

# Package ‘reticulate’

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**Type** Package

**Title** Interface to 'Python'

**Version** 1.7

**Description** Interface to 'Python' modules, classes, and functions. When calling into 'Python', R data types are automatically converted to their equivalent 'Python' types. When values are returned from 'Python' to R they are converted back to R types. Compatible with all versions of 'Python'  $\geq 2.7$ .

**License** Apache License 2.0

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**BugReports** <https://github.com/rstudio/reticulate/issues>

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---

array_reshape	<i>Reshape an Array</i>
---------------	-------------------------

---

### Description

Reshape (reindex) a multi-dimensional array, using row-major (C-style) reshaping semantics by default.

### Usage

```
array_reshape(x, dim, order = c("C", "F"))
```

### Arguments

x	An array
dim	The new dimensions to be set on the array.
order	The order in which elements of x should be read during the rearrangement. "C" means elements should be read in row-major order, with the last index changing fastest; "F" means elements should be read in column-major order, with the first index changing fastest.

### Details

This function differs from e.g. `dim(x) <- dim` in a very important way: by default, `array_reshape()` will fill the new dimensions in row-major (C-style) ordering, while `dim<-()` will fill new dimensions in column-major (Fortran-style) ordering. This is done to be consistent with libraries like NumPy, Keras, and TensorFlow, which default to this sort of ordering when reshaping arrays. See the examples for why this difference may be important.

### Examples

```
## Not run:
# let's construct a 2x2 array from a vector of 4 elements
x <- 1:4

# rearrange will fill the array row-wise
array_reshape(x, c(2, 2))
#      [,1] [,2]
# [1,]  1  2
# [2,]  3  4
# setting the dimensions 'fills' the array col-wise
dim(x) <- c(2, 2)
x
#      [,1] [,2]
# [1,]  1  3
# [2,]  2  4

## End(Not run)
```

---

dict *Create Python dictionary*

---

### Description

Create a Python dictionary object, including a dictionary whose keys are other Python objects rather than character vectors.

### Usage

```
dict(..., convert = FALSE)

py_dict(keys, values, convert = FALSE)
```

### Arguments

...	Name/value pairs for dictionary (or a single named list to be converted to a dictionary).
convert	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <code>py_to_r()</code> function.
keys	Keys to dictionary (can be Python objects)
values	Values for dictionary

### Value

A Python dictionary

### Note

The returned dictionary will not automatically convert it's elements from Python to R. You can do manual conversion with the `py_to_r()` function or pass `convert = TRUE` to request automatic conversion.

---

eng\_python *A reticulate Engine for Knitr*

---

### Description

This provides a reticulate engine for knitr, suitable for usage when attempting to render Python chunks. Using this engine allows for shared state between Python chunks in a document – that is, variables defined by one Python chunk can be used by later Python chunks.

### Usage

```
eng_python(options)
```

**Arguments**

options            Chunk options, as provided by knitr during chunk execution.

**Details**

The engine can be activated by setting (for example)

```
knitr::knit_engines$set(python = reticulate::eng_python)
```

Typically, this will be set within a document's setup chunk, or by the environment requesting that Python chunks be processed by this engine. Note that knitr (since version 1.18) will use the reticulate engine by default when executing Python chunks within an R Markdown document.

---

<code>import</code>	<i>Import a Python module</i>
---------------------	-------------------------------

---

**Description**

Import the specified Python module for calling from R.

**Usage**

```
import(module, as = NULL, convert = TRUE, delay_load = FALSE)
```

```
import_main(convert = TRUE)
```

```
import_builtins(convert = TRUE)
```

```
import_from_path(module, path = ".", convert = TRUE)
```

**Arguments**

<code>module</code>	Module name
<code>as</code>	Alias for module name (affects names of R classes)
<code>convert</code>	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <code>py_to_r()</code> function.
<code>delay_load</code>	TRUE to delay loading the module until it is first used. FALSE to load the module immediately. If a function is provided then it will be called once the module is loaded. If a list containing <code>on_load()</code> and <code>on_error(e)</code> elements is provided then <code>on_load()</code> will be called on successful load and <code>on_error(e)</code> if an error occurs.
<code>path</code>	Path to import from

**Details**

The `import_from_path` function imports a Python module from an arbitrary filesystem path (the directory of the specified python script is automatically added to the `sys.path`).

