R & Python

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R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

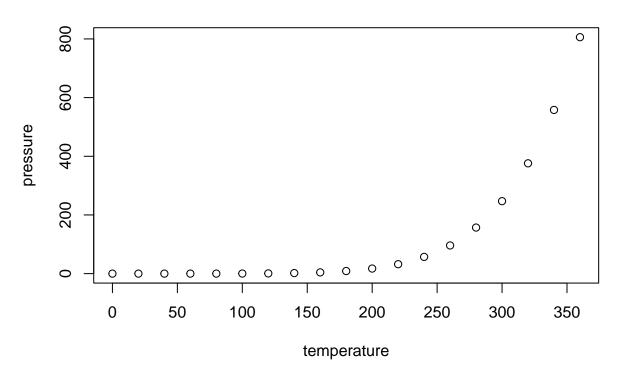
When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

summary(cars)

```
##
        speed
                          dist
##
           : 4.0
                    Min.
                            : 2.00
    Min.
    1st Qu.:12.0
                    1st Qu.: 26.00
##
##
    Median:15.0
                    Median: 36.00
##
    Mean
            :15.4
                    Mean
                            : 42.98
    3rd Qu.:19.0
                    3rd Qu.: 56.00
    Max.
            :25.0
                    Max.
                            :120.00
```

Including Plots

You can also embed plots, for example:



```
library(reticulate)
use_python("C:/Users/mlevi/anaconda3") #Ubicación de Python
#py_install("nombre_paquete_instalar") #Forma de instalar un paquete
os <- import("os") #Variable con la librería os de Python
os$listdir(".") #listo los archivos y directorios de mi ubicación actual
## [1] "01-EjemploRMD.html"
                               "01-EjemploRMD.pdf"
                                                       "01-EjemploRMD.Rmd"
                                                       "add.py"
## [4] "02-Documentacion.pdf"
                               "02-Documentacion.Rmd"
## [7] "prueba1RyPython.pdf"
                                                       "prueba1RyPython_files"
                               "prueba1RyPython.Rmd"
source_python("add.py") #Llama a un fichero de Python
add(10,2) #Llama a la función dentro del fichero
## [1] 12
```

#py_run_file("fichero.py") #Permite invocar al main de una clase

```
#Importar librería numpy y con el convert en FALSE indica que
#no se hagan conversiones automáticas
np <- import("numpy", convert = FALSE)
x <- np$array(c(1:4)) #Convierte un vector de 1 a 4 en array de Python
sum <- x$cumsum()
print(sum) #Imprime el array</pre>
```

```
## [ 1 3 6 10]
py_to_r(sum) #Convierte el array en un vector de R
## [1] 1 3 6 10
a <- np_array(c(1:10), dtype = "float16")</pre>
## [ 1. 2. 3. 4. 5. 6. 7. 8. 9. 10.]
b <- np_array(c(1:10), order = "C")</pre>
## [ 1 2 3 4 5 6 7 8 9 10]
datos <- iris #Dataframe de R (comienza en 1)
head(datos) #Mostramos los primeros datos
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1
              5.1
                         3.5
                                      1.4
                                                 0.2 setosa
## 2
              4.9
                         3.0
                                      1.4
                                                  0.2 setosa
## 3
             4.7
                        3.2
                                      1.3
                                                 0.2 setosa
## 4
              4.6
                         3.1
                                      1.5
                                                  0.2 setosa
## 5
             5.0
                         3.6
                                      1.4
                                                  0.2 setosa
## 6
                         3.9
             5.4
                                      1.7
                                                  0.4 setosa
datos_py <- r_to_py(datos) #Pasamos el dataframe a Python</pre>
import numpy as np
import pandas as pd
r.datos_py.head() #Mostramos los primeros datos (comienzan en 0)
      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 0
              5.1
                            3.5
                                         1.4
                                                      0.2 setosa
## 1
               4.9
                           3.0
                                         1.4
                                                      0.2 setosa
## 2
              4.7
                           3.2
                                                      0.2 setosa
                                         1.3
## 3
               4.6
                           3.1
                                         1.5
                                                      0.2 setosa
               5.0
                                                      0.2 setosa
## 4
                            3.6
                                         1.4
library(Matrix)
N <- 6
set.seed(123)
sparse_mat <- sparseMatrix(</pre>
i = sample(N, N, replace = F),
 j = sample(N, N, replace = F),
 x = runif(N),
 dims = c(N, N)
)
sparse_mat
```

```
## 6 x 6 sparse Matrix of class "dgCMatrix"
##
## [1,] .
                          0.8895393 .
## [2,] . 0.04205953 .
0.899825 .
                                            . 0.3279207
## [5,] 0.9545036 .
                         . 0.2460877 .
## [6,] .
sparse_mat_py <- r_to_py(sparse_mat)</pre>
r.sparse_mat_py
## <6x6 sparse matrix of type '<class 'numpy.float64'>'
## with 6 stored elements in Compressed Sparse Column format>
py_to_r(sparse_mat_py)
## 6 x 6 sparse Matrix of class "dgCMatrix"
##
## [1,] . . 0.8895393 .
## [2,] . 0.04205953 . . .
                          0.8895393 .
## [2,]
## [3,] .
                                            0.899825 .
                                           . 0.3279207
## [5,] 0.9545036 .
                      . . . 0.2460877 .
## [6,] . .
```

Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.