

Workshop Resources

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1. Facilitator: <https://damayanthiherath.net>
2. Introduction to programming: Sections 1-3
3. Biological Data Analysis with Python
4. [Data Visualization More resources](#)
5. [The files for text processing exercise](#)
6. Powerpoint slides
7. Feedback Form

1. Algorithmic Thinking (Time: 10 Mins)

ILO: The participants will get experience in working out an algorithm for a given problem and speaking it out (while working as a group)

Divide the class into 4 by asking them to count to 1 to 4. Ask them to solve below problems and explain the steps

1. How to make a cup of tea?
2. How to organize a set of books in a library by alphabetical order? Imagine you have 10 books.
3. You are given a deck of cards with letters in an order. Reverse them'; Tell us how you did it?
4. You and your friends are planning a picnic. You need to pack a picnic basket that ensures the majority's preferences and needs are met while ensuring the basket doesn't exceed its weight limit. How will you do that?

Communicating Algorithms: paired Activity (20 mins):

ILO: The participants will get hands on writing a pseudocode/drawing a flowchart to present an algorithm for a given problem.

First 10 mins:

Write the pseudocode/flowchart to calculate the rainfall given the inputs, temperature and humidity

Next 10 mins

Complete the rest of it output whether a landslide would occur or not; A landslide will occur if rainfall is more than 200 mm

2, Python Basics

ILO: The participants will get hands on experience on Python environment

```
# This is a simple Python program to print Hello, World!  
print("Hello, World!")
```

Excercise: Display Hello with your name. What if instead of you, it is your friend or someone else and the name changes.?

Print the word 'Nenathambara' . Make the output seem underlined using the asterisk sign '*'. You can use double

quotes or triple quotes.

Type a short para about yourself using double quotes.

Type the same in point form(line after line) using triple quotes and using '\n'

You can only use one print command.Give a heading and underline your essay.

Lets get into our weather analyser. Let us write a program to calculate the rainfall given the temperature and humidity.

```
# Define temperature and humidity
temperature = 25.0 # Example temperature
humidity = 80 # Example humidity
```

Activity 4

ILO : The participants will get hands -on experience on implementing solutions involving Arithmetic, variables and types, Boolean variables and if else conditions, and loops.

Extend the code to calculate the rainfall; Print the value.

Extend the code to output whether there is a landslide risk or not given the rainfall value.

Can you print the numbers less than 500, divisible by 9?

```
>>> for i in range(500):
```

```
    if i%9 == 0:
```

```
        print(i)
```

Here's something a bit more tricky. Can you print the numbers less than 500, divisible by 5 and 7 and 11?

```
>>> for i in range (500):
```

```
if i%5==0:
```

```
    if i%7==0:
```

```
        if i%11 == 0:
```

```
            print(i)
```

Thinking expected- If you want to filter something that meets several requirements, use nested if statements.

*Introduce the word nested

Can you print the numbers less than 500, divisible by 4 and 6?

Can you keep printing the characters in a string until you either print all the characters or print the character “7” if it appears in the string?

```
string = input("Enter a string: ")
```

```
index = 0
```

```
while index < len(string):
```

```
    print(string[index])
```

```
    if string[index] == '7':
```

```
        break
```

```
    index += 1
```

4. You are building a basic landslide monitoring system that asks the user to input rainfall values in each loop. The system will:

- Stop and print an alert if **rainfall exceeds 200 mm**.
- Allow the user to stop the system manually by typing "stop."

Checking Multiple Conditions in a Sequence

```
seq =  
"MGSNKS KPKDASQRRRSLEPAENVHGAGGGAFPASQTPSKPASADGHRGPSAAFAPAAAE"
```

```
if 'GGG' in seq and 'RRR' in seq:
    print('GGG is at position:', seq.find('GGG'))
    print('RRR is at position:', seq.find('RRR'))

if 'WWW' in seq or 'AAA' in seq:
    print('Either WWW or AAA occur in the sequence')

if 'AAA' in seq and not 'PPP' in seq:
    print('AAA occurs in the sequence but not PPP')
```

Exercise

Define a sequence variable called `protein_seq` with a sample protein sequence that contains multiple amino acid motifs (e.g., 'MGGGKKKRRRAAAWWT').

Write code to:

- Check if both 'GGG' and 'KKK' are present in the sequence, and print their positions.
- Print a message if either 'WWW' or 'SSS' is found.
- Check if 'AAA' exists in the sequence but not 'YYY', and print a message if true.

3. Manipulating Text

Read from a text file

`readlines()` reads the file line-by-line into a list.

