

Use Python with R with reticulate :: CHEATSHEET



The `reticulate` package lets you use Python and R together seamlessly in R code, in R Markdown documents, and in the RStudio IDE.

Python in R Markdown

(Optional) Build Python env to use.

Add `knitr::knit_engines$set(python = reticulate::eng_python)` to the setup chunk to set up the reticulate Python engine (not required for `knitr >= 1.18`).

Suggest the Python environment to use, in your setup chunk.

Begin Python chunks with ````{python}`. Chunk options like `echo`, `include`, etc. all work as expected.

Use the `py` object to access objects created in Python chunks from R chunks.

Python chunks all execute within a **single** Python session so you have access to all objects created in previous chunks.

Use the `r` object to access objects created in R chunks from Python chunks.

Output displays below chunk, including matplotlib plots.

```
python.Rmd x
1 ```{r setup, include = FALSE}
2 library(reticulate)
3 virtualenv_create("fmri-proj")
4 py_install("seaborn", envname = "fmri-proj")
5 use_virtualenv("fmri-proj")
6 ```
7 ```{python, echo = FALSE}
8 import seaborn as sns
9 fmri = sns.load_dataset("fmri")
10 ```
11 ```{r}
12 f1 <- subset(py$fmri, region == "parietal")
13 ```
14 ```{python}
15 import matplotlib as mpl
16 sns.lmplot("timepoint", "signal", data=r.f1)
17 mpl.pyplot.show()
18 ```
```

R Console: A scatter plot of signal vs timepoint for the parietal region.

R Markdown: A histogram of signal values.

```
python.R x
1 library(reticulate)
2 py_install("seaborn")
3 use_virtualenv("r-reticulate")
4 ```
5 sns <- import("seaborn")
6 ```
7 fmri <- sns.load_dataset("fmri")
8 dim(fmri)
9 ```
10 # creates tips
11 source_python("python.py")
12 dim(tips)
13 ```
14 # creates tips in main
15 py_run_file("python.py")
16 dim(py$tips)
17 ```
18 py_run_string("print(tips.shape)")
19 ```
```

Object Conversion

Tip: To index Python objects begin at 0, use integers, e.g. `0L`

Reticulate provides automatic built-in conversion between Python and R for many Python types.

R	↔	Python
Single-element vector		Scalar
Multi-element vector		List
List of multiple types		Tuple
Named list		Dict
Matrix/Array		NumPy ndarray
Data Frame		Pandas DataFrame
Function		Python function
NULL, TRUE, FALSE		None, True, False

Or, if you like, you can convert manually with

`py_to_r(x)` Convert a Python object to an R object. Also `r_to_py()`. `py_to_r(x)`

`tuple(..., convert = FALSE)` Create a Python tuple. `tuple("a", "b", "c")`

`dict(..., convert = FALSE)` Create a Python dictionary object. Also `py_dict()` to make a dictionary that uses Python objects as keys. `dict(foo = "bar", index = 42L)`

`np_array(data, dtype = NULL, order = "C")` Create NumPy arrays. `np_array(c(1:8), dtype = "float16")`

`array_reshape(x, dim, order = c("C", "F"))` Reshape a Python array. `x <- 1:4; array_reshape(x, c(2, 2))`

`py_func(f)` Wrap an R function in a Python function with the same signature. `py_func(xor)`

`py_main_thread_func(f)` Create a function that will always be called on the main thread.

`iterate(it, f = base::identity, simplify = TRUE)` Apply an R function to each value of a Python iterator or return the values as an R vector, draining the iterator as you go. Also `iter_next()` and `as_iterator()`. `iterate(iter, print)`

`py_iterator(fn, completed = NULL)` Create a Python iterator from an R function. `seq_gen <- function(x){ n <- x; function() {n <- n + 1; n}}; py_iterator(seq_gen(9))`

Helpers

`py_capture_output(expr, type = c("stdout", "stderr"))` Capture and return Python output. Also `py_suppress_warnings()`. `py_capture_output("x")`

`py_get_attr(x, name, silent = FALSE)` Get an attribute of a Python object. Also `py_set_attr()`, `py_has_attr()`, and `py_list_attributes()`. `py_get_attr(x)`

`py_help(object)` Open the documentation page for a Python object. `py_help(sns)`

`py_last_error()` Get the last Python error encountered. Also `py_clear_last_error()` to clear the last error. `py_last_error()`

`py_save_object(object, filename, pickle = "pickle", ...)` Save and load Python objects with pickle. Also `py_load_object()`. `py_save_object(x, "x.pickle")`

`with(data, expr, as = NULL, ...)` Evaluate an expression within a Python context manager.

```
py <- import_builtins();
with(py$open("output.txt", "w") %as% file,
     file$write("Hello, there!"))
```

Python in R

Call Python from R code in three ways:

IMPORT PYTHON MODULES

Use `import()` to import any Python module. Access the attributes of a module with `$`.

