PROGRAM STUDI INFORMATIKA FAKULTAS TEKNIK DAN INFORMATIKA UNIVERSITAS MULTIMEDIA NUSANTARA SEMESTER GENAP TAHUN AJARAN 2021/2022



IF420 – ANALISIS NUMERIK

Pertemuan ke 3 – More on Python

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Capaian Pembelajaran Mingguan Mata Kuliah (Sub-CPMK):



Sub-CPMK 3: Mahasiswa mampu menerapkan dan memanfaatkan beberapa fitur lanjutan di Python – C3





- Recursion
- Object Oriented Programming (OOP)
- Complexity
- Representation of Numbers
- Errors, Good Programming Practices, and Debugging



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- Reading and Writing Data
- Visualization and Plotting
- Parallel Your Python

Reading and Writing Data



- Storing data and the results of your programming efforts is important for working over multiple sessions and sharing your results with collaborators.
- Since when Python closes, all the variables in the memory are lost, data must be stored
 in some other way.
- Sometimes data must also be readable by or written in a form that can be read by other programs.
- We will discuss several means in storing data, such as TXT files, CSV files, Pickle files, JSON files, and HDF5 files.





- A **text file**, many times with an extension **.txt**, is a file containing only **plain text**. However, programs you write and programs that read your text file will usually expect the text file to be in a certain format; that is, organized in a specific way.
- To work with text files, we need to use open function which returns a file object. It is commonly used with two arguments.

f = open(filename, mode)

Example: Create a text file called test.txt and write a couple lines in it.

```
In [1]: M f = open('test.txt', 'w')
for i in range(5):
    f.write(f"This is line {i}\n")
f.close()
```

```
File Edit Format View Help

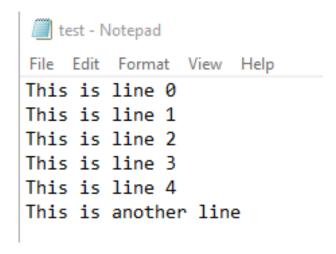
This is line 0
This is line 1
This is line 2
This is line 3
This is line 4
```





Now, let's append some string to the test.txt file. It is very similar to how we write the
file, with only one difference - change the mode to 'a' instead.

```
In [2]: M f = open('test.txt', 'a')
f.write(f"This is another line\n")
f.close()
```



TXT Files



We could read a file from disk and store all the contents to a variable.

```
In [3]: M f = open('./test.txt', 'r')
                                             In [4]: M type(content)
           content = f.read()
           f.close()
                                                Out[4]: str
           print(content)
           This is line 0
           This is line 1
                                         But sometimes we want to read in the contents in
           This is line 2
                                          the files line by line and store it in a list. We could
           This is line 3
           This is line 4
                                          use f.readlines() to achieve this.
           This is another line
In [5]: M f = open('./test.txt', 'r')
                                                                    In [6]: M type(contents)
            contents = f.readlines()
                                                                       Out[6]: list
            f.close()
            print(contents)
            ['This is line 0\n', 'This is line 1\n', 'This is line 2\n', 'This is line
            3\n', 'This is line 4\n', 'This is another line\n']
```

Dealing with numbers and arrays



- Since we are working with numerical methods later, and many times, we work with the numbers or arrays. We could use the above methods to save the numbers or arrays to a file and read it back to the memory. But it is not so convenient this way.
- Instead, commonly we use the numpy package to directly save/read an array.

```
my_arr - Notepad

File Edit Format View Help

# Col1 Col2 Col3
1.20 2.20 3.00
4.14 5.65 6.42
```

We can see, read in the file directly to an array is very simple using the np.loadtxt function and it skips the first header as well.





- There are many scientific data which are stored in the comma-separated values (CSV)
 file format, a delimited text file that uses a comma to separate values.
- It is a very useful format that can store large tables of data (numbers and text) in plain text.
- Each line (row) in the data is **one** data **record**, and each record consists of **one or more fields**, separated by **commas**.
- It also can be opened using Microsoft Excel, and visualize the rows and columns.