`read()` reads the entire file as a single string.

```
# Using readlines() to read all lines into a list
text_file = open('text_file.txt', 'r')
lines = text_file.readlines()
text_file.close()
print(lines) # use a for loop to print each line one by one

# Using read() to read the entire file as one string
text_file = open('text_file.txt')
print(text_file.read())
text_file.close()
```

Exercise 01

Write a program to open and read the contents of the given file called `neuron_data.txt`. Print each line of the file individually, and then close the file. You can use `readlines()` and then use a for loop to print each line.

Writing to a Text File

Use 'w' mode to create a new file or overwrite an existing one.

```
output_file = open('new_file.txt', 'w')
output_file.write('sample text to be written\n')
output_file.close()
```

Exercise 02

Write a program to create and open a new file called “microbe_counts.txt” in write mode. Write a line that says "Total microbe count: 15", then close the file. Reopen the file to read and print the line you wrote.

Cleaning Text Data

The `strip()` method removes extra spaces. You can also use `rstrip()` to remove spaces on the right side only or `lstrip()` on the left.

```
output_file = open('file.txt')
print(output_file.read().strip())
output_file.close()
```

Exercise 03

First create a text file named “friends.txt” and then add some data with white spaces in the beginning of the sentence. Then write a program to open and read the contents of “data.txt”. Use `strip()` to remove any leading or trailing whitespace, then print the cleaned line.

Basics of Formatting

Specifier	Explanation	Example Output
:d	Formats an integer.	<code>f'{42:d}'</code> → 42
:f	Formats a float (default 6 decimal places)	<code>f'{3.14159:f}'</code> → 3.141590
:10.2f	Right-align, width=10, 2 decimal places	<code>f'{3.1:10.2f}'</code> → 3.10
:<10.2f	Left-align, width=10, 2 decimal places	<code>f'{3.1:<10.2f}'</code> → 3.10
:+10.2f	Include a sign (+ or -)	<code>f'{3.1:+10.2f}'</code> → +3.10

:010.2f	Pad with zeros (width=10, 2 decimals)	f"{3.1:010.2f}" → 0000003.10
:.0f	Display as a whole number (rounded)	f"{3.6:.0f}" → 4

```
# Integer formatting
num = 42
print(f"{num:d}")      # Output: 42 (integer)
print(f"{num:5d}")     # Output: '  42' (right-aligned, width=5)
print(f"{num:<5d}")     # Output: '42   ' (left-aligned, width=5)

# Float formatting
pi = 3.14159
print(f"{pi:f}")       # Output: 3.141590 (default float)
print(f"{pi:.2f}")     # Output: 3.14 (2 decimal places)
print(f"{pi:10.2f}")   # Output: '      3.14' (width=10, 2 decimals)
print(f"{pi:<10.2f}")  # Output: '3.14      ' (left-aligned)

# Including signs
neg = -3.14
print(f"{pi:+.2f}")    # Output: +3.14 (positive sign)
print(f"{neg:+.2f}")   # Output: -3.14 (negative sign)

# Zero padding
print(f"{pi:010.2f}")  # Output: 0000003.14 (width=10, padded with zeros)

# Rounded floats
large_num = 12345.6789
print(f"{large_num:.0f}") # Output: 12346 (rounded to nearest whole number)
```

Exercise 04

Format the following values as described:

Print an integer 56 with:

- Right-alignment in a field of width 8.
- Left-alignment in a field of width 8.

Print a float 45.678 with:

- Exactly 2 decimal places.
- Right-aligned in a width of 10, with 2 decimal places.
- Zero-padded, width of 8, and rounded to 1 decimal place.

Calculate the average from a list of numbers

```
# calculate average from float numbers
data = [3.53, 3.47, 3.51, 3.72, 3.43]
average = sum(data) / len(data)
print average

# calculate average from integer numbers
data = [1, 2, 3, 4]
average = float(sum(data)) / len(data)
print average
```

Exercise 05

Print the average of [4.56,8.55,9.25,10.32,23.91]

*****Find items common to two lists*****

```
# Reading lines from both files
lines1 = open("file1.txt").readlines()
lines2 = open("file2.txt").readlines()

# Strip newlines and spaces
lines1 = [line.strip() for line in lines1]
lines2 = [line.strip() for line in lines2]

# Comparing the two lists
for line in lines1:
    if line in lines2:
        print(line, 'same one detected')
    else:
        print(line, 'not detected')
```

Exercise 06

Use the given files “cell_cycle_proteins.txt” and “cancer_cell_proteins.txt” that contain gene IDs.