**Value**

A Python module

**Examples**

```
## Not run:
main <- import_main()
sys <- import("sys")

## End(Not run)
```

---

iterate

*Traverse a Python iterator or generator*

---

**Description**

Traverse a Python iterator or generator

**Usage**

```
iterate(it, f = base::identity, simplify = TRUE)

iter_next(it, completed = NULL)
```

**Arguments**

<code>it</code>	Python iterator or generator
<code>f</code>	Function to apply to each item. By default applies the <code>identity</code> function which just reflects back the value of the item.
<code>simplify</code>	Should the result be simplified to a vector if possible?
<code>completed</code>	Sentinel value to return from <code>iter_next()</code> if the iteration completes (defaults to <code>NULL</code> but can be any R value you specify).

**Details**

Simplification is only attempted all elements are length 1 vectors of type "character", "complex", "double", "integer", or "logical".

**Value**

For `iterate()`, A list or vector containing the results of calling `f` on each item in `x` (invisibly); For `iter_next()`, the next value in the iteration (or the sentinel `completed` value if the iteration is complete).

---

np\_array

*NumPy array*

---

### Description

Create NumPy arrays and convert the data type and in-memory ordering of existing NumPy arrays.

### Usage

```
np_array(data, dtype = NULL, order = "C")
```

### Arguments

data	Vector or existing NumPy array providing data for the array
dtype	Numpy data type (e.g. "float32", "float64", etc.)
order	Memory ordering for array. "C" means C order, "F" means Fortran order.

### Value

A NumPy array object.

---

py

*Interact with the Python Main Module*

---

### Description

The py object provides a means for interacting with the Python main session directly from R. Python objects accessed through py are automatically converted into R objects, and can be used with any other R functions as needed.

### Usage

```
py
```

### Format

An R object acting as an interface to the Python main module.

---

py_available	<i>Check if Python is available on this system</i>
--------------	--

---

**Description**

Check if Python is available on this system

**Usage**

```
py_available(initialize = FALSE)
```

```
py_numpy_available(initialize = FALSE)
```

**Arguments**

initialize	TRUE to attempt to initialize Python bindings if they aren't yet available (defaults to FALSE).
------------	---

**Value**

Logical indicating whether Python is initialized.

**Note**

The `py_numpy_available` function is a superset of the `py_available` function (it calls `py_available` first before checking for NumPy).

---

py_capture_output	<i>Capture and return Python output</i>
-------------------	---

---

**Description**

Capture and return Python output

**Usage**

```
py_capture_output(expr, type = c("stdout", "stderr"))
```

**Arguments**

expr	Expression to capture stdout for
type	Streams to capture (defaults to both stdout and stderr)

**Value**

Character vector with output



---

py_config	<i>Python configuration</i>
-----------	-----------------------------

---

**Description**

Information on Python and Numpy versions detected

**Usage**

```
py_config()
```

**Value**

Python configuration object; Logical indicating whether Python bindings are available

---

py_discover_config	<i>Discover the version of Python to use with reticulate.</i>
--------------------	---

---

**Description**

This function enables callers to check which versions of Python will be discovered on a system as well as which one will be chosen for use with reticulate.

**Usage**

```
py_discover_config(required_module = NULL, use_environment = NULL)
```

**Arguments**

required_module	A optional module name that must be available in order for a version of Python to be used.
use_environment	An optional virtual/conda environment name to prefer in the search

**Value**

Python configuration object.

---

 py\_func

*Wrap an R function in a Python function with the same signature.*


---

### Description

This function could wrap an R function in a Python function with the same signature. Note that the signature of the R function must not contain esoteric Python-incompatible constructs.

### Usage

```
py_func(f)
```

### Arguments

f                    An R function

### Value

A Python function that calls the R function f with the same signature.