- `import(module, as = NULL, convert = TRUE, delay_load = FALSE)` Import a Python module. If `convert = TRUE`, Python objects are converted to their equivalent R types. Also `import_from_path()`. `import("pandas")`
- `import_main(convert = TRUE)` Import the main module, where Python executes code by default. `import_main()`
- `import_builtins(convert = TRUE)` Import Python's built-in functions. `import_builtins()`

SOURCE PYTHON FILES

Use `source_python()` to source a Python script and make the Python functions and objects it creates available in the calling R environment.

- `source_python(file, envir = parent.frame(), convert = TRUE)` Run a Python script, assigning objects to a specified R environment. `source_python("file.py")`

RUN PYTHON CODE

Execute Python code into the **main** Python module with `py_run_file()` or `py_run_string()`.

- `py_run_string(code, local = FALSE, convert = TRUE)` Run Python code (passed as a string) in the main module. `py_run_string("x = 10"); py$x`
- `py_run_file(file, local = FALSE, convert = TRUE)` Run Python file in the main module. `py_run_file("script.py")`
- `py_eval(code, convert = TRUE)` Run a Python expression, return the result. Also `py_call()`. `py_eval("1 + 1")`

Access the results, and anything else in Python's **main** module, with `py`.

- `py` An R object that contains the Python main module and the results stored there. `py$x`



Python in the IDE

Syntax highlighting for Python scripts and chunks.

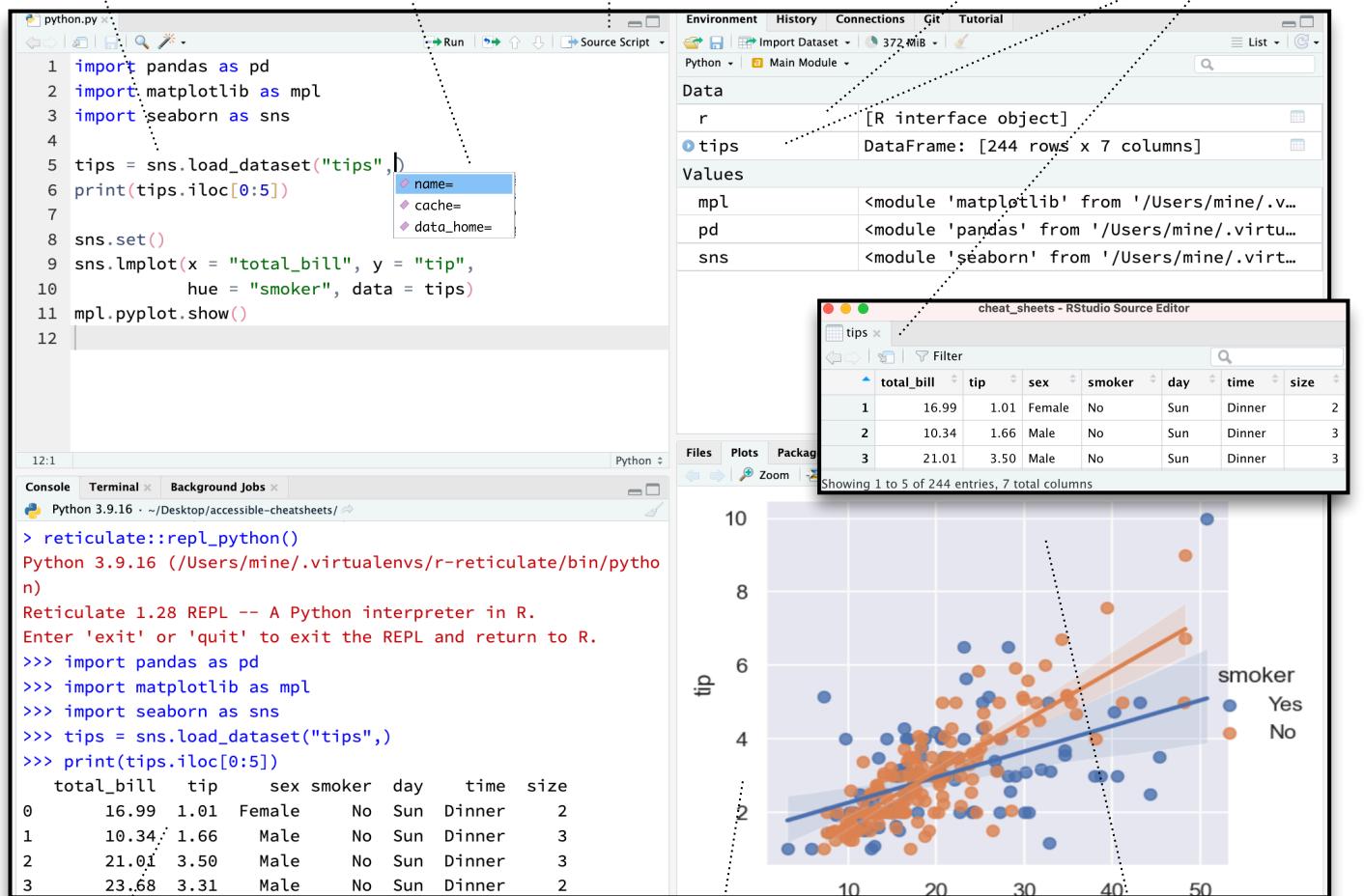
Tab completion for Python functions and objects (and Python modules imported in R scripts).