Write code to print each gene ID from cell_cycle_proteins, indicating if it is also present in cancer_cell_proteins.

Sets

Sets are unordered collections of unique elements. They are useful for removing duplicates, finding intersections, unions, and differences between groups of objects.

```
s1 = set('LDFGJLDFGDGD')
print(s1) # Output: {'J', 'F', 'L', 'G', 'D'}

# Check membership
print('L' in s1) # True
print('Z' not in s1) # True
```

With sets, finding the intersection becomes straightforward:

```
data_a = {1, 2, 3, 4, 5, 6}
data_b = {1, 5, 7, 8, 9}

a_and_b = data_a.intersection(data_b)
print(a_and_b)
```

Exercise 07

Define two sets {21, 11, 2, 64, 35, 60 } and {81, 11, 52, 94, 23, 59 } and print their intersection.

Find Common Elements in Multiple Sets

Using `reduce()` to find common elements in three sets:

```
from functools import reduce

a = {1, 2, 3, 4, 5}
b = {2, 4, 6, 7, 1}
c = {1, 4, 5, 9}

common = reduce(set.intersection, [a, b, c])
print(common)
```

Exercise 08

Define four sets and then find the common elements among them.

Removing Elements from a List

Use `pop()`, `del`, and `remove()` to remove elements:

```
data_a = [1, 2, 3, 4, 5, 6, 7]

data_a.pop() # Removes and returns last element
del data_a[1] # Deletes element at index 1
data_a.remove(3) # Removes first occurrence of the value 3
```

Slicing a List

```
data_a = [1, 2, 3, 4, 5, 6]
print(data_a[:2]) # Output: [1, 2]
```

Exercise 09

Consider the list [23, 55, 67, 94, 35, 77, 12, 74]. Slice it from '67' and create a new list and then pop the last element.

Remove Duplicates from a File

Exercise 10

Open the given file "UniprotID.txt" Then read the content and use a set to remove duplicates from the file.

Reading Tabular Data from a File

Tabular data can be read from a text file and processed into a Python list of lists. This structure makes it easier to manipulate rows and columns.

```
# Read tabular data from a tab-separated text file
table = []

for line in open('lowry_data.txt'):
    table.append(line.strip().split('\t'))

print(table)
```

Exercise 11

Create a text file named `data.txt` with the following content:

Use tab spaces...

```
name age country
Alice 30 USA
Bob 25 UK
Charlie 35 Canada
```

Write a Python script to read the file and print its contents as a list of lists.

****Basic Sorting with sorted() and sort()****

Python provides two main ways to sort data:

- `sort()`: A method that modifies lists in-place
- `sorted()`: A built-in function that works with any iterable and returns a new sorted object

```
# Using sort() method
numbers = [3, 1, 4, 1, 5, 9, 2, 6]
numbers.sort()
print("Sorted list:", numbers)

# Using sorted() function
original = [3, 1, 4, 1, 5, 9, 2, 6]
new_sorted = sorted(original)
print("Original list:", original)
print("New sorted list:", new_sorted)
```

Reverse Sorting

Both `sort()` and `sorted()` accept a `reverse` parameter to sort in descending order.

```
numbers = [1, 5, 2, 8, 3, 9, 7]

# Descending order with sorted()
desc_numbers = sorted(numbers, reverse=True)
print("Descending order:", desc_numbers)
```

Exercise 12

Create two lists:

1. Use `sort()` to sort a list of your favorite numbers in ascending order
2. Use `sort()` to sort a list of your favorite numbers in descending order
3. Use `sorted()` to create a new sorted list from a list of fruit names

- Hint: Try ['banana', 'apple', 'orange', 'kiwi']

Basic Pattern Matching with Regular Expressions

Regular expressions (regex) are powerful tools for pattern matching in text. The `re` module in Python provides methods to work with regex patterns

```
import re

# String to search in
sequence = 'VSVLT MFRYAGWLDRLYMLVGTQLAAIHGVALPLMMLI'

# Compile and search for pattern [ST]Q (S or T followed by Q)
pattern = re.compile('[ST]Q')
match = pattern.search(sequence)

if match:
    # Print 4 characters before and after the match
    print(sequence[match.start() - 4:match.end() + 4])
    print(match.group())
```

Exercise 13

Create a script that searches for the pattern `[AG]T` (A or G followed by T) in the sequence:
`ATGATCGTAGTCGATGCTAGCTAGCTAGT`

Print the match and 2 characters before and after each match.

