

# Package ‘tinycodet’

September 29, 2023

**Title** R package that adds a few tiny functions to help in your programming etiquette

**Version** 0.0.0.9

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**Description** The 'tinycodet' R-

package is a tiny little R package that adds a few functions to help in your coding etiquette. It primarily focuses on 4 things.

1) Safer decimal (in)equality testing, safer atomic conversions, and other functions for safer coding.

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 'tinycodet' R-package has only one dependency, namely 'stringi'.

Most functions in this R-package are fully vectorized and optimized, and have been well documented.

**License** MIT + file LICENSE

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.2.3

**Suggests** rlang,

knitr,

rmarkdown,

tinytest,

pkgdown,

devtools,

fastverse,

gamair

**VignetteBuilder** knitr

**Depends** R (>= 4.1.0)

**Imports** stringi (>= 1.7.12)

**URL** <https://github.com/tony-aw/tinycodet>, <https://tony-aw.github.io/tinycodet>

**BugReports** <https://github.com/tony-aw/tinycodet/issues>

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atomic_conversions	<i>Safer atomic type casting</i>
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## Description

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 `tinycodet` package preserve all attributes - except the "class" attribute.

The functions are as follows:

- `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 = sqrt(.Machine$double.eps))
```

### Arguments

<code>x</code>	vector, matrix, array (or a similar object where all elements share the same class).
<code>...</code>	further arguments passed to or from other methods.
<code>tol</code>	the tolerance.

### Value

The converted object.

### See Also

[tinycodet\\_safer\(\)](#)

### Examples

```
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)
```

**Description**

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

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

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

except for one aspect.

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.

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 and Python.

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

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`

`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

[tinycodet\\_safer\(\)](#)

### Examples

```
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 # 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 <- cbind(x-0.1, x+0.1)
x %d{%} bnd
x %d!{%} bnd

# These operators still work for non-decimal 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
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
x %d>=% y
```

---

import_as	<i>Load R-package + its re-exports + its dependencies + its extensions under one alias</i>
-----------	--

---

## Description

The `import_as()` function imports the namespace of an R package, and optionally also its re-exports, dependencies, 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,
  extensions = NULL,
  lib.loc = .libPaths(),
  loadorder = c("dependencies", "main_package", "extensions")
)
```

## Arguments

- |              |   |
|--------------|---|
| alias        | <p>a syntactically valid non-hidden name giving the alias object where the package(s) are to be loaded into.</p> <p>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.).</p> <p>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 (_).</p>   |
| main_package | <p>a single string, giving the name of the main package to load under the given alias.</p>  |
| re_exports   | <p>logical; Some R packages export functions that are not defined in their own package, but in their direct dependencies; "re-exports", if you will.</p> <p>This argument determines what the <code>import_as</code> function will do with the re-exports of the <code>main_package</code>:</p> <ul style="list-style-type: none"> <li>• If TRUE the re-exports from the <code>main_package</code> are added to the alias, even if <code>dependencies = NULL</code>. This is the default, as it is analogous to the behaviour of base R's <code>::</code> operator.</li> <li>• If FALSE, these re-exports are not added, and the user must specify the appropriate packages in argument <code>dependencies</code>.</li> </ul> |

- |              |  |
|--------------|--|
| dependencies | an optional character vector, giving the names of the dependencies of the main_package to be loaded also under the alias.<br>Defaults to NULL, which means no dependencies are loaded.<br>See <a href="#">pkg_get_deps</a> to quickly get dependencies from a package.   |
| extensions   | an optional character vector, giving the names of the extensions of the main_package to be loaded also under the alias.<br>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).<br>The lib.loc argument would usually be .libPaths().<br>See also <a href="#">loadNamespace</a> .   |
| loadorder    | the character vector<br>c("dependencies", "main_package", "extensions"),<br>or some re-ordering of this character vector, giving the relative load order of the groups of packages.<br>The default setting (which is highly recommended) is the character vector<br>c("dependencies", "main_package", "extensions"),<br>which results in the following load order: <ol style="list-style-type: none"> <li>1. The dependencies, <b>in the order specified by the dependencies argument</b>.</li> <li>2. The main_package (see argument main_package), including re-exports (if re_exports=TRUE).</li> <li>3. The extensions, <b>in the order specified by the extensions argument</b>.</li> </ol> |

