

# Package ‘tinycodet’

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**Title** Functions to Help in your Coding Etiquette

**Version** 0.5.3

**Description** Adds some functions to help in your coding etiquette.

‘tinycodet’ primarily focuses on 4 aspects.

- 1) Safer decimal (in)equality testing,  
standard-evaluated alternatives to with() and aes(),  
and other functions for safer coding.
- 2) A new package import system,  
that attempts to combine the benefits of using a package without attaching it,  
with the benefits of attaching a package.
- 3) Extending the string manipulation capabilities of the ‘stringi’ R package.
- 4) Reducing repetitive code.

Besides linking to ‘Rcpp’, ‘tinycodet’ has only one other dependency, namely ‘stringi’.

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aaa0_tinycodet_help	<i>tinycodet: Functions to Help in your Coding Etiquette</i>
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## Description

Welcome to the 'tinycodet' introduction help page!

'tinycodet' adds some functions to help in your coding etiquette.

It primarily focuses on 4 aspects:

- (1) Safer decimal (in)equality testing, standard-evaluated alternatives to with() and aes(), and other functions for safer coding;  
see [tinycodet\\_safer](#).

(2) A new package import system, that attempts to combine the benefits of using a package without attaching it, 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](#).

Please check the Change-log (see links below) regularly for updates (such as bug fixes).

'tinycodet' adheres to the [tinyverse](#) philosophy. Besides linking to 'Rcpp', 'tinycodet' only has one other dependency: 'stringi'. No other dependencies, thus avoiding "dependency hell". Most functions in this R-package are vectorized and optimised.

## Author(s)

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## References

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

## See Also

Useful links:

- 'tinycodet' GitHub main page and Read-Me: <https://github.com/tony-aw/tinycodet/>
- 'tinycodet' package website: <https://tony-aw.github.io/tinycodet/>
- Report bugs at: <https://github.com/tony-aw/tinycodet/issues/>
- Changelog: <https://github.com/tony-aw/tinycodet/blob/main/NEWS.md/> or <https://tony-aw.github.io/tinycodet/news/index.html>
- The 'fastverse', which is related to the 'tinyverse': <https://github.com/fastverse/fastverse/>

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aaa1\_tinycodet\_safer    *Overview of the 'tinycodet' "Safer" Functionality*

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## Description

To help make your code safer, the 'tinycodet' R-package introduces a few functions:

- [Safer decimal \(in\)equality testing](#).
- The [lock\\_TF](#) function to set and lock T and F to TRUE and FALSE, respectively.

- The `%<-c%` operator to assign locked constants.
- Standard evaluated versions of some common expression-evaluation functions: `with_pro` and `aes_pro`.
- `safer_partialmatch` to set options for safer dollar, arguments, and attribute matching.

## 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
```

---

aaa2\_tinycodet\_import *Overview of the 'tinycodet' Import System*

---

## 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) Enable functions only in current/local 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) Minimise 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 `package1::some_function1()`;  
`package2::some_function2()`;  
`package3::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**: Import a main package, and optionally its re-exports + its direct dependencies + its direct 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**.

The import system also includes general helper functions:

- The **x.import** functions: helper functions specifically for the 'tinycodet' import system.
- The **pversion\_** functions: check mismatch between loaded package version and package version in library path.
- The **pkgs** - functions: general helper functions regarding packages.

See the examples section below to get an idea of how the 'tinycodet' import system works in practice. More examples can be found on the website (<https://tony-aw.github.io/tinycodet/>)

## Details

### When to Use or Not to Use the 'tinycodet' Import System

The 'tinycodet' import system is helpful particularly for packages that have at least one of the following properties:

- The namespace of the package(s) conflicts with other packages.
- The namespace of the package(s) conflicts with core R, or with those of recommended R packages.
- The package(s) have function names that are generic enough, such that it is not obvious which function came from which package.

See examples below.

There is no necessity for using the 'tinycodet' import system with every single package. One can safely attach the 'stringi' package, for example, as 'stringi' uses a unique and immediately recognisable naming scheme (virtually all 'stringi' functions start with "stri\_"), and this naming scheme does not conflict with core R, nor with most other packages.

Of course, if one wishes to use a package (like 'stringi') **only** within a specific environment, it becomes advantageous to still import the package using the 'tinycodet' import system. In that case the [import\\_LL](#) function would be most applicable.

### Some Additional Comments on the 'tinycodet' Import System

- (S3) Methods will automatically be registered.
- Pronouns, such as the .data and .env pronouns from the 'rlang' package, will work without any prefixes required.
- All functions imported by the [import\\_as](#), [import\\_inops](#), or [import\\_LL](#) functions have a "package" attribute, so you will always know which function came from which package.

### See Also

[tinycodet\\_help](#)

### Examples

```
all(c("dplyr", "powerjoin", "magrittr") %installed in% .libPaths())

# NO packages are being attached in any of the following code

# import 'dplyr' + its re-exports + extension 'powerjoin', under alias "dpr.":
import_as(
  ~ dpr., "dplyr", re_exports = TRUE, extensions = "powerjoin"
)

# 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 %>% dpr.$filter(species == "Droid") %>%
  dpr.$select(name, dpr.$ends_with("color"))

male_penguins <- dpr.$tribble(
  ~name, ~species, ~island, ~flipper_length_mm, ~body_mass_g,
  "Giordan", "Gentoo", "Biscoe", 222L, 5250L,
  "Lynden", "Adelie", "Torgersen", 190L, 3900L,
  "Reiner", "Adelie", "Dream", 185L, 3650L
)

female_penguins <- dpr.$tribble(
  ~name, ~species, ~island, ~flipper_length_mm, ~body_mass_g,
  "Alonda", "Gentoo", "Biscoe", 211, 4500L,
  "Ola", "Adelie", "Dream", 190, 3600L,
  "Mishayla", "Gentoo", "Biscoe", 215, 4750L,
)
dpr.$check_specs()
```

```
dpr.$power_inner_join(
  male_penguins[c("species", "island")],
  female_penguins[c("species", "island")]
)

