# Lecture 9: Intro to Python

## What is R?

- R is a programming language specifically designed for statistics and data analysis
  - Objects for storing data, and functions for interacting with data, are fundamental
  - R is very good at graphics and visualization
  - R is easily extended. Users can write and share their own functions and packages
- We can interact with R through IDEs like RStudio

## Who uses R?

- R is widely used in statistical research and academia
- R is also widely used in applications of statistics, such as
  - Biology and bioinformatics
  - Ecology, forestry, and environmental science
  - Psychology
  - Sociology
- R is used in a variety of areas outside research, by government agencies, pharmaceutical companies, insurance companies, etc.

# What other options exist?

- SAS
- Stata
- SPSS
- Excel
- Python
- Julia
- Matlab
- Many others...

## What is Python

- Python is a general-purpose programming language
- Like R, python has a wide range of packages to extend functionality
- Certain Python packages allow for sophisticated data analysis and modeling
  - SciPy, NumPy
  - scikit-learn, statsmodels, pytorch
  - pandas
  - matplotlib

## R vs. Python

My own, personal, preferences:

R is good for

- Data visualization and wrangling
- Classical statistics
- Statistical inference
- New statistical methods

Python is good for

- General-purpose programming
- Challenging data types (e.g. images)
- Prediction and machine learning

## Why do we teach R?

- Excellent support for the material covered in a statistics degree
- Benefits to using a consistent language across courses
- Valuable for a wide variety of future careers
- The primary research tool for most (if not all) the faculty

## A taste of Python

#### Recall our code from the first class:

```
1 M <- 10 # number of people at the party
2 hats <- 1:M # numbered hats
3 nsim <- 10000 # number of simulations
4 results <- rep(0, nsim) # vector to store the results
5
6 for(i in 1:nsim){
7    randomized_hats <- sample(hats, M, replace = FALSE)
8    results[i] <- sum(randomized_hats == hats) > 0
9 }
10
11 mean(results)
```

```
[1] 0.6296
```

## A taste of Python

Here is the same code, written in Python

```
import numpy as np

M = 10 # number of people at the party
hats = np.arange(M) # numbered hats
nsim = 10000 # number of simulations
results = np.zeros(nsim) # to store the results

for i in range(nsim):
    randomized_hats = np.random.choice(hats, M, replace = False)
    results[i] = np.sum(randomized_hats == hats) > 0

np.mean(results)
```

0.634

What similarities and differences do you notice?

## Step 1: representing the hats

```
1 import numpy as np
2
3 M = 10 # number of people at the party
4 hats = np.arange(M) # numbered hats
5
6 hats
array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
1 hats[0]
0
1 hats[1]
```

- hats is a 1-dimensional array (similar to a vector in R)
- Python is *O-indexed*: the first entry is hats [0]

## Step 2: everyone draws a random hat

```
import numpy as np

M = 10 # number of people at the party
hats = np.arange(M) # numbered hats

randomized_hats = np.random.choice(hats, M, replace = False)

randomized_hats
```

```
array([6, 8, 5, 2, 1, 0, 4, 7, 9, 3])
```

- np.random.choice works like R's sample function
- Booleans in Python are True and False (as opposed to TRUE and FALSE, or T and F)

## Step 3: check who got their original hat

```
import numpy as np
 3 M = 10 # number of people at the party
   hats = np.arange(M) # numbered hats
   randomized hats = np.random.choice(hats, M, replace = False)
   randomized hats
array([5, 6, 0, 3, 1, 8, 9, 7, 2, 4])
 1 randomized hats == hats
array([False, False, False, True, False, False, False, True, False,
      False])
 1 np.sum(randomized hats == hats)
```

NumPy arrays allow for "vectorized" operations, like in R

## Step 4: iteration

```
import numpy as np

M = 10 # number of people at the party
hats = np.arange(M) # numbered hats
nsim = 10000 # number of simulations
results = np.zeros(nsim) # to store the results

for i in range(nsim):
    randomized_hats = np.random.choice(hats, M, replace = False)
    results[i] = np.sum(randomized_hats == hats) > 0

np.mean(results)
```

0.6282

- range(nsim) is similar to 1:nsim in R
- We don't use the curly braces { }. Instead we use whitespace (four spaces is standard, you just have to be consistent)

## **Using Python through RStudio**

 You can make Python chunks in Quarto documents, just like R chunks:

```
1 ```{python}
2
3 ```
```

# **Class activity**

https://sta279-

f23.github.io/class\_activities/ca\_lecture\_9.html