## Details

### On the dependencies 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.
- 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, 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**

[tinycodet\\_import\(\)](#)

**Examples**

```
## 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)
```

---

import_data	<i>Directly return a data-set from a package</i>
-------------	--

---

**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).  
The `lib.loc` argument would usually be `.libPaths()`.  
See also [loadNamespace](#).

### Value

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

### See Also

[tinycodet\\_import\(\)](#)

### Examples

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

---

<code>import_inops</code>	<i>(un)expose infix operators from package namespace in the current environment</i>
---------------------------	---

---

### 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 package 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 `tinycodet` import system will be removed from the current environment; "regular" (i.e. user-specified) infix operators will not be touched.

### Usage

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

### Arguments

`expose`, `unexpose`

either one of the following:

- an alias object 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 tinycodet import system or not. Only infix operators exposed by the tinycodet import system will be removed.

## 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).

If using argument `unexpose`:

The infix operators specified in the given package or alias, exposed by `import_inops()`, will be removed from the current environment (like the Global environment, or the environment within a function).

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

## See Also

[tinycodet\\_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
)
```

## Arguments

exclude	<p>a character vector, giving the infix operators NOT to expose to the current environment.</p> <p>This can be handy to prevent overwriting any (user defined) infix operators already present in the current environment.</p>
include.only	<p>a character vector, giving the infix operators to expose to the current environment, and the rest of the operators will not be exposed.</p> <p>This can be handy to prevent overwriting any (user defined) infix operators already present in the current environment.</p>
overwrite	<p>logical, indicating if it is allowed to overwrite existing infix operators.</p> <ul style="list-style-type: none"> <li>• If TRUE (default), a warning is given when operators existing in the current environment are being overwritten, but the function continuous nonetheless.</li> <li>• If FALSE, an error is produced when the to be exposed operators already exist in the current environment, and the function is halted.</li> </ul>
inherits	<p>logical; indicating whether enclosed environments, especially package namespaces, should also be taken into account (TRUE), or not (FALSE). Defaults to FALSE.</p> <p>See also <a href="#">exists</a>.</p>

## 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()`, `tinycodet_import()`

**Examples**

```
## 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")

## 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**

<code>package</code>	a single string, giving the name of the package to take functions from.
<code>selection</code>	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 <a href="#">loadNamespace</a> .
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

## 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 functions 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(...)
```

or use it directly without re-assignment like so:

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

## See Also

[tinycodet\\_import\(\)](#)

## Examples

```
## 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(tinycodet ~ .internal_paste, .libPaths())
fun("hello", "world")
# Or using directly:
import_int(
  tinycodet ~ .internal_paste, .libPaths()
)("hello", "world")

## 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:

```
mtcars$mpg[mtcars$cyl>6] <- mtcars$mpg[mtcars$cyl>6]^2
```

Can now be re-written as:

```
\code{mtcars$mpg[mtcars$cyl>6\] %:=% \(\x)x^2}
```

## Usage

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

## Arguments

<code>x</code>	a variable.
<code>f</code>	a (possibly anonymous) function to be applied in-place on <code>x</code> . 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**

[tinycodet\\_dry\(\)](#)