mypaste <- function(x, y) {
  import_LL("stringi", selection = "stri_c")
  stringi::stri_c(x, y)
}
mypaste("hello ", "world")
```

---

aaa3\_tinycodet\_strings

*Overview of the 'tinycodet' Extension of 'stringi'*


---

## Description

Virtually every programming language, even those primarily focused on mathematics, will at some point have to deal with strings. R's atomic classes boil down to some form of either numbers or characters. R's numerical functions are generally very fast. But R's native string functions are somewhat slow, do not have a unified naming scheme, and are not as comprehensive as R's impressive numerical functions. The primary R-package that fixes this is 'stringi'.

'stringi' is arguably the fastest and most comprehensive string manipulation package available at the time of writing. Many string related packages fully depend on 'stringi' (see its reverse-dependencies on CRAN).

As string manipulation is so important to programming languages, 'tinycodet' adds a little bit new functionality to 'stringi'.

'tinycodet' adds the following functions to extend 'stringi':

- Find  $i^{th}$  pattern occurrence ([stri\\_locate\\_ith](#)), or  $i^{th}$  text boundary ([stri\\_locate\\_ith\\_boundaries](#)).
- [Concatenate a character matrix row- or column-wise](#).

'tinycodet' adds the following operators, to complement the already existing 'stringi' operators:

- Infix operators for [string arithmetic](#).
- Infix operators for [string sub-setting](#), which get or remove the first and/or last n characters from strings.
- Infix operators for [detecting patterns](#), and [strfind\(\)](#)<- for locating/extracting/replacing found patterns.

And finally, 'tinycodet' adds the somewhat separate [strcut\\_-functions](#), to cut strings into pieces without removing the delimiters.

### Regarding Vector Recycling in the 'stringi'-based Functions

Generally speaking, vector recycling is supported as 'stringi' itself supports it also.

There are, however, a few exceptions.

First, matrix inputs (like in `strcut_loc` and [string sub-setting operators](#)) will generally not be recycled.

Second, the `i` argument in [stri\\_locate\\_ith](#) does not support vector recycling.

Scalar recycling is virtually always supported.

### 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](https://doi.org/10.18637/jss.v103.i02)

### See Also

[tinycodet\\_help](#), [s\\_pattern](#)

### Examples

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

# detect if there are digits:
x %s{}% "\\d"

# find second last digit:
loc <- stri_locate_ith(x, i = -2, regex = "\\d")
stringi::stri_sub(x, from = loc)

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

# sort rows of matrix using the fast %row~% operator:
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol = ncol(mat))
sorted <- mat %row~% rank
sorted[is.na(sorted)] <- ""

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



## Description

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

- The [transform\\_if](#) function
- [Atomic type casting with names and dimensions preserved](#).
- The [subset\\_if](#) operators and the [in-place unreal modifier operator](#).
- The [general in-place \(mathematical\) modification operator](#).

## 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
```

## 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.

**See Also**[tinycodet\\_help\(\)](#)

atomic\_typecast

*Atomic Type Casting With Names and Dimensions Preserved***Description**

Atomic type casting in R is generally performed using the functions [as.logical](#), [as.integer](#), [as.double](#), [as.character](#), [as.complex](#), and [as.raw](#).

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

The functions provided here by the 'tinycodet' package preserve the dimensions, dimnames, and names.

The functions are as follows:

- `as_bool()`: converts object to atomic type logical (TRUE, FALSE, NA).
- `as_int()`: converts object to atomic type integer.
- `as_dbl()`: converts object to atomic type double (AKA decimal numbers).
- `as_chr()`: converts object to atomic type character.
- `as_cplx()`: converts object to atomic type complex.
- `as_raw()`: converts object to atomic type raw.

**Usage**`as_bool(x, ...)``as_int(x, ...)``as_dbl(x, ...)``as_chr(x, ...)``as_cplx(x, ...)``as_raw(x, ...)`**Arguments**

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

**Value**

The converted object.

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

```
# matrix example ====
x <- matrix(rnorm(30), ncol = 5)
colnames(x) <- month.name[1:5]
rownames(x) <- month.abb[1:6]
names(x) <- c(letters[1:20], LETTERS[1:10])
print(x)

as_bool(x)
as_int(x)
as_dbl(x)
as_chr(x)
as_cplx(x)
as_raw(x)

#####

# factor example ====
x <- factor(month.abb, levels = month.abb)
names(x) <- month.name
print(x)

as_bool(as_int(x) > 6)
as_int(x)
as_dbl(x)
as_chr(x)
as_cplx(x)
as_raw(x)
```

decimal\_truth

*Safer Decimal Number (In)Equality Testing Operators***Description**

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

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

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

except for 2 aspects:

1. The decimal number (in)equality operators assume that if the absolute difference between any 2 numbers x and y is smaller than the Machine tolerance, `sqrt(.Machine$double.eps)`, then x and y should be consider to be equal.

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.

- Only numeric input is allowed, so characters are not coerced to numbers.  
I.e. `1 < "a"` gives TRUE, whereas `1 %d<% "a"` gives an error.  
For character equality testing, see `%s==%` from the 'stringi' package.

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

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`.

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

### 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

is_wholenumber(x, tol = sqrt(.Machine$double.eps))
```

### Arguments

<code>x, y</code>	numeric vectors, matrices, or arrays.
<code>bnd</code>	either a vector of length 2, or a matrix with 2 columns and 1 row, or else a matrix with 2 columns where <code>nrow(bnd)==length(x)</code> (or can be recycled to be <code>nrow(bnd)==length(x)</code> ). The first element/column of <code>bnd</code> gives the lower bound of the closed interval; The second element/column of <code>bnd</code> gives the upper bound of the closed interval.
<code>tol</code>	a single, strictly positive number close to zero, giving the tolerance.

### Value

For the `%d...%` operators:

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

NOTE: `Inf` by `Inf` and `-Inf` by `-Inf` comparisons with the `%d...%` operators return NA.

For `is_wholenumber()`:

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

NOTE: `Inf`, `-Inf`, `NaN` and `NA` all return `NA` for `is_wholenumber()`.

## 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 work for integers also:
x <- 1L:5L
y <- 1L:5L
x %d==% y
x %d!=% y
x %d<% y
x %d>% y
x %d<=% y
x %d>=% y
#'

x <- 1L:5L
y <- x+1
x %d==% y
x %d!=% y
x %d<% y
x %d>% y
x %d<=% y
x %d>=% y
#'

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

```
# is_wholenumber:
is_wholenumber(1:10 + c(0, 0.1))
```

---

import_as	<i>Import R-package, its Re-exports, Dependencies, and/or Extensions, Under a Single 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, containing the exported functions from the specified packages, will be placed in the current environment.

## Usage

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

## Arguments

- |              |  |
|--------------|--|
| alias        | a syntactically valid non-hidden name giving the alias object where the package(s) are to be imported into.<br>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.).   |
| main_package | a single string, giving the name of the main package to import under the given alias.<br>Core R (i.e. "base", "stats", etc.) is not allowed.   |
| re_exports   | TRUE or FALSE. <ul style="list-style-type: none"> <li>If <code>re_exports = TRUE</code> the re-exports from the <code>main_package</code> (including those exported from Core R) are added to the alias together with the main package.<br/>This is the default, as it is analogous to the behaviour of base R's <code>::</code> operator.</li> <li>If <code>re_exports = FALSE</code>, these re-exports are not added together with the main package.<br/>The user can still import the packages under the alias from which the re-exported functions came from, by specifying them in the <code>dependencies</code> argument.</li> </ul> |
| dependencies | an optional character vector, giving the names of the dependencies of the <code>main_package</code> to be imported also under the alias.<br>Defaults to <code>NULL</code> , which means no dependencies are imported under the alias.<br>See <a href="#">pkg_get_deps</a> to quickly get dependencies from a package.<br>Core R (i.e. "base", "stats", etc.) is not allowed.   |

extensions	an optional character vector, giving the names of the extensions of the main_package to be imported also under the alias. Defaults to NULL, which means no extensions are imported under the alias. Core R (i.e. "base", "stats", etc.) is not allowed.
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> .
import_order	the character vector c("dependencies", "main_package", "extensions"), or some re-ordering of this character vector, giving the relative import order of the groups of packages. See Details section for more information.

## Details

### Expanded Definitions of Some Arguments

- "Re-exports" are functions that are defined in the dependencies of the main\_package, but are re-exported in the namespace of the main\_package.  
Unlike the Dependencies argument, functions from core R are included in re-exports.
- "Dependencies" are here defined as any R-package appearing in the "Depends", "Imports", or "LinkingTo" fields of the Description file of the main\_package. So no recursive dependencies.
- "Extensions" are reverse-dependencies that actually extend the functionality of the main\_package. Programmatically, some package "E" is considered an extension of some "main\_package", if the following is TRUE:  
"main\_package" %in% `pkg_get_deps_minimal("E")`

### Why Aliasing Multiple Packages is Useful

To use an R-package with its extension packages or dependencies, whilst avoiding the disadvantages of attaching a package (see [tinycodet\\_import](#)), one would traditionally use the `::` operator like so:

```
main_package::some_function1()
dependency1::some_function2()
extension1::some_function3()
```

This becomes cumbersome as more packages are needed and/or as the package name(s) become longer.

The `import_as()` function avoids this issue by allowing multiple **related** packages to be imported under a single alias, allowing one to code like this:

```
import_as(
  ~ alias., "main_package",
  dependencies = "dependency1", extensions = "extension1",
```

```

    lib.loc = .libPaths()
  )
  alias.$some_function1()
  alias.$some_function2()
  alias.$some_function3()

```

Thus importing a package, or multiple directly related packages, under a single alias, which `import_as()` provides, avoids the above issues. Importing a package under an alias is referred to as "aliasing" a package.

### Alias Naming Recommendation

To keep package alias object names 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 (\_).

### Regarding `import_order`

The order of the character vector given in the `dependencies` and `extensions` arguments matters. If multiple packages share objects with the same name, the objects of the package named last will overwrite those of the earlier named packages.

The `import_order` argument defaults to the character vector `c("dependencies", "main_package", "extensions")`, which is the recommended setting.

This setting results in the following importing 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 extensions, **in the order specified by the `extensions` argument**.

### Other Details

Packages that appear in the "Suggests" or "Enhances" fields of packages are not considered dependencies or extensions.

No more than 10 packages (ignoring re-exports) are allowed to be imported under a single alias.

### Value

A locked environment object, similar to the output of [loadNamespace](#), with the name as specified in the `alias` argument, will be created.

This object, referred to as the "(package) alias object", will contain the exported functions from the specified package(s).

The alias object will be placed in the current environment.

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 "unimport" the package alias object, simply remove it (i.e. `rm(list = "alias.")`).

## See Also

[tinycodet\\_import](#)

## Examples

```
all(c("data.table", "tidytable") %installed in% .libPaths())

import_as( # this creates the 'tdt.' object
  "tdt.", "tidytable", dependencies = "data.table"
)
# same as:
import_as(
  ~ tdt., "tidytable", dependencies = "data.table"
)
```

---

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

<code>package</code>	a single string, giving the name of the R-package.
<code>dataname</code>	a single string, giving the name of the data set.
<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> .

## Value

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

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

```
d <- import_data("datasets", "cars")
head(d)
```

---

import_inops	<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.

`import_inops(unexpose = ...)` "unexposes" (i.e. removes) the infix operators specified in a package or an alias object from the current environment.

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-defined) infix operators will not be touched.

To attach all infix operators from a package to the global namespace, one can use the [pkg\\_lsf](#) function like so:

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

**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. Core R (i.e. "base", "stats", etc.) is not allowed.

`lib.loc`

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

Only used when supplying a string to `expose` / `unexpose`, and ignored when supplying an alias object to `expose` / `unexpose` (the library is path already stored 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

### Why Exposing Infix Operators Is Useful

To use a function from an R-package, while avoiding the disadvantages of attaching a package (see [tinycodet\\_import](#)), one would traditionally use the `::` operator like so:

```
packagename::function_name()
```

This is, however, cumbersome with infix operators, as it forces one to code like this:

```
packagename::`%op%`(x,y)
```

Exposing infix operators to the current environment, using the `import_inops()` function, allows one to use infix operators without using cumbersome code, and without having to attach the infix operators globally.

### Other 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.

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.

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

## See Also

[tinycodet\\_import](#), `import_inops.control()`, `report_inops()`

## Examples

```
import_inops(expose = "stringi") # expose infix operators from package
import_inops(unexpose = "stringi") # remove the exposed infix operators from environment
```

```
import_as(~ stri., "stringi")
import_inops(expose = stri.) # expose infix operators from alias
import_inops(unexpose = stri.) # unexposed infix operators from current environment
```

```
# additional arguments (only used when exposing, not unexposing):
import_inops(expose = "stringi", exclude = "%s==%")
```

```

import_inops(unexpose = "stringi")
import_inops(expose = "stringi", overwrite = FALSE)
import_inops(unexpose = "stringi")

import_as(~ stri., "stringi")
import_inops(expose = stri., include.only = "%s==%")
import_inops(unexpose = stri.)
import_inops(expose = stri., overwrite = FALSE)
import_inops(unexpose = stri.)

```

---

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      | a character vector, giving the infix operators NOT to expose to the current environment.<br>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.<br>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. <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 continues 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`      logical.

When exposing infix operators, [import\\_inops](#) checks if infix operators with the same names are already present in the current environment.

If `inherits = FALSE`, only the current environment is checked for existing operators.

If `inherits = TRUE`, enclosed environments, most notably package namespaces, are also checked for existing operators.

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

## Examples

```
# additional arguments (only used when exposing, not unexposing):
import_as(~ stri., "stringi")
import_inops(expose = stri., include.only = "%s==%")
import_inops(unexpose = stri.)
import_inops(expose = "stringi", exclude = "%s==%")
import_inops(unexpose = "stringi")
import_inops(expose = stri., overwrite = FALSE)
import_inops(unexpose = stri.)
import_inops(expose = "stringi", overwrite = FALSE)
import_inops(unexpose = "stringi")
```

## 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 exposing functions inside a local environment.

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

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

1. `import_int()` includes the `lib.loc` argument.
2. `import_int()` 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. Core R (i.e. "base", "stats", etc.) is not allowed.
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 <code>lib.loc</code> argument would usually be <code>.libPaths()</code> . 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: <code>package_name ~ function_name</code> Core R (i.e. "base", "stats", etc.) is not allowed.

## 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, and locked.

To unexpose 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**

```
# Using import_LL ====
import_LL(
  "stringi", "stri_sub"
)
# the stri_sub() function now cannot be modified, only used or removed, because it's locked:
bindingIsLocked("stri_sub", environment()) # TRUE

mypaste <- function(x, y) {
  import_LL("stringi", selection = "stri_c")
  stri_c(x, y)
}
mypaste("hello ", "world")

# 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")
```

## Description

The `x %:=% f` operator performs 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:

```
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

*Lock T, Lock F, or Create Locked Constants***Description**

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(env)
```

```
X %<-c% A
```

**Arguments**

env	an optional environment to give, determining in which environment T and F should be locked. When not specified, the current environment is used.
X	a syntactically valid unquoted name of the object to be created.
A	any kind of object to be assigned to X.

**Details**

Note that following statement

```
x %<-c% 2+2
print(x)
```

returns

```
[1] 2
```

due to R's precedence rules. Therefore, in such cases, the right hand side of `X %<-c% A` need to be surrounded with brackets. I.e.:

```
x %<-c% (2 + 2)
```

Note that the `lock_TF()` function and `%s<-c%` operator create constants through [lockBinding](#).

The constants are protected from modification by copy, but they are **not** protected from modification by reference (see for example `collapse::setv`).

**Value**

For `lock_TF()`:

Two constants, namely T and F, set to TRUE and FALSE respectively, are created in the specified or else 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

*Additional 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 checks for every value of numeric vector n if it can be considered a number belonging to type numtype.

The `s %=strtype% strtype` operator checks for every value of character vector s if it can be seen as a certain strtype.

**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. See Details section for supported types.
s	a character vector.
strtype	a single string giving the string type to be checked. See Details section for supported types.

### Details

For argument numtype, the following options are supported:

- "~0": zero, or else a number whose absolute value is smaller than the Machine tolerance ( $\sqrt{.Machine\$double.eps}$ ).
- "B": binary numbers (exactly 0 or exactly 1);
- "prop": proportions - numbers between 0 and 1 (exactly 0 or 1 is also allowed);
- "I": Integers;
- "odd": odd integers;
- "even": even integers;
- "R": Real numbers;
- "unreal": infinity, NA, or NaN;

For argument strtype, 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.

## 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

*Row- or 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**

<code>x</code>	a matrix
<code>mat</code>	a matrix with the same dimensions as <code>x</code> , giving the ordering ranks of every element of matrix <code>x</code> .

**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(seq_along(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.

Note that these operators strip all attributes except dimensions.

**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(seq_along(x)) |> matrix(ncol = ncol(x))
x %row~% mat # randomly shuffle every row independently
x %col~% mat # randomly 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(seq_along(x)) |> matrix(ncol = ncol(x))
x %row~% mat # randomly shuffle every row independently
x %col~% mat # randomise shuffle every column independently

```

---

pkgs

---

*Miscellaneous Package Related Functions*


---

## Description

The `pkgs %installed in% lib.loc` operator checks if one or more given packages (`pkgs`) exist in the given library paths (`lib.loc`), without loading the packages.

The syntax of this operator forces the user to make it syntactically explicit where to look for installed R-packages.

As `pkgs %installed in% lib.loc` does not even load a package, the user can safely use it without fearing any unwanted side-effects.

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

The `pkg_get_deps_minimal()` function is the same as `pkg_get_deps()`, except with `base`, `recom`, `rstudioapi`, `shared_tidy` all set to `FALSE`, and the default value for `deps_type` is `c("Depends", "Imports")`.

The `pkg_ls()` 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 package using `library`, like so:

```

y <- pkg_ls("packagename", type = "inops")
library(packagename, include.only = y)

```

## Usage

```
pkgs %installed in% lib.loc
```

```
pkg_get_deps(
```

```

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

pkg_get_deps_minimal(
  package,
  lib.loc = .libPaths(),
  deps_type = c("Depends", "Imports")
)

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

```

## Arguments

pkgs	a character vector with the package name(s).
lib.loc	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> .
package	a single string giving the package name.
deps_type	a character vector, giving the dependency types to be used. The order of the character vector given in <code>deps_type</code> affects the order of the returned character vector; see Details sections.
base	logical, indicating whether base/core R should be included (TRUE), or not included (FALSE).
recom	logical, indicating whether the pre-installed 'recommended' R-packages should be included (TRUE), or not included (FALSE).
rstudioapi	logical, indicating whether the 'rstudioapi' R-package should be included (TRUE), or not included (FALSE).
shared_tidy	logical, indicating whether the shared dependencies of the 'tidyverse' should be included (TRUE), or not included (FALSE). <b>Details:</b> Some of the (often many) dependencies 'tidyverse' packages have are shared across the majority of the 'tidyverse'. The "official" list of shared dependencies in the 'tidyverse' currently is the following: 'rlang', 'lifecycle', 'cli', 'glue', and 'withr'.
type	The type of functions to list. Possibilities: <ul style="list-style-type: none"> <li>"inops" or "operators": Only infix operators.</li> <li>"regfuns": Only regular functions (thus excluding infix operators).</li> <li>"all": All functions, both regular functions and infix operators.</li> </ul>

## Details

For `pkg_get_deps()`:

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:

For example, `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.

The names give the package names.

The value `TRUE` indicates a package is installed in `lib.loc`.

The value `FALSE` indicates a package is not installed in `lib.loc`.

The value `NA` indicates a package is not actually a separate package, but base/core 'R' (i.e. 'base', 'stats', etc.).

For `pkg_get_deps()` and `pkg_get_deps_minimal()`:

A character vector of direct dependencies, without duplicates.

For `pkg_ls()`:

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

## References

O'Brien J., 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-d>

## See Also

[tinycodet\\_import](#)

## Examples

```
"dplyr" %installed in% .libPaths()
```

```
pkg_get_deps_minimal("dplyr")
pkgs <- pkg_get_deps("dplyr")
pkgs %installed in% .libPaths()
pkg_ls("dplyr", "all")
```



---

pversion

*Check for Package Versions Mismatch*

---

### Description

The `pversion_check4mismatch()` function checks if there is any mismatch between the currently loaded packages and the packages in the specified library path.

The `pversion_report()` function gives a table of all specified packages, with their loaded and installed versions, regardless if there is a mismatch or not.

### Usage

```
pversion_check4mismatch(pkgs = NULL, lib.loc = .libPaths())
```

```
pversion_report(pkgs = NULL, lib.loc = .libPaths())
```

### Arguments

<code>pkgs</code>	a character vector with the package name(s). Packages that are not actually loaded will be ignored. Base/core R will also be ignored. If <code>NULL</code> , all loaded packages (see <a href="#">loadedNamespaces</a> ) excluding core/base R will be checked.
<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> .

### Value

For `pversion_check4mismatch()`:

If no mismatch between loaded versions and those in `lib.loc` were found, returns `NULL`.

Otherwise it returns a `data.frame`, with the loaded version and library version of the specified packages.

For `pversion_report()`:

Returns a `data.frame`, with the loaded version and library version of the specified packages, as well as a logical column indicating whether the two versions are equal (`TRUE`), or not equal (`FALSE`).

### See Also

[tinycodet\\_import](#)

**Examples**

```
"dplyr" %installed in% .libPaths()

import_as(~dpr., "dplyr")
pversion_check4mismatch()
pversion_report()
```

report\_inops

*Report Infix Operators***Description**

The `report_inops()` function returns a data.frame listing the infix operators defined in the current environment, 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 is used.

**Value**

A data.frame. The first column gives the infix operator names. The second column gives the package the operator came from, or NA if it did not come from a package.

**See Also**

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

**Examples**

```
report_inops()

`%paste%` <- function(x,y)paste0(x,y)

report_inops()

import_inops("stringi")

report_inops()
```

---

safer_partialmatch	<i>Set Safer Dollar, Arguments, and Attribute Matching</i>
--------------------	--

---

## Description

The `safer_partialmatch()` function simply calls the following:

```
options(  
  warnPartialMatchDollar = TRUE,  
  warnPartialMatchArgs = TRUE,  
  warnPartialMatchAttr = TRUE  
)
```

Thus it forces 'R' to give a warning when partial matching occurs when using the dollar (\$) operator, or when other forms of partial matching occurs.

The `safer_partialmatch()` function is intended for when running R interactively (see [interactive](#)).

## Usage

```
safer_partialmatch()
```

## Value

Sets the options. Returns nothing.