---

 py\_function\_custom\_scaffold

*Custom Scaffolding of R Wrappers for Python Functions*


---

### Description

This function can be used to generate R wrapper for a specified Python function while allowing to inject custom code for critical parts of the wrapper generation, such as process the any part of the docs obtained from [py\\_function\\_docs\(\)](#) and append additional roxygen fields. The result from execution of python\_function is assigned to a variable called python\_function\_result that can also be processed by postprocess\_fn before writing the closing curly braces for the generated wrapper function.

### Usage

```
py_function_custom_scaffold(python_function, r_function = NULL,
  additional_roxygen_fields = NULL, process_docs_fn = function(docs) docs,
  process_param_fn = function(param, docs) param,
  process_param_doc_fn = function(param_doc, docs) param_doc,
  postprocess_fn = function() { }, file_name = NULL)
```

**Arguments**

python_function	Fully qualified name of Python function or class constructor (e.g. tf\$layers\$average_pooling1d)
r_function	Name of R function to generate (defaults to name of Python function if not specified)
additional_roxygen_fields	A list of additional roxygen fields to write to the roxygen docs, e.g. list(export = "", rdname = "gene")
process_docs_fn	A function to process docs obtained from reticulate::py_function_docs(python_function).
process_param_fn	A function to process each parameter needed for python_function before executing python_function.
process_param_doc_fn	A function to process the roxygen docstring for each parameter.
postprocess_fn	A function to inject any custom code in the form of a string before writing the closing curly braces for the generated wrapper function.
file_name	The file name to write the generated wrapper function to. If NULL, the generated wrapper will only be printed out in the console.

**Examples**

```
## Not run:

library(tensorflow)
library(stringr)

# Example of a `process_param_fn` to cast parameters with default values
# that contains "L" to integers
process_int_param_fn <- function(param, docs) {
  # Extract the list of parameters that have integer values as default
  int_params <- gsub(
    " = [-]?[0-9]+L",
    "",
    str_extract_all(docs$signature, "[A-z]+ = [-]?[0-9]+L")[[1]])
  # Explicitly cast parameter in the list obtained above to integer
  if (param %in% int_params) {
    param <- paste0("as.integer(", param, ")")
  }
  param
}

# Note that since the default value of parameter `k` is `1L`. It is wrapped
# by `as.integer()` to ensure it's casted to integer before sending it to `tf$nn$top_k`
# for execution. We then print out the python function result.
py_function_custom_scaffold(
  "tf$nn$top_k",
  r_function = "top_k",
  process_param_fn = process_int_param_fn,
  postprocess_fn = function() { "print(python_function_result)" })
```

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

---

py\_get\_attr                    *Get an attribute of a Python object*

---

**Description**

Get an attribute of a Python object

**Usage**

```
py_get_attr(x, name, silent = FALSE)
```

**Arguments**

x	Python object
name	Attribute name
silent	TRUE to return NULL if the attribute doesn't exist (default is FALSE which will raise an error)

**Value**

Attribute of Python object

---

py\_has\_attr                    *Check if a Python object has an attribute*

---

**Description**

Check whether a Python object x has an attribute name.

**Usage**

```
py_has_attr(x, name)
```

**Arguments**

x	A python object.
name	The attribute to be accessed.

**Value**

TRUE if the object has the attribute name, and FALSE otherwise.

---

py\_help

*Documentation for Python Objects*

---

**Description**

Documentation for Python Objects

**Usage**

py\_help(object)

**Arguments**

object            Object to print documentation for

---

py\_id

*Unique identifier for Python object*

---

**Description**

Get a globally unique identifier for a Python object.

**Usage**

py\_id(object)

**Arguments**

object            Python object

**Value**

Unique identifier (as integer) or NULL

**Note**

In the current implementation of CPython this is the memory address of the object.

---

py\_install                      *Install Python packages*

---

### Description

Install Python packages into a virtualenv or conda env.

### Usage

```
py_install(packages, envname = "r-reticulate", method = c("auto",
  "virtualenv", "conda"), conda = "auto", ...)
```

### Arguments

packages	Character vector with package names to install
envname	Name of environment to install packages into
method	Installation method. By default, "auto" automatically finds a method that will work in the local environment. Change the default to force a specific installation method. Note that the "virtualenv" method is not available on Windows.
conda	Path to conda executable (or "auto" to find conda using the PATH and other conventional install locations).
...	Additional arguments passed to <a href="#">conda_install()</a> or <a href="#">virtualenv_install()</a> .