**Examples**

```
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)
```

---

lock	<i>Lock T, lock F, or create locked constants</i>
------	---

---

**Description**

One can re-assign the values T and F. One can even run things like T <- FALSE and F <- 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
```

**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**[tinycodet\\_safer\(\)](#)**Examples**

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

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% y` 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 be 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 <a href="#">Logic</a> .
n	a numeric vector.
numtype	a single string giving the numeric type to be checked. The following options are supported: <ul style="list-style-type: none"> <li>• "~0": zero, or else a number whose absolute value is smaller than the Machine tolerance (<code>sqrt(.Machine\$double.eps)</code>).</li> <li>• "B": binary numbers (exactly 0 or exactly 1);</li> <li>• "prop": proportions - numbers between 0 and 1 (exactly 0 or 1 is also allowed);</li> <li>• "I": Integers;</li> <li>• "odd": odd integers;</li> <li>• "even": even integers;</li> <li>• "R": Real numbers;</li> <li>• "unreal": infinity, NA, or NaN;</li> </ul>
s	a character vector.
strtype	a single string giving the string type to be checked. The following options are supported: <ul style="list-style-type: none"> <li>• "empty": checks if the string only consists of empty spaces.</li> <li>• "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.</li> <li>• "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 actual number 5.0.</li> <li>• "special": checks if the string consists of only special characters.</li> </ul>

**Value**

A logical vector.

**Examples**

```
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"
n[n %=numtype% "B"]
n[n %=numtype% "prop"]
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"]

```

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 %col% x** is 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.

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

then the code

```
x %row% mat
```

will randomly shuffle the elements of every row of **x**, 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 of **x**, 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

[tinycodet\\_misc\(\)](#)

## Examples

```
# numeric matrix ====

x <- matrix(sample(1:25), nrow=5)
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)
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)
print(x)
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~% mat # sort elements of every column independently
x %col~% -mat # reverse-sort elements of every column independently

x <- matrix(sample(letters, 25), nrow=5)
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

```

---

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()` 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

<code>pkgs</code>	a character vector with the package name(s).
<code>lib.loc</code>	character vector specifying library search path (the location of R library trees to search through). The <code>lib.loc</code> argument would usually be <code>.libPaths()</code> . See also <a href="#">loadNamespace</a> .
<code>package</code>	a single string giving the package name.
<code>deps_type</code>	a character vector, giving the dependency types to be used. Defaults to <code>c("LinkingTo", "Depends", "Imports")</code> . The order of the character vector given in <code>deps_type</code> affects the order of the returned character vector; see Details sections.
<code>base</code>	logical, indicating whether base/core R should be included (TRUE), or not included (FALSE; the default).
<code>recom</code>	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.
<code>rstudioapi</code>	logical, indicating whether the <code>rstudioapi</code> R package should be included (TRUE), or not included (FALSE; the default).
<code>type</code>	The type of functions to list. Possibilities: <ul style="list-style-type: none"> <li>• <code>"inops"</code> or <code>"operators"</code>: Only infix operators.</li> <li>• <code>"regfuns"</code>: Only regular functions (thus excluding infix operators).</li> <li>• <code>"all"</code>: All functions, both regular functions and infix operators.</li> </ul>

## 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

elegantly extract R package dependencies of a package not listed on CRAN. Stack Overflow. (1 September 2023). <https://stackoverflow.com/questions/30223957/elegantly-extract-r-package-dependencies-of-a-package-not-listed-on-cran>

## See Also

`tinycodet_import()`

## Examples

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

## End(Not run)
```

---

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 operators. When not specified, the current environment (like the global environment, or the environment within a function) is used.
-----	--

## Value

A data.frame.

**See Also**[tinycodet\\_misc\(\)](#)**Examples**

```
## Not run:
report_inops()

## End(Not run)
```

---

source_selection	<i>Source selection</i>
------------------	-------------------------

---

**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 script file.

**Usage**

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

**Arguments**

<code>lst</code>	a named list, giving the arguments to be passed to the <a href="#">source</a> function. The <code>local</code> argument should not be included in the list.
<code>select</code>	a character vector, giving the exact names of the functions or objects appearing in the script, to expose to the current environment.
<code>regex</code>	a character vector of regex patterns (see <a href="#">about_search_regex</a> ). 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, <code>mymethod.numeric()</code> and <code>mymethod.character()</code> from generic <code>mymethod()</code> , one could specify <code>regex = "^mymethod"</code> .