Source Python scripts.

Execute Python code line by line with **Cmd + Enter** (**Ctrl + Enter**).

View Python objects in the Environment Pane.

View Python objects in the Data Viewer.



A Python REPL opens in the console when you run Python code with a keyboard shortcut. Type **exit** to close.

Python REPL

A REPL (Read, Eval, Print Loop) is a command line where you can run Python code and view the results.

1. Open in the console with **repl_python()**, or by running code in a Python script with **Cmd + Enter** (**Ctrl + Enter**).
2. Type commands at **>>>** prompt.
3. Press **Enter** to run code.
4. Type **exit** to close and return to R console.

```
Console Terminal x Background Jobs x
> reticulate::repl_python()
Python 3.9.16 (/Users/mine/.virtualenvs/r-reticulate/bin/python)
Reticulate 1.28 REPL -- A Python interpreter in R.
Enter 'exit' or 'quit' to exit the REPL and return to R.

>>> import seaborn as sns
>>> tips = sns.load_dataset("tips")
>>> tips.shape
(244, 7)
>>> exit
> |
```



Configure Python

Reticulate binds to a local instance of Python when you first call **import()** directly or implicitly from an R session. To control the process, find or build your desired Python instance. Then suggest your instance to reticulate. **Restart R to unbind**.

Find Python

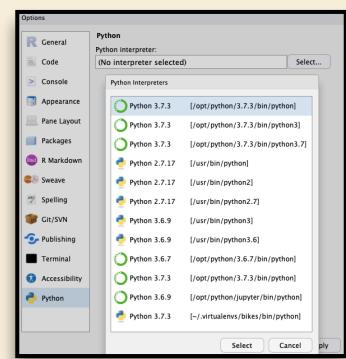
- **install_python(version, list = FALSE, force = FALSE)** Download and install Python. `install_python("3.9.16")`
- **py_available(initialize = FALSE)** Check if Python is available on your system. Also **py_module_available()** and **py_numpy_module()**. `py_available()`
- **py_discover_config()** Return all detected versions of Python. Use **py_config()** to check which version has been loaded. `py_config()`
- **virtualenv_list()** List all available virtual environments. Also **virtualenv_root()**. `virtualenv_list()`
- **conda_list(conda = "auto")** List all available conda environments. Also **conda_binary()** and **conda_version()**. `conda_list()`

Suggest an env to use

Set a default Python interpreter in the RStudio IDE Global or Project Options.

Go to **Tools > Global Options... > Python** for Global Options.

Within a project, go to **Tools > Project Options... > Python**.



Otherwise, to choose an instance of Python to bind to, reticulate scans the instances on your computer in the following order, **stopping at the first instance that contains the module called by import()**.

1. The instance referenced by the environment variable **RETICULATE PYTHON** (if specified). **Tip: set in .Renviron file.**

- **Sys.setenv(RETICULATE PYTHON = PATH)** Set default Python binary. Persists across sessions! Undo with **Sys.unsetenv()**. `Sys.setenv(RETICULATE PYTHON = "/usr/local/bin/python")`

2. The instances referenced by **use_** functions if called before **import()**. Will fail silently if called after **import** unless **required = TRUE**.

- **use_python(python, required = FALSE)** Suggest a Python binary to use by path. `use_python("/usr/local/bin/python")`

- **use_virtualenv(virtualenv = NULL, required = FALSE)** Suggest a Python virtualenv. `use_virtualenv("~/myenv")`

- **use_condaenv(condaenv = NULL, conda = "auto", required = FALSE)** Suggest a conda env to use. `use_condaenv(condaenv = "r-nlp", conda = "/opt/anaconda3/bin/conda")`

3. Within virtualenvs and conda envs that carry the same name as the imported module. e.g. `~/anaconda/envs/nltk` for `import("nltk")`

4. At the location of the Python binary discovered on the system PATH (i.e. `Sys.which("python")`)

5. At customary locations for Python, e.g. `/usr/local/bin/python, /opt/local/bin/python...`

Create a Python env

- **virtualenv_create(envname = NULL, ...)** Create a new virtual environment. `virtualenv_create("r-pandas")`
- **conda_create(envname = NULL, ...)** Create a new conda environment. `conda_create("r-pandas", packages = "pandas")`

Install Packages

Install Python packages with R (below) or the shell:
pip install SciPy
conda install SciPy

- **py_install(packages, envname, ...)** Installs Python packages into a Python env. `py_install("pandas")`
- **virtualenv_install(envname, packages, ...)** Install a package within a virtualenv. Also **virtualenv_remove()**. `virtualenv_install("r-pandas", packages = "pandas")`
- **conda_install(envname, packages, ...)** Install a package within a conda env. Also **conda_remove()**. `conda_install("r-pandas", packages = "plotly")`