BIOS-584 Python Programming (Non-Bios Student)

Week 05

Instructor: Tianwen Ma, Ph.D.

Department of Biostatistics and Bioinformatics,
Rollins School of Public Health,
Emory University

Lecture Overview

- Use random numbers and simulations to solve problems
- Draw random samples from different distributions
- Create subplots to compare different simulations
- Practice nested loops
- Write for loops with if/else statements
 - Simulate discrete random variables
- week-05-application-simulation.ipynb

Modifications of this slide

- For this lecture, I only give hints on the code.
- You write your own code to fill in the blanks.
- The slide is longer because I show them step-bystep.
- Pages with upyter are related to in-class coding.
- Raise hands if you have questions.
 - You can simply say "I don't understand this step."

Simulate random variables

- We use random numbers to simulate different distributions
- Previously we have learned random module to generate random numbers
- We can use functions within numpy module to simulate different distributions.
- np.random.distribution_name(parameters)
 - Normal, uniform, binomial, Chi-square, etc.
- We will start from continuous random variables.



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- Suppose we perform three simulations:
 - Normal: mean=1, variance=4
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 - Uniform: [-5, 5].
- Define each random vector as vec_normal, vec_chisq, and vec_uniform.



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- Print out all summary statistics in one sentence per distribution using .format() syntax.
- Avoid duplicating print functions by using a for loop.
 - Involve indexing the element of a list (and a nested list).

Multiple plots in a row (subplots)

- Previously, we learned how to create multiple plots in multiple figures.
- We can create multiple plots in the same figure using plt.subplots() function.
- This function returns a figure and an array of axes
 - fig, axes = plt.subplots(nrows, ncols)
 - You can change the output with other names
 - nrows: the number of rows of subplots
 - ncols: the number of columns of subplots
 - Within each axis, using the plot() function.



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- Display the plot and set other optional parameters

- We met range() function and np.arange() function to create a sequence of numbers.
- Let's introduce it formally!
- Syntax: list(range(start, stop, step)) to create a sequence of numbers.
 - start: The first number in the sequence
 - stop: The last number in the sequence (not included)
 - step: difference between adjacent numbers in the sequence.
 - If start is omitted, it defaults to 0.
 - If step is omitted, it defaults to 1.

- For range(), it only supports integers, while np.arange(), it supports floating numbers.
- The boundary setting is a little weird, but it follows the principle of half-open intervals (including the starting value but excludes the stop value).
- Since Python's index starts from 0, this design makes it easier to work with lists and arrays.
 - list[0:3] or list[:3] return the first three elements in the list.
 - list[1:] returns elements excluding the first one.



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- Find the first 10 elements of int_ls and even_ls.



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- Find the first 10 elements of int_ls and even_ls.
- Find the elements excluding the first two elements of int_ls and even_ls.

Nested Loops

- We can nest loops inside each other to iterate over multiple dimensions.
- The inner loop runs to completion for each iteration of the outer loop
- This is useful when we need to iterate over a matrix or a list of lists.
- The syntax is simple: Write a loop inside another loop

Nested Loops

- The inner loop is indented twice: once for the outer loop and once for the inner loop.
- Given i=0, j iterates 0,
 1, 2.
- Repeat for i=1,2,3.
- You can modify the code section.

```
for i in range(4):
    for j in range(3):
        print("i = {}, j = {}".format(i, j))

i = 0, j = 0
i = 0, j = 1
i = 0, j = 2
i = 1, j = 0
i = 1, j = 1
i = 1, j = 2
i = 2, j = 0
i = 2, j = 0
i = 2, j = 1
i = 2, j = 2
i = 3, j = 0
i = 3, j = 1
i = 3, j = 2
```

Central Limit Theorem (CLT)

- The CLT is a fundamental concept in statistics.
- It states that the distribution of the mean (or sum) of many independent, identically distributed random variables approaches to a normal distribution, regardless of the original distribution
- This is true even if the original distribution is NOT normal.

Real-world examples

- Quality control in manufacturing:
- A manager can take a random sample of products to assess the proportion of defective items produced.
- The CLT allowed them to use this sample to make an informed estimate of the defect rate for the entire plant.

Real-world examples

- Epidemiologists study how common illnesses or health behaviors are in populations using surveys.
- For example, to see how many adults in an area are obese, they select a random sub-population and measure people's height and weight and calculate the BMI.
- The CLT allowed them to use this sample to make an informed estimate of the obesity rate for the entire poupation.