We use the np.savetxt function to save the data to a csv file.

```
In [11]: M import numpy as np

data = np.random.random((100,5))

np.savetxt('test.csv', data, fmt = '%.2f', delimiter=',', header = 'c1, c2, c3, c4, c5')
```

A	Α	В	С
1	# c1, c2, c3, c4, c5		
2	0.94,0.71,0.55,1.00,0.51		
3	0.70,1.00,0	0.68,0.13,0.	.92
4	0.89,0.00,0	0.33,0.28,0.	.72
5	0.67,0.22,0	0.16,0.73,0.	.24
6	0.78,0.49,0	0.91,0.43,0.	43
7	0.06,0.27,0	0.27,0.39,0.	.88
8	0.66,0.65,0	0.38,0.45,0.	.69
9	0.17,0.45,0	0.73,0.88,0.	.01
10	0.12,0.22,0	0.59,0.38,0.	.59
11	0.87,0.64,0	0.56,0.97,0.	.67

We could read in the csv file using the np.loadtxt function.

Pickle Files



- In this section, we introduce another way to store the data to the disk pickle.
- We talked about saving data into text file or csv file. But in certain cases, we want to store dictionaries, tuples, lists, or any other data type to the disk and use them later or send them to some colleagues.
- Pickle can be used to serialize Python object structures, which refers to the process of converting an object in the memory to a byte stream that can be stored as a binary file on disk.
- When we load it back to a Python program, this binary file can be de-serialized back to a Python object.
- To use a pickle, we need to import the module first.

Pickle Files



- To use pickle to serialize an object, we use the pickle.dump function, which takes two
 arguments: the first one is the object, and the second argument is a file object returned
 by the open function.
- Note here the mode of the open function is 'wb' which indicates write binary file.

```
In [13]: M import pickle
In [14]: M dict_a = {'A':0, 'B':1, 'C':2}
    pickle.dump(dict_a, open('test.pkl', 'wb'))
```

• Now let's load the pickle file we just saved on the disk back using the pickle.load function.

```
In [15]: | my_dict = pickle.load(open('./test.pkl', 'rb'))
    my_dict
Out[15]: {'A': 0, 'B': 1, 'C': 2}
```





- JSON stands for JavaScript Object Notation.
- A JSON file usually ends with extension ".json".
- Unlike pickle, which is Python dependent, JSON is a language-independent data format, which makes it attractive to use.
- Besides, it is usually takes less space on the disk and the manipulation is faster than
 pickle. Therefore, it is a good option to store your data using JSON.
- The text in JSON is done through quoted string containing value in key-value pairs within {}. It is actually very similar to the dictionary we saw in Python.

"state": "California",

"postal": "94720"

"student 1",
"student 2",

"student 3"

"array":[1, 2, 3]

},

"list":[

],

```
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```

- In Python, the easiest way to handle JSON is to use **json library**. There are some other libraries such as **simplejson**, **jyson**, etc. Here, we will use **json** which is natively supported by Python to write and load JSON files.
- To use json to serialize an object, we use the json.dump function.
- To load the JSON file, we use json.load function.

HDF5 Files



- In scientific computing, sometimes, we need to store large amounts of data with quick access, the file formats we introduced before are not going to cut it.
- You will soon find there are many cases, HDF5 (Hierarchical Data Format) is the solution. It is a powerful binary data format with no limit on the file size. It provides parallel IO (input/output), and carries out a bunch of low level optimizations under the hood to make the queries faster and storage requirements smaller.
- An HDF5 file saves two types of objects: datasets, which are array-like collections of data (like NumPy arrays), and groups, which are folder-like containers that hold datasets and other groups.
- The so called **hierarchical** in HDF5 refers to the fact that the data could be saved like a file system, with **folder-like structures**, such as folder, subfolder (in HDF5, it is called group, subgroup). Groups operate like dictionaries with the keys and values, with the **keys** are **names** of the groups, and the **values** are the **subgroups** or **datasets**.

HDF5 Files



- In order to use read/write HDF5 in Python, there are some packages or wrappers to serve the purposes. The most common two packages are **PyTables** and **h5py**.
- We will only introduce the **h5py** here. You can install h5py using **conda** (hope you still remember how to do that, if you forget, please go back to Chapter 1).
- Example: Suppose we deployed some instruments to monitor the accelerations and GPS location in Bay Area, CA. We deployed two accelerometers at Berkeley and Oakland as well as one GPS station at San Fransisco. And they record data at different sampling rates, with the accelerometer at Berkeley sample the data every 0.04 s, and 0.01 s for the sensor at Oakland. The GPS samples the location every 60 seconds in San Fransisco. Now we want to store the two types of data into a HDF5 as well as some attributes indicate where the data is recorded, start time of the recording, station name and the sampling interval.