### Details

On Linux and OS X the "virtualenv" method will be used by default ("conda" will be used if virtualenv isn't available). On Windows, the "conda" method is always used.

### See Also

[conda-tools](#), [virtualenv-tools](#)

---

py\_is\_null\_xptr                      *Check if a Python object is a null externalptr*

---

### Description

Check if a Python object is a null externalptr

### Usage

```
py_is_null_xptr(x)

py_validate_xptr(x)
```

**Arguments**

x Python object

**Details**

When Python objects are serialized within a persisted R environment (e.g. .RData file) they are deserialized into null externalptr objects (since the Python session they were originally connected to no longer exists). This function allows you to safely check whether whether a Python object is a null externalptr.

The `py_validate` function is a convenience function which calls `py_is_null_xptr` and throws an error in the case that the `xptr` is `NULL`.

**Value**

Logical indicating whether the object is a null externalptr

---

`py_iterator` *Create a Python iterator from an R function*

---

**Description**

Create a Python iterator from an R function

**Usage**

```
py_iterator(fn, completed = NULL)
```

**Arguments**

fn R function with no arguments.  
 completed Special sentinel return value which indicates that iteration is complete (defaults to `NULL`)

**Details**

Python generators are functions that implement the Python iterator protocol. In Python, values are returned using the `yield` keyword. In R, values are simply returned from the function.

In Python, the `yield` keyword enables successive iterations to use the state of previous iterations. In R, this can be done by returning a function that mutates it's enclosing environment via the `<<-` operator. For example:

```
sequence_generator <- function(start) {
  value <- start
  function() {
    value <<- value + 1
    value
  }
}
```

Then create an iterator using `py_iterator()`:

```
g <- py_iterator(sequence_generator(10))
```

### Value

Python iterator which calls the R function for each iteration.

### Ending Iteration

In Python, returning from a function without calling `yield` indicates the end of the iteration. In R however, `return` is used to yield values, so the end of iteration is indicated by a special return value (NULL by default, however this can be changed using the `completed` parameter). For example:

```
sequence_generator <-function(start) {
  value <- start
  function() {
    value <<- value + 1
    if (value < 100)
      value
    else
      NULL
  }
}
```

### Threading

Some Python APIs use generators to parallelize operations by calling the generator on a background thread and then consuming it's results on the foreground thread. The `py_iterator()` function creates threadsafe iterators by ensuring that the R function is always called on the main thread (to be compatible with R's single-threaded runtime) even if the generator is run on a background thread.

---

<code>py_last_error</code>	<i>Get or clear the last Python error encountered</i>
----------------------------	---

---

### Description

Get or clear the last Python error encountered

### Usage

```
py_last_error()
```

```
py_clear_last_error()
```

### Value

For `py_last_error()`, a list with the type, value, and traceback for the last Python error encountered (can be NULL if no error has yet been encountered).



---

py_len	<i>Length of Python object</i>
--------	--------------------------------

---

**Description**

Get the length of a Python object (equivalent to the Python len() built in function).

**Usage**

```
py_len(x)
```

**Arguments**

x	Python object
---	---------------

**Value**

Length as integer

---

py_list_attributes	<i>List all attributes of a Python object</i>
--------------------	---

---

**Description**

List all attributes of a Python object

**Usage**

```
py_list_attributes(x)
```

**Arguments**

x	Python object
---	---------------

**Value**

Character vector of attributes

---

py\_module\_available    *Check if a Python module is available on this system.*

---

**Description**

Check if a Python module is available on this system.

**Usage**

```
py_module_available(module)
```

**Arguments**

module	Name of module
--------	----------------

**Value**

Logical indicating whether module is available

---

py\_run                    *Run Python code*

---

**Description**

Execute code within the the `__main__` Python module.