**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.

## Value

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

## See Also

[tinycodet\\_misc](#), `base::source()`

## Examples

```
exprs <- expression({
  helloworld = function()print("helloworld")
  goodbyeworld <- function() print("goodbye world")
  `s+test` <- function(x,y) stringi::`s+`(x,y)
  `s*test` <- function(x,y) stringi::`s*`(x,y)
  mymethod <- function(x) UseMethod("mymethod", x)
  mymethod.numeric <- function(x)x * 2
  mymethod.character <- 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(){
  source_selection(list(exprs=exprs), regex = "^mymethod", fixed = c("%", ":="))
  ls() # list all objects residing within the function definition
}
temp.fun()
```

```
temp.fun <- function(){
  source_selection(list(exprs=exprs), select = c("helloworld", "goodbyeworld"))
  ls() # list all objects residing within the function definition
}
temp.fun()
```



---

strcut_loc	<i>Cut strings</i>
------------	--------------------

---

## 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`.

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

- |                       |  |
|-----------------------|--|
| <code>str</code>      | a string or character vector.  |
| <code>loc</code>      | Either one of the following: <ul style="list-style-type: none"> <li>• the result from the <a href="#">stri_locate_ith</a> function.</li> <li>• a matrix of 2 integer columns, with <code>nrow(loc)==length(str)</code>, giving the location range of the middle part.</li> <li>• a vector of length 2, giving the location range of the middle part.</li> </ul>  |
| <code>fill_loc</code> | logical, indicating what should be done if for some row <code>i</code> , <code>loc[i, ]</code> is <code>c(NA, NA)</code> . <ul style="list-style-type: none"> <li>• If <code>TRUE</code>, <code>c(NA, NA)</code> in <code>loc[i, ]</code> is translated to <code>c(1, nc[i])</code>, where <code>nc[i]</code> is the number of characters of <code>str[i]</code></li> <li>• If <code>FALSE</code>, <code>strcut_loc()</code> will return <code>c(NA, NA, NA)</code> for when <code>loc[i, ]</code> is <code>c(NA, NA)</code>.</li> </ul> |
| <code>brk</code>      | a single string, giving one of the following: <ul style="list-style-type: none"> <li>• <code>"chr"</code> or <code>"character"</code>: attempts to split string into individual characters.</li> <li>• <code>"line"</code> or <code>"line_break"</code>: attempts to split string into individual lines (NOTE: this is somewhat locale dependent).</li> </ul>  |

- "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!).

about search: boundaries

... additional settings for [stri\\_opts\\_brkiter](#)

### Value

For the `strcut_loc()` function:

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

- 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

[tinycodet\\_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))
strcut_loc(x, c(NA, NA), fill_loc = TRUE)
strcut_loc(x, c(NA, NA), fill_loc = FALSE)

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")
```

---

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 row-wise (`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.

**See Also**

[tinycodet\\_strings\(\)](#)

**Examples**

```
# re-ordering characters in strings ====

x <- c("Hello world", "Goodbye world")
print(x)
mat <- strcut_brk(x, "chr")
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row% rank
print(sorted)
stri_join_mat(sorted, margin=1)
stri_join_mat(sorted, margin=2)

#####

# re-ordering words ====

x <- c("Hello everyone", "Goodbye everyone")
print(x)
mat <- strcut_brk(x, "word")
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
```