```
In [19]: M import numpy as np
             import h5py

    # Generate random data for recording

In [20]:
             acc 1 = np.random.random(1000)
             station_number 1 = '1'
             # unix timestamp
             start time 1 = 1542000276
             # time interval for recording
             dt 1 = 0.04
                                                 In [21]: M | hf = h5py.File('station.hdf5', 'w')
             location 1 = 'Berkeley'
             acc 2 = np.random.random(500)
             station number 2 = '2'
             start time 2 = 1542000576
             dt 2 = 0.01
             location 2 = 'Oakland'
```



```
In [22]: M | hf['/acc/1/data'] = acc 1
             hf['/acc/1/data'].attrs['dt'] = dt 1
             hf['/acc/1/data'].attrs['start time'] = start time 1
             hf['/acc/1/data'].attrs['location'] = location 1
             hf['/acc/2/data'] = acc 2
             hf['/acc/2/data'].attrs['dt'] = dt 2
             hf['/acc/2/data'].attrs['start time'] = start time 2
             hf['/acc/2/data'].attrs['location'] = location 2
             hf['/gps/1/data'] = np.random.random(100)
             hf['/gps/1/data'].attrs['dt'] = 60
             hf['/gps/1/data'].attrs['start time'] = 1542000000
             hf['/gps/1/data'].attrs['location'] = 'San Francisco'
```

Read in the HDF5 files

```
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```

We can see reading a HDF5 is also easy with h5py. After we read in the HDF5 to hf_in, we could see what groups are in the HDF5 using the keys function.

Out[29]: array([0.47186583, 0.47308128, 0.46795467, 0.01968574, 0.94769089,

0.49005002, 0.7099151 , 0.76995257, 0.89032606, 0.23462921])

```
• Then we could get access to the group In [30]: M list(data_1.attrs)

members and see what contains in the subgroups as the hf_in['acc'], or In [31]: M data_1.attrs['dt']

directly specify the path to the datasets as hf_in['acc/1/data'] and get_In [32]: M data_1.attrs['location']

the array data.

Out[32]: 'Berkeley'
```

Visualization and Plotting



- Visualizing data is usually the best way to convey important engineering and science ideas and information, especially if the information is made up of many, many numbers.
- The ability to visualize and plot data quickly and in many different ways is one of Python's most powerful features.
- Python has numerous graphics functions that enable you to efficiently display plots, surfaces, volumes, vector fields, histograms, animations, and many other data plots.