## See Also

[tinycodet\\_safer](#)

## Examples

```
interactive()  
  
safer_partialmatch()  
data(iris)  
head(iris)  
iris$Sepal.Length <- iris$Sepal.Length^2  
head(iris)
```

---

source_selection	<i>Source Specific Objects from Script</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> . <a href="#">about search: regex</a>
<code>fixed</code>	a character vector of fixed patterns (see <a href="#">about_search_fixed</a> ). 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, <code>mymethod.numeric()</code> and <code>mymethod.character()</code> from generic <code>mymethod()</code> , one could specify <code>fixed = "mymethod"</code> . <a href="#">about search: fixed</a>

## 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.

**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

*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`.

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\(..., tokens\\_only = FALSE\)](#)) cuts every string into individual text breaks (like character, word, line, or sentence boundaries).

## Usage

```
strcut_loc(str, loc)
```

```
strcut_brk(str, type = "character", tolist = FALSE, n = -1L, ...)
```

## 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>type</code>	either one of the following: <ul style="list-style-type: none"> <li>a single string giving the break iterator type (i.e. "character", "line_break", "sentence", "word", or a custom set of ICU break iteration rules).</li> <li>a list with break iteration options, like a list produced by <a href="#">stri_opts_brkiter</a>.</li> </ul> <p><b>about search: boundaries</b></p>
<code>tolist</code>	logical, indicating if <code>strcut_brk</code> should return a list (TRUE), or a matrix (FALSE, default).
<code>n</code>	see <a href="#">stri_split_boundaries</a> .
<code>...</code>	additional arguments to be passed to <a href="#">stri_split_boundaries</a> .

## Details

The `strcut_` functions provide a short and concise way to cut strings into pieces, without removing the delimiters, which is an operation that lies at the core of virtually all boundaries-operations in 'stringi'.

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 return a matrix by default.

## Value

For `strcut_loc()`:

A character matrix with `length(str)` rows and 3 columns, where for every row `i` it holds the following:

- the first column contains the sub-string **before** `loc[i,]`, or NA if `loc[i,]` contains NA;
- the second column contains the sub\_string at `loc[i,]`, or the uncut string if `loc[i,]` contains NA;
- the third and last column contains the sub-string **after** `loc[i,]`, or NA if `loc[i,]` contains NA.

For `strcut_brk(..., tolist = FALSE)`:

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

Empty places are filled with NA.

For `strcut_brk(..., tolist = TRUE)`:

A list with `length(str)` elements, where each element is a character vector containing the cut string.

## 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))
strcut_loc(x, c(5, NA))
strcut_loc(x, c(NA, 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)
strcut_brk(test, n = 1)
strcut_brk(test, "line", tolist = TRUE)
strcut_brk(test, "word", tolist = TRUE)
strcut_brk(test, "sentence", tolist = TRUE)

brk <- stringi::stri_opts_brkiter(
  type = "line"
)
strcut_brk(test, brk)
```

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 within 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 within 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](#).

**Value**

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

**See Also**

[tinycodet\\_strings](#)

**Examples**

```
#####

# Basic example

x <- matrix(letters[1:25], ncol = 5, byrow = TRUE)
print(x)
stri_join_mat(x, margin = 1)

x <- matrix(letters[1:25], ncol = 5, byrow = FALSE)
print(x)
stri_join_mat(x, margin = 2)
```



```
#####
# sorting characters in strings ====

x <- c(paste(sample(letters), collapse = ""),
      paste(sample(letters), collapse = ""))
print(x)
mat <- strcut_brk(x)
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank
sorted[is.na(sorted)] <- ""
print(sorted)
stri_join_mat(sorted, margin = 1)
stri_join_mat(sorted, margin = 2)

#####

# sorting words ====

x <- c("2nd 3rd 1st", "Goodbye everyone")
print(x)
mat <- strcut_brk(x, "word")
rank <- stringi::stri_rank(as.vector(mat)) |> matrix(ncol=ncol(mat))
sorted <- mat %row~% rank
sorted[is.na(sorted)] <- ""
stri_c_mat(sorted, margin = 1, sep = " ") # <- alias for stri_join_mat
stri_c_mat(sorted, margin = 2, sep = " ")

#####

# randomly shuffling 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 <- sample(1:length(mat)) |> matrix(ncol = ncol(mat))
sorted <- mat %row~% rank
sorted[is.na(sorted)] <- ""
stri_paste_mat(sorted, margin = 1) # <- another alias for stri_join_mat
stri_paste_mat(sorted, margin = 2)
```

---

stri\_locate\_ith

Locate  $i^{th}$  Pattern Occurrence or Text Boundary

---

## Description

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

The `stri_locate_ith_boundaries()` function locates the  $i^{th}$  text boundary (like character, word, line, or sentence boundaries).

## Usage

```
stri_locate_ith(str, i, ..., regex, fixed, coll, charclass)
stri_locate_ith_regex(str, pattern, i, ..., opts_regex = NULL)
stri_locate_ith_fixed(str, pattern, i, ..., opts_fixed = NULL)
stri_locate_ith_coll(str, pattern, i, ..., opts_collator = NULL)
stri_locate_ith_charclass(str, pattern, i, merge = TRUE, ...)
stri_locate_ith_boundaries(str, i, ..., opts_brkiter = NULL)
```

## Arguments

<code>str</code>	a string or character vector.
<code>i</code>	<p>an integer scalar, or an integer vector of appropriate length (vector recycling is not supported).</p> <p>Positive numbers count occurrences from the left/beginning of the strings. Negative numbers count occurrences from the right/end of the strings. I.e.:</p> <ul style="list-style-type: none"> <li>• <code>stri_locate_ith(str, i = 1, ...)</code> gives the position (range) of the first occurrence of a pattern.</li> <li>• <code>stri_locate_ith(str, i = -1, ...)</code> gives the position (range) of the last occurrence of a pattern.</li> <li>• <code>stri_locate_ith(str, i = 2, ...)</code> gives the position (range) of the second occurrence of a pattern.</li> <li>• <code>stri_locate_ith(str, i = -2, ...)</code> gives the position (range) of the second-last occurrence of a pattern.</li> </ul> <p>If <code>abs(i)</code> is larger than the number of pattern occurrences <code>n</code>, the first (if <code>i &lt; -n</code>) or last (if <code>i &gt; n</code>) instance will be given.</p> <p>For example: suppose a string has 3 instances of some pattern; then if <code>i &gt;= 3</code> the third instance will be located, and if <code>i &lt;= -3</code> the first instance will be located.</p>
<code>...</code>	<p>more arguments to be supplied to <a href="#">stri_locate_all</a> or <a href="#">stri_locate_all_boundaries</a>. Do not supply the arguments <code>omit_no_match</code> or <code>get_length</code>, as they are already specified internally. Supplying these arguments anyway will result in an error.</p>
<code>pattern, regex, fixed, coll, charclass</code>	<p>a character vector of search patterns, as in <a href="#">stri_locate_all</a>.</p> <p>about search: <code>regex</code>  about search: <code>fixed</code>  about search: <code>coll</code>  about search: <code>charclass</code></p>

opts_regex, opts_fixed, opts_collator, opts_brkiter	<p>named list used to tune up the selected search engine's settings.          see <a href="#">stri_opts_regex</a>, <a href="#">stri_opts_fixed</a>, <a href="#">stri_opts_collator</a>, and <a href="#">stri_opts_brkiter</a>.          NULL for the defaults.          about search: <a href="#">regex</a>          about search: <a href="#">fixed</a>          about search: <a href="#">coll</a>          about search: <a href="#">charclass</a>          about search: <a href="#">boundaries</a></p>
merge	<p>logical, indicating if charclass locations should be merged or not.  <b>Details:</b>          For the charclass pattern type, the <code>stri_locate_ith()</code> function gives the start and end of <b>consecutive</b> characters by default, just like <a href="#">stri_locate_all</a>.          To give the start and end positions of single characters, much like <a href="#">stri_locate_first</a> or <a href="#">stri_locate_last</a>, set <code>merge = FALSE</code>.</p>

## Details

The 'stringi' functions only support operations on the first, last, or all occurrences of a pattern. The `stri_locate_ith()` function allows locating the  $i^{th}$  occurrence of a pattern. This allows for several workflows for operating on the  $i^{th}$  pattern occurrence. See also the examples section.

### Extract $i^{th}$ Occurrence of a Pattern

For extracting the  $i^{th}$  pattern occurrence:

Locate the the  $i^{th}$  occurrence using `stri_locate_ith()`, and then extract it using, for example, [stri\\_sub](#).

### Replace/Transform $i^{th}$ Occurrence of a Pattern

For replacing/transforming the  $i^{th}$  pattern occurrence:

1. Locate the the  $i^{th}$  occurrence using `stri_locate_ith()`.
2. Extract the occurrence using [stri\\_sub](#).
3. Transform or replace the extracted sub-strings.
4. Return the transformed/replaced sub-string back, using again [stri\\_sub](#).

### Capture Groups of $i^{th}$ Occurrence of a Pattern

The `capture_groups` argument for `regex` is not supported within `stri_locate_ith()`.

To capture the groups of the  $i^{th}$  occurrences:

1. Use `stri_locate_ith()` to locate the  $i^{th}$  occurrences without group capture.
2. Extract the occurrence using [stri\\_sub](#).
3. Get the matched group capture on the extracted occurrences using [stri\\_match](#).

## 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.

If an empty string or empty pattern is supplied, a warning is given and a matrix with 0 rows is returned.

## Note

### Long Vectors

The `stri_locate_ith`-functions do not support long vectors (i.e. character vectors with more than  $2^{31} - 1$  strings).

### Performance

The performance of `stri_locate_ith()` is close to that of [stri\\_locate\\_all](#).

## See Also

[tinycodet\\_strings](#)

## Examples

