Jse Python with R with reticulate :: CHEAT SHEET

rhe **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

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

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

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

Output displays below chunk, including matplotlib plots.

R Console

py_run_string("print(tips.shape)") source_python("python.py") py_run_file("python.py") creates tips in main dim(py\$tips) dim(tips) 11 14 15 16 17

data=r.f1)

sns)lmplot("timepoint","signal",

mpl.pyplot.show()

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import matplotlib as mpl

Object Conversion

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

between Python and R for many Python types.

<u> </u>	Python	u
Single-element vector	Scalar	
Multi-element vector	List	
List of multiple types	Tuple	
Named list	Dict	
Matrix/Array	NumPy ndarray	>
Data Frame	Pandas DataFrame	ame
Function	Python function	'n
NULL, TRUE, FALSE	None, True, False	lse

Or, if you like, you can convert manually with py_to_r(x) Convert a Python object to

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

an R object. Also **r_to_py**. $py_to_r(x)$

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)

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

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

each value of a Python iterator or return the values iterate(..., convert = FALSE) Apply an R function to as an R vector, draining the iterator as you go. Also iter_next and as_iterator. iterate(iter, print) $np_array(data, dtype = NULL, order = "C")$ 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; $array_reshape(x, c(2, 2))$

Helpers

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

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")

py <- import_builtins(); with(py\$open("output.txt", "w") %as% file, { file\$write("Hello, there!")}} **with**(data, expr, as = NULL, ...) Evaluate an expression within a Python context manager.

Python in R code **→** Source •

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→ Run →

Call Python from R in three ways:

IMPORT PYTHON MODULES

use_virtualenv("r-reticulate")

"fmri-proj")

py_install("seaborn", envname =

use_virtualenv("fmri-proj")

o: 0

virtualenv_create("fmri-proj") 1 * ```{r setup, include = FALSE}

2 library(reticulate)

py_install("seaborn" 1 library(reticulate)

sns <- import("seaborn")

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

Python objects are converted to their equivalent R types. Also import_from_path.import("pandas") import(module, as = NULL, convert = TRUE, delay_load = FALSE) Import a Python module. If convert = TRUE,

fmri <- sns\$load_dataset("fmri")</pre>

dim(fmri)

H

fmri = sns.load_dataset("fmri")

```{python, echo = FALSE}

import seaborn as sns

creates tips

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f1 <- subset(py\$fmri) region == "parietal")

1. In

- Import the main module, where Python executes code by default. import\_main() import\_main(convert = TRUE)
- import\_builtins(convert = TRUE)
  Import Python's built-in functions.

#### **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, environment. source\_python("file.py") assigning objects to a specified R

#### **RUN PYTHON CODE**

module with **py\_run\_file**() or **py\_run\_string**(). Execute Python code into the **main** Python

- module.  $py\_run\_string("x=10"); py$x$ py\_run\_string(code, local = FALSE, convert = TRUE) Run Python code (passed as a string) in the main
- py\_run\_file(file, local = FALSE, convert = **TRUE) Run Python file in the main** module. py\_run\_file("script.py")
- a Python expression, return the result. py\_eval(code, convert = TRUE) Run 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