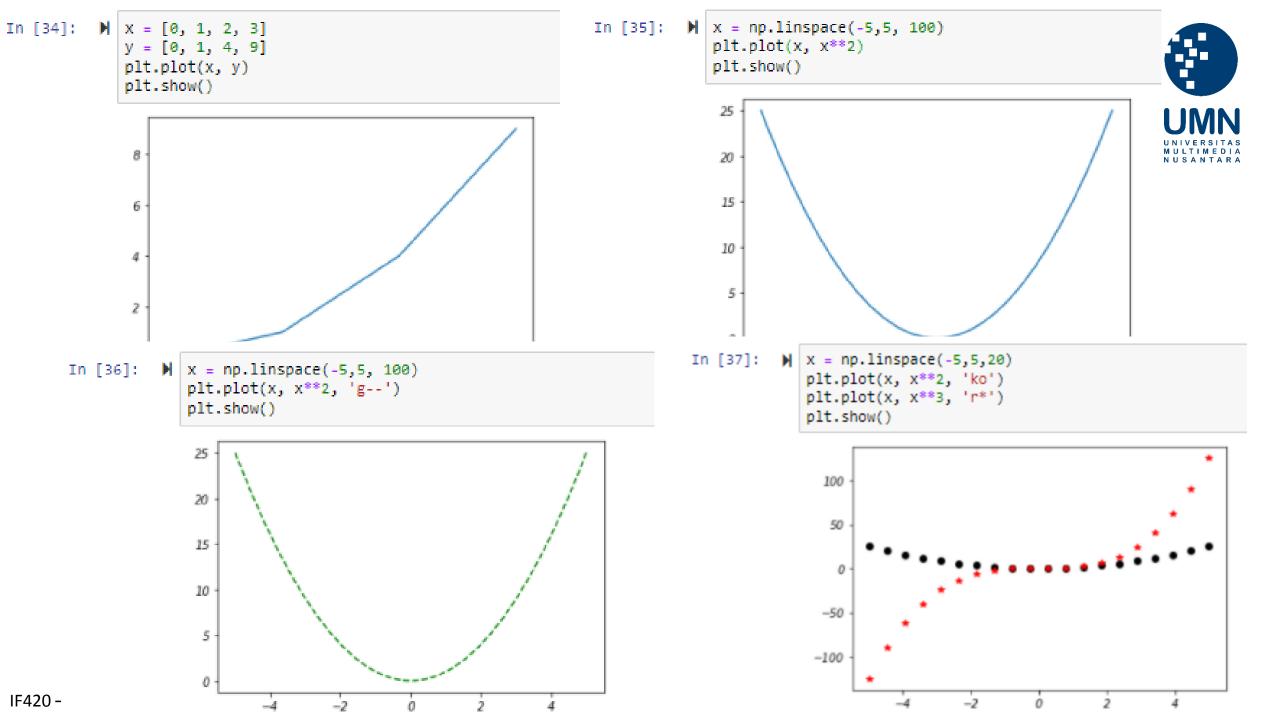
2P Plotting



- In Python, the matplotlib is the most important package that to make a plot, you can have a look of the matplotlib gallery and get a sense of what could be done there.
- Usually the first thing we need to do to make a plot is to import the matplotlib package.
- In Jupyter notebook, we could show the figure directly within the notebook and also have the interactive operations like pan, zoom in/out, and so on using the magic command %matplotlib notebook.

```
In [33]: M import numpy as np
import matplotlib.pyplot as plt
%matplotlib notebook
```

• The basic plotting function is plot(x,y). The plot function takes in two lists/arrays, x and y, and produces a visual display of the respective points in x and y.



```
In [38]: M plt.figure(figsize = (10,6))

x = np.linspace(-5,5,20)
plt.plot(x, x**2, 'ko')
plt.plot(x, x**3, 'r*')
plt.title(f'Plot of Various Polynomials from {x[0]} to {x[-1]}')
plt.xlabel('x axis', fontsize = 18)
plt.ylabel('Y axis', fontsize = 18)
plt.show()
```



Plot of Various Polynomials from -5.0 to 5.0

- We can see that we could change any part
 of the figure, such as the x and y axis label
 size by specify a fontsize argument in the
 plt.xlabel function.
- But there are some pre-defined styles that we could use to automatically change the style. Here is the list of the styles.

```
50
axis
   -100
                                                X axis
```

2P Plotting



X axis

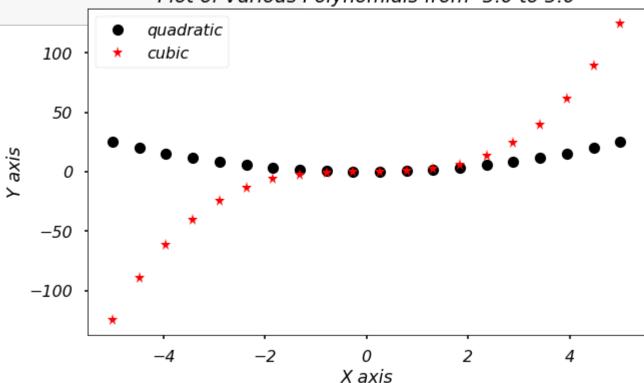
```
M plt.style.use('seaborn-poster')
In [40]:
          plt.figure(figsize = (10,6))
In [41]:
             x = np.linspace(-5,5,20)
             plt.plot(x, x**2, 'ko')
             plt.plot(x, x**3, 'r*')
             plt.title(f'Plot of Various Polynomials from \{x[0]\}\ to \{x[-1]\}')
                                                                                 Plot of Various Polynomials from -5.0 to 5.0
             plt.xlabel('x axis')
             plt.ylabel('Y axis')
             plt.show()
                                                                    100
                                                                     50
                                                                Y axis
                                                                       0
                                                                    -50
                                                                   -100
                                                                                           -2
```

```
In [42]: | plt.figure(figsize = (10,6))

x = np.linspace(-5,5,20)
plt.plot(x, x**2, 'ko', label = 'quadratic')
plt.plot(x, x**3, 'r*', label = 'cubic')
plt.title(f'Plot of Various Polynomials from {x[0]} to {x[-1]}')
plt.xlabel('X axis')
plt.ylabel('Y axis')
plt.legend(loc = 2)
plt.show()
Plot of Various Polynomials from -5.0 to 5.0
```