```
#####

# practical example: transform regex pattern ====

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

# locate ith (second and second-last) vowel locations:
p <- rep("A|E|I|O|U", 2) # vowels
loc <- stri_locate_ith(x, c(2, -2), regex = p, case_insensitive = TRUE)
print(loc)

# extract ith vowels:
extr <- stringi::stri_sub(x, loc)
print(extr)

# transform & replace ith vowels with numbers:
repl <- chartr("aeiou", "12345", extr)
stringi::stri_sub(x, loc) <- repl

# result (notice ith vowels are now numbers):
print(x)

#####

# practical example: group-capture regex pattern ====

# input character:
# first group: c(breakfast=eggs, breakfast=bacon)
# second group: c(lunch=pizza, lunch=spaghetti)
x <- c('breakfast=eggs;lunch=pizza',
```

```

      'breakfast=bacon;lunch=spaghetti',
      'no food here') # no group here
print(x)

# locate ith=2nd group:
p <- '(\w+)=(\w+)'
loc <- stri_locate_ith(x, i = 2, regex = p)
print(loc)

# extract ith=2nd group:
extr <- stringi::stri_sub(x, loc)
print(extr)

# capture ith=2nd group:
stringi::stri_match(extr, regex = p)

#####

# practical example: replace words using boundaries ====

# input character vector:
x <- c("good morning and good night",
      "hello ladies and gentlemen")
print(x)

# report ith word locations:
loc <- stri_locate_ith_boundaries(x, c(-3, 3), type = "word")
print(loc)

# extract ith words:
extr <- stringi::stri_sub(x, from=loc)
print(extr)

# transform and replace words (notice ith words have inverted case):
tf <- chartr(extr, old = "a-zA-Z", new = "A-Za-z")
stringi::stri_sub(x, loc) <- tf

# result:
print(x)

#####

# find pattern ====

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)

#####

# 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)
```

```
#####
```

```
# Boundaries ====
```

```
test <- c(
```

```
paste0("The\u00a0above-mentioned    features are very useful. ",
       "Spam, spam, eggs, bacon, and spam. 123 456 789"),
       "good morning, good evening, and good night"
)
loc <- stri_locate_ith_boundaries(test, i = c(1, -1), type = "word")
stringi::stri_sub(test, from = loc)
```

str\_arithmetic

*String Arithmetic Operators***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 character/pattern defined in `p` occurs in each element of `x`.

The `x %s//% brk` operator counts how often the text boundary specified in list `brk` 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.

The `x %ss% p` operator splits the strings in `x` by a delimiter character/pattern defined in `p`, and removes `p` in the process.

For cutting strings by text boundaries, or around a location, see [strcut\\_brk](#) and [strcut\\_loc](#).

**Usage**

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

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

```
x %s//% brk
```

```
x %ss% p
```

**Arguments**

`x` a string or character vector.

`p` either a list with 'stringi' arguments (see [s\\_pattern](#)), or else a character vector with regular expressions.

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

brk                    a list with break iteration options, like a list produced by [stri\\_opts\\_brkiter](#).  
[about search: boundaries](#)

## Value

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

The %s/% and %s//% both return an integer vector of the same length as x.

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

The %ss% operator returns a list of the split strings - or, if simplify = TRUE / simplify = NA, returns a matrix of the split strings.

## 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 %ss% p # split x around vowels, removing the vowels in the process
x %ss% s_regex(p, simplify = NA) # same as above, but in matrix form
```

```
test <- c(
  paste0("The\u00a0above-mentioned    features are very useful. ",
    "Spam, spam, eggs, bacon, and spam. 123 456 789"),
  "good morning, good evening, and good night"
)
test %s//% list(type = "character")
```

```
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.

x <- c(paste(letters, collapse = ", "), paste(LETTERS, collapse = ", "))
print(x)
x %ss% " ", "
t(x %ss% s_fixed(" ", simplify = NA))
```

str\_search

*'stringi' Pattern Search Operators*

## Description

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

When supplying a list on the right hand side (see [s\\_pattern](#)), one can optionally include the list element `at = "start"` or `at = "end"`:

- Supplying `at = "start"` will check if the pattern appears at the start of a string (like [stri\\_startswith](#)).
- Supplying `at = "end"` will check if the pattern appears at the end of a string (like [stri\\_endswith](#)).

The `x %s!{%}% p` operator is the same as `x %s{%}% p`, except it checks for **absence** of the pattern, rather than presence.

For string (in)equality operators, see [%s==%](#) from the `'stringi'` package.

`strfind()` <- locates, extracts, or replaces found patterns.

It complements the other string-related operators, and uses the same [s\\_pattern](#) API.

It functions as follows:

- `strfind()` finds all pattern matches, and returns the extractions of the findings in a list, just like [stri\\_extract\\_all](#).
- `strfind(..., i = "all" )`, finds all pattern matches like [stri\\_locate\\_all](#).
- `strfind(..., i = i)`, where `i` is an integer vector, locates the  $i^{th}$  occurrence of a pattern, and reports the locations in a matrix, just like [stri\\_locate\\_ith](#).
- `strfind() <- value` finds pattern matches in variable `x`, replaces the pattern matches with the character vector specified in `value`, and assigns the transformed character vector back to `x`.