```

sorted <- mat %row% rank
print(sorted)
stri_c_mat(sorted, margin=1) # <- alias for stri_join_mat
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 <- 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)

```

---

stri_locate_ith	<i>Locate <math>i^{th}</math> Pattern Occurrence</i>
-----------------	--

---

## 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

<code>str</code>	a string or character vector.
<code>i</code>	a number, or a numeric vector of the same length as <code>str</code> . 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 >= 3` the third instance will be located,

and if `i <= -3` the first instance will be located.

... more arguments to be supplied to [stri\\_locate](#).  
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](#).

about search: `regex`

about search: `fixed`

about search: `coll`

about search: `charclass`

## Details

### Special note regarding charclass

The `stri_locate_ith()` function is based on [stri\\_locate\\_all](#). This generally gives results consistent with [stri\\_locate\\_first](#) or [stri\\_locate\\_last](#), but the exception is when `charclass` pattern is used.

Where the functions [stri\\_locate\\_first](#) or [stri\\_locate\\_last](#) give the location of the first or last single character matching the `charclass` (respectively), [stri\\_locate\\_all](#) gives the start and end of **consecutive** characters.

The `stri_locate_ith()` is in this aspect more in line with [stri\\_locate\\_all](#), as it gives the  $i^{th}$  set of consecutive characters.

## 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

[tinycodet\\_strings\(\)](#)

## Examples

```
# simple pattern ====
```

```
x <- rep(paste0(1:10, collapse=""), 10)
print(x)
out <- stri_locate_ith(x, 1:10, regex = as.character(1:10))
cbind(1:10, out)
```

```
x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- rep("a|e|i|o|u", 2)
out <- stri_locate_ith(x, c(-1, 1), regex=p)
```

```

print(out)
substr(x, out[,1], out[,2])

#####

# ignore case pattern ====

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- rep("A|E|I|O|U", 2)
out <- stri_locate_ith(x, c(1, -1), regex=p, case_insensitive=TRUE)
substr(x, out[,1], out[,2])

#####

# multi-character pattern ====

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
# multi-character pattern:
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])

#####

# 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)

```

---

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.

`tinycodet` adds some convenience functions based on the `stri_opts_` - functions in `stringi`:

- `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 `stringi`-functions when it matches their arguments.

## Usage

```
stri_rgx(
  p,
  case_insensitive,
  comments,
  dotall,
  multiline,
  time_limit,
  stack_limit,
  ...
)

stri_fxd(p, case_insensitive, overlap, ...)

stri_cll(
  p,
```

```

    locale,
    strength,
    alternate_shifted,
    french,
    uppercase_first,
    case_level,
    numeric,
    normalization,
    ...
)

stri_chrcls(p, ...)

```

### Arguments

**p** a character vector giving the pattern to search for.  
[about search: regex](#)  
[about search: fixed](#)  
[about search: coll](#)  
[about search: charclass](#)

**case\_insensitive** see [stri\\_opts\\_regex](#) and [stri\\_opts\\_fixed](#).

**comments, dotall, multiline** see [stri\\_opts\\_regex](#).

**time\_limit, stack\_limit** see [stri\\_opts\\_regex](#).

**...** additional arguments not part of the `stri_opts` - functions to be passed here.  
 For example: `max_count`

**overlap** see [stri\\_opts\\_fixed](#).

**locale, strength, alternate\_shifted** see [stri\\_opts\\_collator](#).

**french, normalization, numeric** see [stri\\_opts\\_collator](#).

**uppercase\_first, case\_level** see [stri\\_opts\\_collator](#).

### Value

A list.

### See Also

[tinycodet\\_strings\(\)](#)

### Examples

```

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 <- 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=""))
print(x)
x %s/% list(regex = rep("A|E|I|O|U", 2), case_insensitive = TRUE)
x %s/% stri_rgx(rep("A|E|I|O|U", 2), case_insensitive = TRUE)

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- list(regex = c("A", "A"), case_insensitive=TRUE)
x %s{%} p
x %s!{%} p

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- list(fixed = c("A", "A"), case_insensitive=TRUE)
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 of an infix operator.