You can add a legend to your plot by using the legend function and add a label argument in the plot function. The legend function also takes argument of loc to indicate where to put the legend, try to change it from 0 to 10.



```
In [43]: M plt.figure(figsize = (10,6))

x = np.linspace(-5,5,100)
plt.plot(x, x**2, 'ko', label = 'quadratic')
plt.plot(x, x**3, 'r*', label = 'cubic')
plt.title(f'Plot of Various Polynomials from {x[0]} to {x[-1]}')
plt.xlabel('x axis')
plt.ylabel('Y axis')
```



 Finally, you can further customize the appearance of your plot to change the limits of each axis using the xlim or ylim function.

plt.legend(loc = 2)

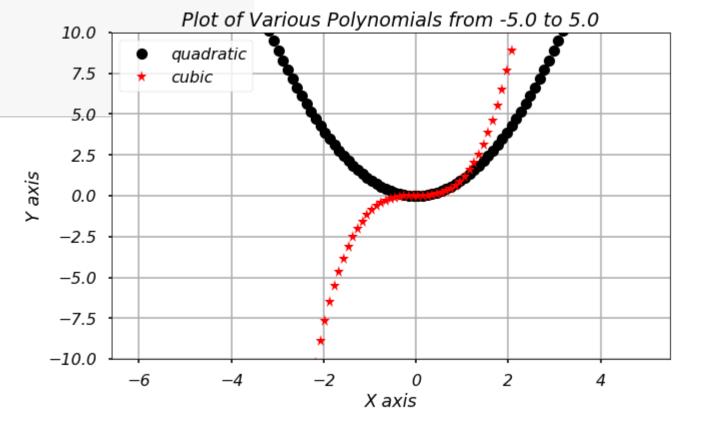
plt.xlim(-6.6)

plt.grid()

plt.show()

plt.ylim(-10,10)

 Also, you can use the grid function to turn on the grid of the figure.



2P Plotting



- We can create a table of plots on a single figure using the subplot function.
- The subplot function takes three inputs: the number of rows of plots, the number of columns of plots, and to which plot all calls to plotting functions should plot.
- You can move to a different subplot by calling the subplot again with a different entry for the plot location.
- There are several other plotting functions that plot x versus y data. Some of them are scatter, bar, loglog, semilogx, and semilogy.