Finding Multiple Matches

The `re` module provides several methods for finding multiple matches:

- `findall()`: Returns a list of all matches
- `group()`: Returns the matched string
- `span()`: Returns start and end positions
- `start()`: Returns starting position
- `end()`: Returns ending position

```
import re

sequence = 'RQSAMGSNKS KPKDASQRRRSLEPAENVHGAGGGAFPASQRPSKP'
pattern = re.compile('R.[ST][^P]') # R followed by any char, then S or T, then not P

# Using findall
matches = pattern.findall(sequence)
print("All matches:", matches)
```

Exercise 14

Write a script to find all occurrences of the pattern `[AG]{2}[CT]` (two A's or G's followed by C or T) in the sequence: `AAGCTAAGTCGAGGCTTAGCTAGGC`

Print both the matches.

Debugging:

Here's a code that someone learning coding typed but doesn't run. Can you figure out what they wanted to do with this code? Can you fix the code?

Sample code (erroneous):

```
price=20
money=30
if price<=money:
print("can buy")
else:
print("can not buy")
```

Exercise:

You left your computer for a moment and a grade 1 kid came along and scrambled everything. Now you've forgotten what you were trying to do as well. Can you make sense of the codes here and get a working algorithm?

Sample code (erroneous):

```
length=20
height=40
```

```
print("length>=height")
if length>=height:print("length<height")
else:
```

Data Visualization with Python

Matplotlib is a Python package, which provides a wide variety of plot types such as lines, bars, pie charts, and histograms.

Seaborn is a Python visualization library based on Matplotlib and provides a high-level interface for drawing attractive statistical graphics.

matplotlib.pyplot provides an implicit, MATLAB-like, way of plotting. It also opens figures on your screen, and acts as the figure GUI manager.

Line plot

```
from matplotlib.pyplot import figure, plot, savefig

xdata = [1, 2, 3, 4]
ydata = [1.25, 2.5, 5.0, 10.0]

figure()
plot(xdata, ydata)
```

https://matplotlib.org/2.1.1/api/_as_gen/matplotlib.pyplot.plot.html

```

import math
from matplotlib.pyplot import figure, plot, text, axis
figure()

xdata = [0.1 * i for i in range(100)]
ydata = [math.sin(j) for j in xdata]

plot(xdata, ydata, 'kd', linewidth = 1)
text(4.8, 0, "$y = \sin(x)$", horizontalalignment = 'center', fontsize = 20)
axis([0, 3 * math.pi, -1.2, 1.2])

```

Histograms

```

from pickle import TRUE
from matplotlib.pyplot import plot, figure, title, xlabel, ylabel, hist, axis, grid

data = [1, 1, 9, 1, 3, 5, 8, 2, 1, 5, 11, 8, 3, 4, 2, 5]
n_bins = 5

figure()
num, bins, patches = hist(data, n_bins, density = TRUE, histtype = 'bar', facecolor = 'green', alpha = 0.75)

title('Histogram')
xlabel('value')
ylabel('frequency')
axis()
grid(True)

```

Bar plots

```
from matplotlib.pyplot import figure, title, xlabel, ylabel, xticks, bar, legend, axis

nucleotides = ["A", "G", "C", "U"]

counts = [    [606, 1024, 759, 398],
           [762, 912, 639, 591],
           ]

figure()
title('RNA nucleotides in the ribosome')
xlabel('RNA')
ylabel('base count')

x1 = [2.0, 4.0, 6.0, 8.0]
x2 = [x - 0.5 for x in x1]

xticks(x1, nucleotides)

bar(x1, counts[1], width = 0.5, color = "#cccccc", label = "E.coli 23S")
bar(x2, counts[0], width = 0.5, color = "#808080", label = "T.thermophilus 23S")

legend()
axis([1.0, 9.0, 0, 1200])
```

```
# scatterplot with error bars
x1 = [1.1, 1.2, 1.3, 1.4, 1.5]
y1 = [10, 15, 10, 15, 17]
err1 = (2, 3, 4, 1, 2)
width = 0.05
bar(x1, y1,width, color = 'r', yerr = err1, ecolor = "black")
```

boxplots

```
from matplotlib.pyplot import plot, boxplot
import matplotlib.pyplot as plt
import numpy as np

# Creating dataset
np.random.seed(10)
data = np.random.normal(100, 20, 200)
# Creating plot
boxplot(data)
```

Resources:

1. <https://docs.python.org/3/tutorial/index.html>
2. Lessons 3,4,6 of

https://www.youtube.com/playlist?list=PLqAsWscG_yDP6U_dJWhuSuQWMvv2XbJ9W

3. Matplotlib tutorials:

<https://matplotlib.org/stable/tutorials/index.html>