**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.

about search: [regex](#)  
 about search: [fixed](#)  
 about search: [coll](#)  
 about search: [charclass](#)

## Value

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

## See Also

[tinycodet\\_strings\(\)](#)

## Examples

```
x <- 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 <- 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=""))
print(x)
y <- "a"
# pattern that ignores case:
p <- list(regex=rep("A|E|I|O|U", 2), case_insensitive = TRUE)
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.
```

## Description

String subsetting operators.

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

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

## Usage

```
x %sget% ss
```

```
x %strim% ss
```

## Arguments

<code>x</code>	a character vector.
<code>ss</code>	a vector of length 2, or a matrix with 2 columns with <code>nrow(ss)==length(x)</code> . The object <code>ss</code> 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 <code>ss</code> gives the number of characters counting from the left side to be extracted/removed from <code>x</code> . The second element/column of <code>ss</code> gives the number of characters counting from the right side to be extracted/removed from <code>x</code> .

## Details

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

## Value

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`.

## See Also

[tinycodet\\_strings\(\)](#)

## Examples

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

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

```

---

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.

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`

## Usage

`x %[if]% cond`

`x %[!if]% cond`

`x %unreal =% repl`

**Arguments**

<code>x</code>	a vector, matrix, or array.
<code>cond</code>	a (possibly anonymous) function that returns a logical vector of the same length/dimensions as <code>x</code> . For example: <code>\(x)x&gt;0</code> .
<code>repl</code>	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**

[tinycodet\\_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)
```

---

tinycodet\_dry

---

*The tinycodet "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 `tinycodet` 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**

```
tinycodet_dry()
```

**See Also**

[tinycodet\\_help\(\)](#)

**Examples**

```
object <- matrix(c(-9:8, NA, NA) , ncol=2)

# in base R:
ifelse( # repetitive, and gives unnecessary warning
  is.na(object>0), -Inf,
  ifelse(
    object>0, log(object), object^2
  )
)
mtcars$mpg[mtcars$cyl>6] <- (mtcars$mpg[mtcars$cyl>6])^2 # long

# with tinycodet:
object |> transform_if(\(x)x>0, log, \(x)x^2, \(x) -Inf) # compact & no warning
mtcars$mpg[mtcars$cyl>6] %:=% \(x)x^2 # short
```

---

`tinycodet_help`

*The tinycodet help page*

---

**Description**

Welcome to the tinycodet introduction help page!

The 'tinycodet' R-package is a tiny little R package that adds some functions to help in your coding etiquette.

It primarily focuses on 4 things:

(1) Safer decimal (in)equality testing, safer atomic conversions, and other functions for safer coding;

see [tinycodet\\_safer](#).

(2) A new package import system, that combines the benefits of aliasing a package with the benefits of attaching a package;

see [tinycodet\\_import](#)

(3) Extending the string manipulation capabilities of the stringi R package;

see [tinycodet\\_strings](#).

(4) Reducing repetitive code;

see [tinycodet\\_dry](#).

And some miscellaneous functionality; see [tinycodet\\_misc](#).

The tinycodet R-package has only one dependency, namely `stringi`. No other dependencies, thus avoiding "dependency hell". 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/tinycodet>

## Usage

`tinycodet_help()`

## References

The badges shown in the documentation of this R package were made using the services of: <https://shields.io/>

---

<code>tinycodet_import</code>	<i>The tinycodet import system</i>
-------------------------------	------------------------------------

---

## Description

The tinycodet R package introduces a new package import system.

One can use a package **without attaching** the package - for example by using the `::` operator.

Or, one can explicitly **attach** a package - for example by using the `library` function.