**Usage**

```
py_run_string(code, local = FALSE, convert = TRUE)
```

```
py_run_file(file, local = FALSE, convert = TRUE)
```

```
py_eval(code, convert = TRUE)
```

**Arguments**

code	Code to execute
local	Whether to create objects in a local/private namespace (if FALSE, objects are created within the main module).
convert	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <code>py_to_r()</code> function.
file	Source file

**Value**

For `py_eval()`, the result of evaluating the expression; For `py_run_string()` and `py_run_file()`, the dictionary associated with the code execution.

---

py\_save\_object            *Save and load Python objects with pickle*

---

**Description**

Save and load Python objects with pickle

**Usage**

```
py_save_object(object, filename, pickle = "pickle")
```

```
py_load_object(filename, pickle = "pickle")
```

**Arguments**

object	Object to save
filename	File name
pickle	The implementation of pickle to use (defaults to "pickle" but could e.g. also be "cPickle")

---

py\_set\_attr            *Set an attribute of a Python object*

---

**Description**

Set an attribute of a Python object

**Usage**

```
py_set_attr(x, name, value)
```

**Arguments**

x	Python object
name	Attribute name
value	Attribute value

---

py_set_seed	<i>Set Python and NumPy random seeds</i>
-------------	--

---

### Description

Set various random seeds required to ensure reproducible results. The provided seed value will establish a new random seed for Python and NumPy, and will also (by default) disable hash randomization.

### Usage

```
py_set_seed(seed, disable_hash_randomization = TRUE)
```

### Arguments

seed	A single value, interpreted as an integer
disable_hash_randomization	Disable hash randomization, which is another common source of variable results. See <a href="https://docs.python.org/3.4/using/cmdline.html#envvar-PYTHONHASHSEED">https://docs.python.org/3.4/using/cmdline.html#envvar-PYTHONHASHSEED</a>

### Details

This function does not set the R random seed, for that you should call `set.seed()`.

---

py_str	<i>An S3 method for getting the string representation of a Python object</i>
--------	--

---

### Description

An S3 method for getting the string representation of a Python object

### Usage

```
py_str(object, ...)
```

### Arguments

object	Python object
...	Unused

### Details

The default implementation will call `PyObject_Str` on the object.

### Value

Character vector

---

py\_suppress\_warnings    *Suppress Python warnings for an expression*

---

**Description**

Suppress Python warnings for an expression

**Usage**

```
py_suppress_warnings(expr)
```

**Arguments**

expr                    Expression to suppress warnings for

**Value**

Result of evaluating expression

---

py\_unicode              *Convert to Python Unicode Object*

---

**Description**

Convert to Python Unicode Object

**Usage**

```
py_unicode(str)
```

**Arguments**

str                    Single element character vector to convert

**Details**

By default R character vectors are converted to Python strings. In Python 3 these values are unicode objects however in Python 2 they are 8-bit string objects. This function enables you to obtain a Python unicode object from an R character vector when running under Python 2 (under Python 3 a standard Python string object is returned).

---

r-py-conversion	<i>Convert between Python and R objects</i>
-----------------	---

---

**Description**

Convert between Python and R objects

**Usage**

```
r_to_py(x, convert = FALSE)
```

```
py_to_r(x)
```

**Arguments**

x	A Python object.
convert	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <a href="#">py_to_r()</a> function.

**Value**

An R object, as converted from the Python object.

---

repl_python	<i>Run a Python REPL</i>
-------------	--------------------------

---

**Description**

This function provides a Python REPL in the R session, which can be used to interactively run Python code. All code executed within the REPL is run within the Python main module, and any generated Python objects will persist in the Python session after the REPL is detached.

**Usage**

```
repl_python(module = NULL, quiet = getOption("reticulate.repl.quiet",
  default = FALSE))
```

**Arguments**

module	An (optional) Python module to be imported before the REPL is launched.
quiet	Boolean; print a startup banner when launching the REPL? If TRUE, the banner will be suppressed.

## Details

When working with R and Python scripts interactively, one can activate the Python REPL with `repl_python()`, run Python code, and later run `exit` to return to the R console.

## See Also

[py](#), for accessing objects created using the Python REPL.