Central Limit Theorem (CLT)

- The larger the sample size is, the closer the distribution of the sample mean is to a normal distribution.
- We use simulation studies to understand the central limit theorem.
- To simplify, let's simulate the CLT using the uniform distribution.

Logic Flow (before iterations)



- We pre-specify the number of iterations
 - iteration_num: The number of repetitions

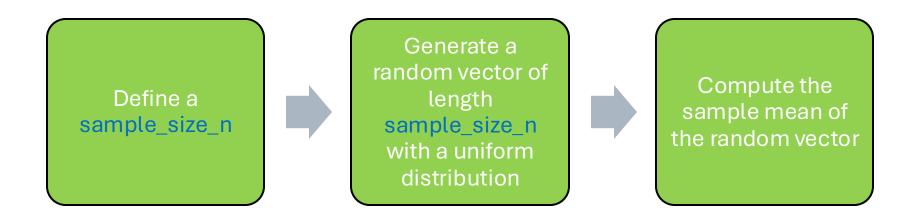
Logic Flow (before iterations)



- We pre-specify the number of iterations
 - iteration_num: The number of repetitions
- We initialize an empty list to save the sample mean
 - sample_size_n: within each repetition, the length of a random vector with a uniform distribution

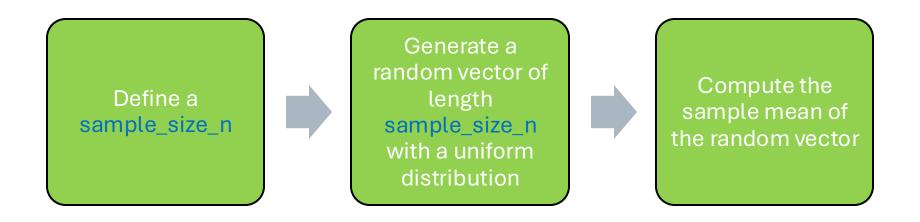
Logic Flow within each iteration





Logic Flow within each iteration





The above process is repeated iteration_num times. You append the sample mean to the vec_unif1 list.

Subplot after four scenarios



 Repeat the steps in the previous two slides to create unif_vec10, unif_vec50, and unif_vec100.

Subplot after four scenarios



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- Create a 2x2 subplot to display histograms
 - The first row: unif_vec1 (left) and unif_vec10 (right)
 - The second row: unif_vec50 (left) and unif_vec100 (right)

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- Create a 2x2 subplot to display histograms
 - The first row: unif_vec1 (left) and unif_vec10 (right)
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- Display the final subplot



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- The inner for loop is to create a random vector to store the sample mean of uniform vectors per sample size
- The outer for loop is to repeat for each sample size, i.e., 1, 10, 50, and 100.



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- Within each sample size, initialize an empty list unif_vec_sample_size_ls
- Repeat iteration_num times and for each iteration, generate a uniform vector and calculate the sample mean
- Append the sample mean to unif_vec_sample_size_ls



 After the completion of inner for loop, append unif_vec_sample_size_ls to unif_vec_ls_ls.



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- Display the four histograms using a 1x4 subplot
 - Left to right order: 1, 10, 50, and 100

for loop with if/else statements

- In addition to nested for loop, we can write if/else statement within the for loop.
- The syntax looks like:

```
for iter_id in total_ls:
    if condition1:
        statement1
    else:
        statement2
```

- We illustrate this function via simulating a discrete random variable.
 - We start with a binary variable.



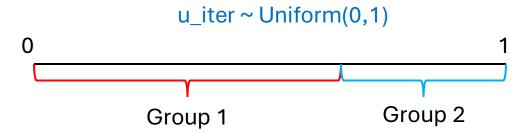
- Suppose we want to randomize a clinical trial of two treatment groups to a N_total patients with a 2:1 ratio
- How can we generate the treatment group id to make it random and approximately follow the 2:1 ratio?
- We define two treatment groups as 1 and 2 such that the number of group 1 is 2 times the number of group 2.



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 - We generate a u_iter from Uniform(0, 1).
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 - The < or <= does not make any difference.
- We replicate it for N_total times and store the treatment id using .append().
- Print out the actual numbers of patients who receive treatment 1 and 2 using .format().

- When N_total is moderate or small, the actual ratio may not be exactly 2:1.
- However, if you use bigger N_total, the actual ratio will approach to 2:1.

- Q: What if we have three treatment groups with a 2:2:1 ratio?
 - Expand your current one cutoff to two cutoffs.
- You will work on this in HW5.