The advantages and disadvantages of using **without attaching** a package versus **attaching** a package, at least those relevant here, are compactly presented in the following list:

(1) Prevent masking functions from other packages:

`use without attach`: Yes(advantage); `attaching`: No(disadvantage);

(2) Prevent masking core R functions:

`use without attach`: Yes(advantage); `attaching`: No(disadvantage);

(3) Clarify which function came from which package:

`use without attach`: Yes(advantage); `attaching`: No(disadvantage);

(4) Place/expose functions only in current environment instead of globally:

`use without attach`: Yes(advantage); `attaching`: No(disadvantage);

(5) Prevent namespace pollution:

`use without attach`: Yes(advantage); `attaching`: No(disadvantage);

(6) Minimize typing - especially for infix operators

(i.e. typing `package::`%op%`(x, y)` instead of `x %op% y` is cumbersome):

`use without attach`: No(disadvantage); `attaching`: Yes(advantage);

(7) Use multiple related packages, without constantly switching between package prefixes

(i.e. doing `packagename1::some_function1()`;

`packagename2::some_function2()`;

`packagename3::some_function3()` is chaotic and cumbersome):

[use without attach](#): No(disadvantage); [attaching](#): Yes(advantage);

What tinycodet 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](#): Load a main package + its re-exports + its dependencies + its extensions under a single alias. This essentially combines the attaching advantage of using multiple related packages (item 7 on the list), whilst keeping most advantages of using without attaching 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 (item 6 on the list), whilst simultaneously avoiding the disadvantage of attaching functions from a package globally (item 4 on the list).
- [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](#).

All `import_`-functions have the `lib.loc` argument to specify the library path to load packages from, thus allowing straight-forward project isolation.

See the examples section below to get an idea how the tinycodet import system works in practice.

## Usage

```
tinycodet_import()
```

## See Also

[tinycodet\\_help](#)

## Examples

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

# exposing infix operators from "magrittr" 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)
```



---

tinycodet_misc	<i>The tinycodet miscellaneous functionality</i>
----------------	--

---

**Description**

Some additional functions provided by the tinycodet 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**

```
tinycodet_misc()
```

**See Also**

[tinycodet\\_help\(\)](#)

---

tinycodet_safer	<i>The tinycodet "safer" functionality</i>
-----------------	--

---

**Description**

To help make your code safer, the tinycodet 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**

```
tinycodet_safer()
```

**See Also**

[tinycodet\\_help\(\)](#)

**Examples**

```
x <- c(0.3, 0.6, 0.7)
y <- c(0.1*3, 0.1*6, 0.1*7)
x == y # gives FALSE, but should be TRUE
x %d==% y # here it's done correctly
```

---

tinycodet_strings	<i>The tinycodet extension of 'stringi'</i>
-------------------	---

---

## Description

The tinycodet 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 [detecting patterns](#).
- The tinycodet 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.
- Although the functions are written in R, they have been optimized to be in the same order of speed as the other stringi functions.

## Usage

```
tinycodet_strings()
```

## References

Gagolewski M., *stringi: Fast and portable character string processing in R*, Journal of Statistical Software 103(2), 2022, 1–59, doi:10.18637/jss.v103.i02

## See Also

[tinycodet\\_help\(\)](#), [stri\\_rgx\(\)](#)

## Examples

```
# character vector:
x <- c("3rd 1st 2nd", "5th 4th 6th")
print(x)

# detect if there are digits:
x %s{}% "[[:digits]]"

# cut x into matrix of individual words:
x <- strcut_brk(x, "word")

# re-order matrix using the fast %row~% operator:
```

```
mat <- stringi::stri_rank(as.vector(x)) |> matrix(ncol=ncol(x))
sorted <- x %row% mat