This is somewhat similar to [stri\\_replace](#), though the replacement is done in-place.

## Usage

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

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

```
strfind(x, p, ..., i, rt)
```

```
strfind(x, p, ..., i, rt) <- value
```

## Arguments

x	<p>a string or character vector.</p> <p>For <code>strfind()</code>&lt;-, x must obviously be the variable containing the character vector/string, since <code>strfind()</code>&lt;- performs assignment in-place.</p>
p	<p>either a list with 'stringi' arguments (see <a href="#">s_pattern</a>), or else a character vector with regular expressions.</p> <p>See also the Details section.</p> <p>about search: <a href="#">regex</a></p> <p>about search: <a href="#">fixed</a></p> <p>about search: <a href="#">coll</a></p> <p>about search: <a href="#">charclass</a></p>
...	additional arguments to be specified.
i	<p>either one of the following can be given for i:</p> <ul style="list-style-type: none"> <li>• if i is not given or NULL, <code>strfind()</code> extracts all found pattern occurrences.</li> <li>• if i is the string "all", <code>strfind()</code> locates all found pattern occurrences.</li> <li>• if i is an integer, <code>strfind()</code> locates the <math>i^{th}</math> pattern occurrences.</li> </ul> <p>See the i argument in <a href="#">stri_locate_ith</a> for details.</p> <p>For <code>strfind()</code> &lt;- value, i must not be specified.</p>
rt	<p>use rt to specify the Replacement Type that <code>strfind()</code>&lt;- should perform. Either one of the following can be given for rt:</p> <ul style="list-style-type: none"> <li>• if rt is not given, NULL or "vec", <code>strfind()</code>&lt;- performs regular, vectorized replacement of <b>all</b> occurrences.</li> <li>• if rt = "dict", <code>strfind()</code>&lt;- performs dictionary replacement of <b>all</b> occurrences.</li> <li>• if rt = "first", <code>strfind()</code>&lt;- replaces only the first occurrences.</li> <li>• if rt = "last", <code>strfind()</code>&lt;- replaces only the last occurrences.</li> </ul> <p>Note: rt = "first" and rt = "last" only exist for convenience; for more specific locational replacement, use <a href="#">stri_locate_ith</a> or <code>strfind(..., i)</code> with numeric i (see the Examples section).</p> <p>For <code>strfind()</code>, rt must not be specified.</p>
value	a character vector giving the replacement values.

## Details

### Right-hand Side List for the %s{}% and %s!{}% Operators

When supplying a list to the right-hand side of the %s{}% and %s!{}% operators, one can add the argument `at`.

If `at` = "start", the operators will check if the pattern is present/absent at the start of the string.

If `at` = "end", the operators will check if the pattern is present/absent at the end of the string.

Unlike [stri\\_startswith](#) or [stri\\_endswith](#), regex is supported by the %s{}% and %s!{}% operators.

See examples below.

## Vectorized Replacement vs Dictionary Replacement

- Vectorized replacement:  
x, p, and value are of the same length (or recycled to become the same length).  
All occurrences of pattern p[j] in x[j] is replaced with value[j], for every j.
- Dictionary replacement:  
p and value are of the same length, and their length is independent of the length of x.  
For every single string in x, all occurrences of pattern p[1] are replaced with value[1],  
all occurrences of pattern p[2] are replaced with value[2], etc.

Notice that for single replacement, i.e. `rt = "first"` or `rt = "last"`, it makes no sense to distinguish between vectorized or dictionary replacement, since then only a single occurrence is being replaced per string.

See examples below.

## Value

For the `x %s{}% p` and `x %s!{}% p` operators:  
Return logical vectors.

For `strfind()`:  
Returns a list with extractions of all found patterns.

For `strfind(..., i = "all")`:  
Returns a list with all found pattern locations.

For `strfind(..., i = i)` with integer vector `i`:  
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.

For `strfind() <- value`:  
Returns nothing, but performs in-place replacement (using R's default in-place semantics) of the found patterns in variable `x`.

## Note

`strfind()<-` performs in-place replacement.  
Therefore, the character vector or string to perform replacement on, must already exist as a variable.  
So take for example the following code:

```
strfind("hello", p = "e") <- "a" # this obviously does not work
```

```
y <- "hello"
strfind(y, p = "e") <- "a" # this works fine
```

In the above code, the first `strfind()<-` call does not work, because the string needs to exist as a variable.

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

```
# example of %s{}% and %s!{}% ====

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)

#####

# Example of %s{}% and %s!{}% with "at" argument ====

x <- c(paste0(letters, collapse = ""),
      paste0(rev(letters), collapse = ""), NA)
p <- s_fixed("abc", at = "start")
x %s{}% p
stringi::stri_startswith(x, fixed = "abc") # same as above

p <- s_fixed("xyz", at = "end")
x %s{}% p
stringi::stri_endswith(x, fixed = "xyz") # same as above

p <- s_fixed("cba", at = "end")
x %s{}% p
stringi::stri_endswith(x, fixed = "cba") # same as above

p <- s_fixed("zyx", at = "start")
x %s{}% p
stringi::stri_startswith(x, fixed = "zyx") # same as above
```

```
#####
```

```
# Example of transforming ith occurrence ====
```

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

# report ith (second and second-last) vowel locations:
p <- s_regex( # vowels
  rep("A|E|I|O|U", 2),
  case_insensitive = TRUE
)
loc <- strfind(x, p, i = c(2, -2))
print(loc)

# extract ith vowels:
extr <- stringi::stri_sub(x, from = loc)
print(extr)

# replace ith vowels with numbers:
repl <- chartr("aeiou", "12345", extr) # transformation
stringi::stri_sub(x, loc) <- repl
print(x)
```

```
#####
```

```
# Example of strfind for regular vectorized replacement ====
```

```
x <- rep('The quick brown fox jumped over the lazy dog.', 3)
print(x)
p <- c('quick', 'brown', 'fox')
rp <- c('SLOW', 'BLACK', 'BEAR')
x %s{}% p
strfind(x, p)
strfind(x, p) <- rp
print(x)
```

```
#####
```

```
# Example of strfind for dictionary replacement ====
```

```
x <- rep('The quick brown fox jumped over the lazy dog.', 3)
print(x)
p <- c('quick', 'brown', 'fox')
rp <- c('SLOW', 'BLACK', 'BEAR')
# thus dictionary is:
# quick => SLOW; brown => BLACK; fox => BEAR
strfind(x, p, rt = "dict") <- rp
print(x)
```

```
#####
```

```
# Example of strfind for first and last replacement ====
```

```
x <- rep('The quick brown fox jumped over the lazy dog.', 3)
print(x)
p <- s_fixed("the", case_insensitive = TRUE)
rp <- "One"
strfind(x, p, rt = "first") <- rp
print(x)
```

```
x <- rep('The quick brown fox jumped over the lazy dog.', 3)
print(x)
p <- s_fixed("the", case_insensitive = TRUE)
rp <- "Some Other"
strfind(x, p, rt = "last") <- rp
print(x)
```

---

str\_subset\_ops

---

String Subsetting Operators

---

## Description

String subsetting operators.

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

`%s><%` is an alias for `%sget%`.

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

`%s<>%` is an alias for `%strim%`.

## Usage