## Examples

```
## Not run:

# enter the Python REPL, create a dictionary, and exit
repl_python()
dictionary = {'alpha': 1, 'beta': 2}
exit

# access the created dictionary from R
py$dictionary
# $alpha
# [1] 1
#
# $beta
# [1] 2

## End(Not run)
```

## Description

R interface to Python modules, classes, and functions. When calling into Python R data types are automatically converted to their equivalent Python types. When values are returned from Python to R they are converted back to R types. The reticulate package is compatible with all versions of Python  $\geq 2.7$ . Integration with NumPy requires NumPy version 1.6 or higher.

---

source_python	<i>Read and evaluate a Python script</i>
---------------	--

---

**Description**

Evaluate a Python script within the Python main module, then make all public (non-module) objects within the main Python module available within the specified R environment.

**Usage**

```
source_python(file, envir = parent.frame(), convert = TRUE)
```

**Arguments**

file	Source file
envir	The environment to assign Python objects into (for example, <code>parent.frame()</code> or <code>globalenv()</code> ). Specify NULL to not assign Python objects.
convert	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <a href="#">py_to_r()</a> function.

**Details**

To prevent assignment of objects into R, pass NULL for the `envir` parameter.

---

tuple	<i>Create Python tuple</i>
-------	----------------------------

---

**Description**

Create a Python tuple object

**Usage**

```
tuple(..., convert = FALSE)
```

**Arguments**

...	Values for tuple (or a single list to be converted to a tuple).
convert	TRUE to automatically convert Python objects to their R equivalent. If you pass FALSE you can do manual conversion using the <a href="#">py_to_r()</a> function.

**Value**

A Python tuple



**Note**

The returned tuple will not automatically convert it's elements from Python to R. You can do manual conversion with the `py_to_r()` function or pass `convert = TRUE` to request automatic conversion.

---

use_python	<i>Configure which version of Python to use</i>
------------	---

---

**Description**

Configure which version of Python to use

**Usage**

```
use_python(python, required = FALSE)
```

```
use_virtualenv(virtualenv, required = FALSE)
```

```
use_condaenv(condaenv, conda = "auto", required = FALSE)
```

**Arguments**

python	Path to Python binary
required	Is this version of Python required? If TRUE then an error occurs if it's not located. Otherwise, the version is taken as a hint only and scanning for other versions will still proceed.
virtualenv	Directory of Python virtualenv
condaenv	Name of Conda environment
conda	Conda executable. Default is "auto", which checks the PATH as well as other standard locations for Anaconda installations.

---

virtualenv-tools	<i>Interface to virtualenv</i>
------------------	--------------------------------

---

**Description**

R functions for managing Python **virtual environments**

**Usage**

```

virtualenv_list()

virtualenv_root()

virtualenv_create(envname)

virtualenv_install(envname, packages, ignore_installed = FALSE)

virtualenv_remove(envname, packages = NULL, confirm = interactive())

```

**Arguments**

envname	Name of virtual environment
packages	Character vector with package names to install or remove.
ignore_installed	Ignore any previously installed versions of packages
confirm	Confirm before removing packages or virtual environments

**Details**

Virtual environments are by default located at `~/.virtualenvs`. You can change this behavior by defining the `WORKON_HOME` environment variable.

Virtual environment functions are not supported on Windows (the use of [conda environments](#) is recommended on Windows).

**Value**

`virtualenv_list()` returns a character vector with the names of available virtual environments.  
`virtualenv_root()` returns the root directory for virtual environments.

---

```
with.python.builtin.object
```

*Evaluate an expression within a context.*

---

**Description**

The `with` method for objects of type `python.builtin.object` implements the context manager protocol used by the Python `with` statement. The passed object must implement the **context manager** (`__enter__` and `__exit__` methods).

**Usage**

```

## S3 method for class 'python.builtin.object'
with(data, expr, as = NULL, ...)

```

**Arguments**

<code>data</code>	Context to enter and exit
<code>expr</code>	Expression to evaluate within the context
<code>as</code>	Name of variable to assign context to for the duration of the expression's evaluation (optional).
<code>...</code>	Unused

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