# Statistics 360: Advanced R for Data Science Lecture 11

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### R and Python: References

- ► The reticulate website: https://rstudio.github.io/reticulate/
- Python setup: https://docs.python.org/3/using/index.html
- Python tutorial: https://docs.python.org/3/tutorial/

### Calling Python from R

- Use cases:
  - ► Tap into a growing set of tools for data science, such as scikit-learn https://scikit-learn.org/stable/ , keras https://keras.io/ , . . .
  - Workflow requires substantial computations in **both** languages; e.g., fit a neural network in Python, plot results using ggplot2
- Not recommended:
  - Using R/RStudio as a development environment for Python.
  - Better to go all-in with Python and use Jupyter Notebooks, Spyder, etc. as your IDE.

### Prerequisites

Install and load the reticulate package

```
# install.packages("reticulate")
library(reticulate)
```

- ► Install Python
  - See the Python setup link on the references slide, or the Python distributions on the next slide
  - Warning for Mac users: Macs with the Apple M1 processor can run both M1 and x86 (Intel processor) binaries. Make sure your R and Python installations are for the same processor, or you will get errors when you try to start Python from R.
- ▶ Install the Python packages that you need.
  - More on this below.

## Python packages

- Python packages are like R packages.
- Install with pip or conda.
  - ➤ So far I've only been able to find a miniconda installer to work with my M1 version of R, so I'll discuss conda-based installs.
- conda comes with the Python distributions anaconda and miniconda https:
  - //docs.conda.io/projects/conda/en/latest/user-guide/install
    - miniconda is smaller and faster to install, but you will need to install packages yourself
    - anaconda includes all the python packages you will need for data science but takes more disk space and probably includes stuff you will never use
- Once you have conda install packages from the command line with conda install <package>.

## Python environments

- ▶ Python environments, like RStudio projects, are used to compartmentalize your work with Python.
  - ► A complete Python installation, including its own Python executable and packages.
  - Create from the command line with conda create --name
    <env\_name>
  - ► Then "activate" with conda activate <env\_name> and "de-activate" with conda deactivate.

### Installing packages with reticulate, Part I

- See https: //rstudio.github.io/reticulate/articles/python\_packages.html
- ▶ We'll first install into the "base" Python environment

```
library(reticulate)
#conda_install(packages=c("pandas", "scikit-learn")) # commented out to
```

### Installing packages with reticulate, Part II

- Now install into another Python environment
- ▶ Note: Not currently working on my system, but it used to!

#### library(reticulate)

### Using a conda environment

► For each R session in which you want to use your conda environment:

```
library(reticulate)
use_condaenv("r-reticulate",required=TRUE)
# Can then call py_config() to see which Python version is being
```

## Python embedded in RMarkdown

Example from https://pandas.pydata.org/pandasdocs/stable/user\_guide/10min.html

```
# code chunk header is ```{python} rather than ```{r}
import numpy as np
import pandas as pd

df = pd.DataFrame(np.random.randn(3,4),columns=['A','B','C','D']

df

## A B C D

## 0 0.116952 -0.644985 -0.884050 -0.649658

## 1 -0.221388 0.596401 0.335031 0.087408

## 2 -0.473363 -1.736616 -0.293679 1.929203
```

## Importing Python packages (modules)

You can also import Python packages into R and call their functions directly.

```
npr <- import("numpy.random")
pd <- import("pandas") # import is from reticulate
df <- pd$DataFrame(npr$randn(3L,4L),columns=c('A','B','C','D'))
df

## A B C D
## 1 0.3965882 1.32744536 -0.1481296 -0.2214696
## 2 -1.3726563 -1.46410869 -0.6774934 -0.4406373
## 3 -1.3183013 -0.07735525 -0.1054971 -0.7835564
```

### Notes

- Access Python functions from an imported package with \$.
- ► The randn() function requires integer arguments have to use 3L and 4L to pass integers.
  - reticulate converts R vectors of length 1 to Python scalars.
  - ▶ In general reticulate will try to convert to/from appropriate data types. See the list at https:
    - //rstudio.github.io/reticulate/index.html#type-conversions
- ► I used the numpy random number generator, but passed column names as an R character vector.
  - reticulate converts this to a python list.

### Sourcing Python scripts

- ► Source with source\_python(). By default, objects created by the Python script are made available as R objects in your workspace, and also in the hidden environment py.
- Pass an environment if you want to store somewhere else.

```
# source_python("lec11_1.py") # default
my_py_env <- new.env()
source_python("lec11_1.py",envir=my_py_env)
ls(my_py_env)

## [1] "df" "r"
my_py_env$df

## A B C D
## 1 -0.7867444 0.4330144 1.03313035 -1.415685
## 2 1.6289353 -0.5534447 -0.12533629 -1.269741
## 3 -0.9566918 -1.6008279 0.03205155 1.076558</pre>
```

## Another example

```
source_python("lec11_2.py")
MSE
## [1] 2859.69
ddat <- data.frame(Y=diabetes_y, diabetes_X)</pre>
library(ggplot2)
ggplot(ddat,aes(x=X1,y=Y)) + geom_point() + geom_smooth()
  300 -
 200 -
  100 -
           -0.10
                                 -0.05
                                                                             0.05
                                                                                                   0.10
                                                       0.00
X1
```

## Python REPL

- You can also start the Python interpreter and compute interactively.
  - Useful for debugging your Python scripts

```
# repl_python()
# type your Python commands
# objects will be available in R through py object
# exit to quit
```