```
x %sget% ss
```

```
x %strim% ss
```

```
x %s><% ss
```

```
x %s<>% 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

---

Conditional Sub-setting and In-place Replacement of Unreal Values

---

## 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> .<br>For example: <code>\(x)x&gt;0</code> . |
| <code>repl</code> | the replacement value.  |

## Value

For the `x %[if]% cond` and `x %[!if]% cond` 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)
```

s\_pattern

*Pattern Specifications for String Related Operators***Description**

The `%s-%`, `%s!%`, `%ss%` operators, as well as the string search operators ([str\\_search](#)), perform pattern matching for some purpose, where the pattern is given in the second argument (p).

When a character vector or string is given as the second argument (p), 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':

- `s_regex(p, ...)` is equivalent to `list(regex = p, ...)`
- `s_fixed(p, ...)` is equivalent to `list(fixed = p, ...)`
- `s_coll(p, ...)` is equivalent to `list(coll = p, ...)`
- `s_chrcls(p, ...)` is equivalent to `list(charclass = p, ...)`

With the ellipsis (...) being passed to the appropriate 'stringi'-functions when it matches their arguments.

'stringi' infix operators start with "%s", though they all have an alias starting with "%stri". In analogy to that, the above functions start with "s\_" rather than "stri\_", as they are all meant for operators only.

## Usage

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

s_fixed(p, case_insensitive, overlap, ...)

s_coll(
  p,
  locale,
  strength,
  alternate_shifted,
  french,
  uppercase_first,
  case_level,
  numeric,
  normalization,
  ...
)

s_chrcls(p, ...)
```

## Arguments

<code>p</code>	a character vector giving the pattern to search for. about search: <a href="#">regex</a> about search: <a href="#">fixed</a> about search: <a href="#">coll</a> about search: <a href="#">charclass</a>
<code>case_insensitive</code>	see <a href="#">stri_opts_regex</a> and <a href="#">stri_opts_fixed</a> .

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: the at argument for the [str\\_search](#) operators.  
 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 with arguments to be passed to the appropriate operators.

### 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/% s_regex(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(fixed = c("A", "A"), case_insensitive = TRUE)
x %s{%}% p
x %s!{%}% p
p <- s_fixed(c("A", "A"), case_insensitive = TRUE)
x %s{%}% p
x %s!{%}% p

x <- c(paste0(letters[1:13], collapse = ""),
      paste0(letters[14:26], collapse = ""), NA)
p <- s_fixed("abc", at = "start")
x %s{%}% p
stringi::stri_startswith(x, fixed = "abc") # same as above

```

```
p <- s_fixed("xyz", at = "end")
x %s{}% p
stringi::stri_endswith(x, fixed = "xyz") # same as above
```

transform\_if

*transform\_if: Conditional Sub-set Transformation of Atomic objects*

## 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.

For a more `ifelse`-like function where `yes`, `no`, and `other` are vectors, see `kit::iif`.

## 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
```

---

with\_pro

---

*Standard Evaluated Versions of Some Common Expression-Evaluation Functions*


---

**Description**

The `with_pro()` and `aes_pro()` functions are standard-evaluated versions of the expression-evaluation functions [with](#) and `ggplot2::aes`, respectively.

These alternative functions are more programmatically friendly:

They use proper standard evaluation, through the usage of one-sided formulas, instead of non-standard evaluation, tidy evaluation, or similar programmatically unfriendly evaluations.

**Usage**

```
with_pro(data, form)
```

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

**Arguments**

<code>data</code>	a list, environment, or data.frame.
<code>form</code>	a one-sided formula giving the expression to evaluate in <code>with_pro</code> . If the formula has an environment, that environment is used to find any variables or objects not present in <code>data</code> .
<code>...</code>	arguments to be passed to <code>ggplot2::aes</code> , but given as one-sided formulas.

**Details**

The `aes_pro()` function is the standard evaluated alternative to `ggplot2::aes`. Due to the way `aes_pro()` is programmed, it should work even if the tidy evaluation technique changes in 'ggplot2'. To support functions in combinations with references of the variables, the input used here are formula inputs, rather than string inputs. See the Examples section below.

**Value**

For `with_pro()`: see [with](#).  
For `aes_pro()`: see `ggplot2::aes`.

**Non-Standard Evaluation**

Non-Standard Evaluation (sometimes abbreviated as "NSE"), is somewhat controversial. Consider the following example:

```
aplot <- "ggplot2"
library(aplot)
```

What package will be attached? It will not be 'ggplot2', nor will an error occur. Instead, the package 'aplot' will be attached.

This is due to evaluating the expression 'aplot' as a quoted expression, instead of evaluating the contents (i.e. string or formula) of the variable. In other words: Non-Standard Evaluation.

Regular Standard Evaluation does not have the above problem.

**Note**

The `with_pro()` function, like the original [with](#) function, is made for primarily for convenience. When using modelling or graphics functions with an explicit data argument (and typically using [formulas](#)), it is typically preferred to use the data argument of that function, rather than to use either

with(data, ...) or with\_pro(data, ...).

## See Also

[tinycodet\\_safer](#)

## Examples

```
requireNamespace("ggplot2")

d <- import_data("ggplot2", "mpg")

# mutate data:
myform <- ~ displ + cyl + cty + hwy
d$mysum <- with_pro(d, myform)
summary(d)

# plotting data:
x <- ~ cty
y <- ~ sqrt(hwy)
color <- ~ drv

ggplot2::ggplot(d, aes_pro(x, y, color = color)) +
  ggplot2::geom_point()
```

---

x.import

*Helper Functions for the 'tinycodet' Package Import System*

---

## Description

The `help.import()` function finds the help file for functions or topics, including exposed functions/operators as well as functions in a package alias object.

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 or all special attribute(s) from an alias object returned by [import\\_as](#).

## Usage

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

```
is.tinyimport(x)
```

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

## Arguments

<code>...</code>	further arguments to be passed to <a href="#">help</a> .
<code>i</code>	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>
<code>alias</code>	the alias object as created by the <a href="#">import_as</a> function.
<code>x</code>	an existing object (i.e. an assigned variable or a locked constant) to be tested.
<code>which</code>	The attributes to list. If <code>NULL</code> , all attributes will be returned. Possibilities: <code>"pkgs"</code> , <code>"conflicts"</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.

For example:

```
import_as(~ mr., "magrittr")
import_inops(mr.)
help.import(i = mr.$add)
help.import(i = `%>%`)
help.import(i = "add", alias = mr.)
help.import(topic = "%>%", package = "magrittr")
help.import("%>%", package = "magrittr") # same as previous line
```

## 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 imported in the alias object.
- `main_package`: a string giving the name of the main package. Re-exported functions, if present, are taken 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 imported 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 = "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 imported.

For conflicting objects, the last imported ones are used for the ordering.

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

## See Also

[tinycodet\\_import](#)

## Examples

```
import_as(~ to., "tinycodet")
import_inops(to.)
`%s==%` <- stringi::`%s==%`

is.tinyimport(to.) # returns TRUE
is.tinyimport(`%:=%`) # returns TRUE
is.tinyimport(`%s==%`) # returns FALSE: not imported by tinycodet import system

attr.import(to., which = "conflicts")
```

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