# join elements of every row into a single character vector:
stri_c_mat(sorted, margin=1, sep=" ")
```

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**

<code>x</code>	a vector, matrix, or array.
<code>cond</code>	either an object of class <code>logical</code> with the same length as <code>x</code> , or a (possibly anonymous) function that returns an object of class <code>logical</code> with the same length as <code>x</code> . For example: <code>\(x)x&gt;0</code> .
<code>yes</code>	the (possibly anonymous) transformation function to use when function <code>cond(x)==TRUE</code> / logical <code>cond==TRUE</code> . Alternatively, one can also supply an atomic scalar. If argument <code>yes</code> is not specified, it defaults to <code>\(x)x</code> .
<code>no</code>	the (possibly anonymous) transformation function to use when function <code>cond(x)==FALSE</code> / logical <code>cond==FALSE</code> . Alternatively, one can also supply an atomic scalar. If argument <code>no</code> is not specified, it defaults to <code>\(x)x</code> .
<code>other</code>	the (possibly anonymous) transformation function to use when function <code>cond(x)</code> / logical <code>cond</code> returns <code>NA</code> . Alternatively, one can also supply an atomic scalar. If argument <code>other</code> is not specified, it defaults to <code>NA</code> . Note that function <code>other(x)</code> is run or scalar <code>other</code> is returned when function <code>cond(x)</code> or logical <code>cond</code> is <code>NA</code> , not necessarily when <code>x</code> itself is <code>NA</code> .

## 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

[tinycodet\\_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 tinycodet package import system*

---

## Description

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

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](#), [import\\_inops](#), or [import\\_LL](#) function.

The `attr.import()` function gets one specific special attributes or all special attributes from an alias object returned by [import\\_as](#).

**Usage**

```
help.import(..., i, alias)

is.tinyimport(x)

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

**Arguments**

...	further arguments to be passed to <a href="#">help</a> .
i	either one of the following: <ul style="list-style-type: none"> <li>• a function (use back-ticks when the function is an infix operator). Examples: <code>myfun</code>, <code>`%operator%`</code>, <code>myalias.\$some_function</code>. If a function, the <code>alias</code> argument is ignored.</li> <li>• a string giving the function name or topic (i.e. <code>"myfun"</code>, <code>"thistopic"</code>). If a string, argument <code>alias</code> must be specified also.</li> </ul>
alias	the alias object as created by the <a href="#">import_as</a> function.
x	the object/function to be tested.
which	The attributes to list. If <code>NULL</code> , all attributes will be returned. Possibilities: <code>"pkgs"</code> , <code>"conflicts"</code> , <code>"versions"</code> , <code>"args"</code> , and <code>"ordered_object_names"</code> .

**Details**

For `help.import(...)`:  
Do not use the `topic` / `package` and `i` / `alias` argument sets together. It's either one set or the other.

**Value**

For `help.import()`:  
Opens the appropriate help page.

For `is.tinyimport()`:  
Returns `TRUE` if the function is produced by [import\\_as](#), [import\\_inops](#), or [import\\_LL](#), 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.

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.

For conflicting objects, the last load is used for the ordering.

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

[tinycodet\\_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)
```

**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**

<code>x</code>	a string or character vector.
<code>p</code>	either a list with <code>stringi</code> arguments (see <a href="#">stri_rgx</a> ), or else a character vector of the same length as <code>x</code> with regular expressions. <a href="#">about search: regex</a> <a href="#">about search: fixed</a> <a href="#">about search: coll</a> <a href="#">about search: charclass</a>

**Value**

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

**See Also**

[tinycodet\\_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[x %s{}% "a"] <- 1
x[x %s!{}% "a"] <- 1
```

```

print(x)

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

#####

# ignore case pattern ====

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- list(regex = c("A", "A"), case_insensitive=TRUE)
x %s{}% p
x %s!{}% p
which(x %s{}% p)
which(x %s!{}% p)
x[x %s{}% p]
x[x %s!{}% p]
x[x %s{}% p] <- "hello"
x[x %s!{}% p] <- "hello"
print(x)

#####

# multi-character pattern ====

x <- c(paste0(letters[1:13], collapse=""), paste0(letters[14:26], collapse=""))
print(x)
p <- list(regex = rep("AB", 2), case_insensitive=TRUE)
x %s{}% p
x %s!{}% p
which(x %s{}% p)
which(x %s!{}% p)
x[x %s{}% p]
x[x %s!{}% p]
x[x %s{}% p] <- "CD"
x[x %s!{}% p] <- "CD"
print(x)

```



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