For every value where cond(x)==FALSE / cond==FALSE, function no(x) is run or scalar no is returned.
- For every value where cond(x)==NA / cond==NA, function other(x) is run or scalar other is returned.

Usage

```
transform_if(x, cond, yes = function(x) x, no = function(x) x, other = NA)
```

Arguments

cond either an object of class logical with the same length as x,

or a (possibly anonymous) function that returns an object of class logical with

the same length as x. For example: (x)x>0.

yes the (possibly anonymous) transformation function to use when function cond(x) = TRUE

/ logical cond==TRUE.

Alternatively, one can also supply an atomic scalar. If argument yesis not specified, it defaults to $\(x)x$.

no the (possibly anonymous) transformation function to use when function cond(x) = FALSE

/ logical cond==FALSE.

Alternatively, one can also supply an atomic scalar. If argument no is not specified, it defaults to $\(x)x$.

other the (possibly anonymous) transformation function to use when function cond(x)

/ logical cond returns NA.

Alternatively, one can also supply an atomic scalar. If argument other is not specified, it defaults to NA.

Note that function other(x) is run or scalar other is returned when function

cond(x) or logical cond is NA, not necessarily when x itself is NA.

x.import 43

Details

```
Be careful with coercion! For example the following code: x <- c("a", "b") transform_if(x, \(x)x=="a", as.numeric, as.logical) returns: [1] NA NA due to the same character vector being given 2 incompatible classes.
```

Value

The transformed vector, matrix, or array (attributes are conserved).

See Also

```
tinyoperations_dry()
```

Examples

```
x <- c(-10:9, NA, NA)
object <- matrix(x, ncol=2)
attr(object, "helloworld") <- "helloworld"
print(object)
y <- 0
z <- 1000

object |> transform_if(\(x)x>y, log, \(x)x^2, \(x)-z)
object |> transform_if(object > y, log, \(x)x^2, -z) # same as previous line
```

x.import

Helper functions for the tinyoperations package import system

Description

The help.import() function finds the help file for functions in an alias object or exposed infix operators.

The attr.import() function gets one specific special attributes or all special attributes from an alias object returned by import_as.

The is.tinyimport() function checks if an alias object or an exposed function is of class tinyimport; i.e. if it is an object produced by the import_as or import_inops function.

44 x.import

Usage

```
help.import(..., i, alias)
attr.import(alias, which = NULL)
is.tinyimport(x)
```

Arguments

further arguments to be passed to help.either one of the following:

- a function (use back-ticks when the function is an infix operator). Examples: myfun, `%operator%`, myalias.\$some_function. If a function, the alias argument is ignored.
- a string giving the function name or topic (i.e. "myfun", "thistopic"). If a string, argument alias must be specified also.

alias the alias object as created by the import_as function.

which The attributes to list. If NULL, all attributes will be returned.

Possibilities: "pkgs", "conflicts", "versions", "args", and "ordered_object_names".

x an object produced by import_as or import_inops.

Details

For help.import(...):

Do not use the topic / package and i / alias arguments together. It's either one set or the other.

Value

```
For help.import():
```

Opens the appropriate help page.

```
For is.tinyimport():
```

Returns TRUE if x is produced by functions import_as or import_inops, and returns FALSE if it is not.

```
For attr.import(alias, which = NULL):
```

All special attributes of the given alias object are returned as a list.

```
For attr.import(alias, which = "pkgs"):
```

Returns a list with 3 elements:

- packages_order: a character vector of package names, giving the packages in the order they were loaded in the alias object.
- main_package: a string giving the name of the main package. Re-exported functions, if present, are loaded together with the main package.
- re_exports.pkgs: a character vector of package names, giving the packages from which the re-exported functions in the main package were taken.

x.import 45

```
For attr.import(alias, which = "conflicts"):
```

The order in which packages are loaded in the alias object (see attribute pkgs\$packages_order) matters: Functions from later named packages overwrite those from earlier named packages, in case of conflicts.

The "conflicts" attribute returns a data.frame showing exactly which functions overwrite functions from earlier named packages, and as such "win" the conflicts.

```
For attr.import(alias, which = "versions"):
```

A data.frame, giving the version of every package loaded in the alias, ignoring re-exports.

```
For attr.import(alias, which = "args"):
```

Returns a list of input arguments. These were the arguments supplied to import_as when the alias object in question was created.

```
For attr.import(alias, which = "ordered_object_names"):
```

Gives the names of the objects in the alias, in the order as they were loaded.

Only unique names are given, thus conflicting objects only appear once.

(Note that if argument re_exports is TRUE, re-exported functions are loaded when the main package is loaded, thus changing this order slightly.)

See Also

```
tinyoperations_import()
```

Examples

```
## Not run:
import_as(mr., "magrittr")
import_inops("magrittr")
`:=` <- data.table::`:=`

is.tinyimport(mr.) # returns TRUE
is.tinyimport(`%>%`) # returns TRUE
is.tinyimport(`:=`) # returns FALSE

attr.import(mr.)
attr.import(mr., which="conflicts")

help.import(i=mr.$add)
help.import(i="%>%`)
help.import(i="add", alias=mr.)

## End(Not run)
```

46 %s{}%

%s{}%

String detection operators

Description

The $x %s{}$ p operator checks for every string in character vector x if the pattern defined in p is present.

The x%s!{}% p operator checks for every string in character vector x if the pattern defined in p is NOT present.

Usage

```
x %s{}% p
x %s!{}% p
```

Arguments

x a string or character vector.

p either a list with stringi arguments (see stri_rgx), or else a character vector of the same length as x with regular expressions.

Value

The $x %s{}$ p and $x %s!{}$ p operators return logical vectors.

See Also

```
tinyoperations_strings()
```

Examples

```
# simple pattern ====

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
x %s{}% "a"
x %s!{}% "a"
which(x %s{}% "a")
which(x %s{}% "a")
x[x %s{}% "a"]
x[x %s!{}% "a"]

x %s{}% "1"
which(x %s{}% "1")
which(x %s!{}% "1")
x[x %s{}% "1"]
x[x %s!{}% "1"]</pre>
```

%s{}%

```
# ignore case pattern ====
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))</pre>
print(x)
p <- list(regex = c("A", "A"), case_insensitive=TRUE)</pre>
x %s{}% p
x %s!{}% p
which(x %s{}% p)
which(x %s!{}% p)
x[x %s{}% p]
x[x %s!{}% p]
# multi-character pattern ====
x \leftarrow c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- list(regex = rep("AB", 2), case_insensitive=TRUE)</pre>
x %s{}% p
x %s!{}% p
which(x %s{}% p)
which(x %s!{}% p)
x[x %s{}% p]
x[x %s!{}% p]
```

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