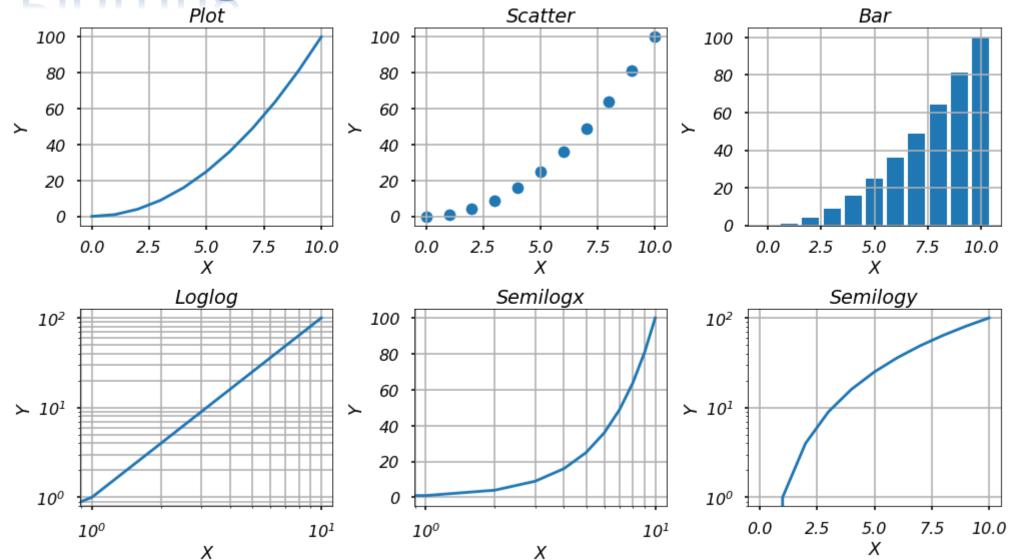
```
In [44]: \mathbf{M} \mid \mathbf{x} = \text{np.arange}(11)
               V = X^{**}2
               plt.figure(figsize = (14, 8))
               plt.subplot(2, 3, 1)
              plt.plot(x,y)
              plt.title('Plot')
              plt.xlabel('X')
               plt.ylabel('Y')
              plt.grid()
               plt.subplot(2, 3, 2)
               plt.scatter(x,y)
               plt.title('Scatter')
               plt.xlabel('X')
              plt.ylabel('Y')
               plt.grid()
               plt.subplot(2, 3, 3)
              plt.bar(x,y)
               plt.title('Bar')
               plt.xlabel('X')
               plt.ylabel('Y')
               plt.grid()
```

```
plt.subplot(2, 3, 4)
plt.loglog(x,y)
plt.title('Loglog')
plt.xlabel('X')
plt.ylabel('Y')
plt.grid(which='both')
plt.subplot(2, 3, 5)
plt.semilogx(x,y)
plt.title('Semilogx')
plt.xlabel('X')
plt.ylabel('Y')
plt.grid(which='both')
plt.subplot(2, 3, 6)
plt.semilogy(x,y)
plt.title('Semilogy')
plt.xlabel('X')
plt.vlabel('Y')
plt.grid()
plt.tight layout()
plt.show()
```



2D Plotting

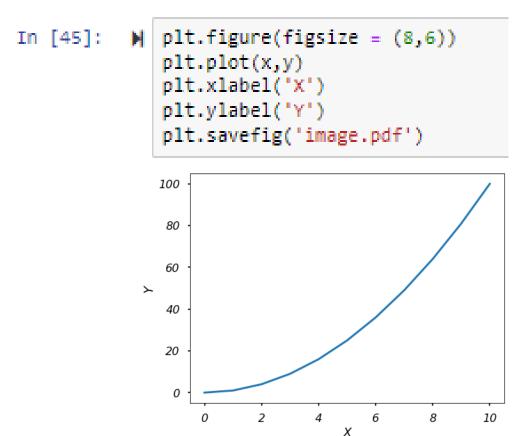


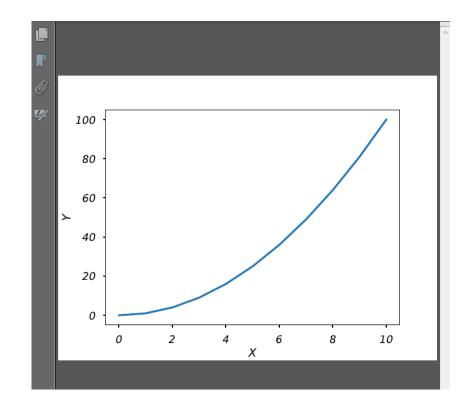






 Sometimes, you want to save the figures as a specific format, such as pdf, jpeg, png, and so on. You can do this with the function plt.savefig.





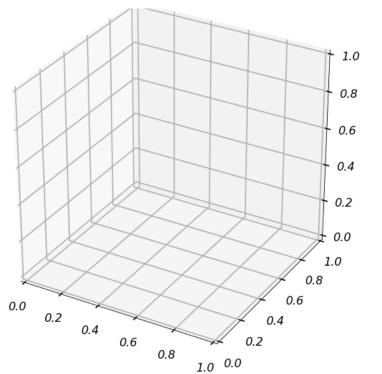
3P Plotting



 In order to plot 3D figures using matplotlib, we need to import the mplot3d toolkit, which adds the simple 3D plotting capabilities to matplotlib.

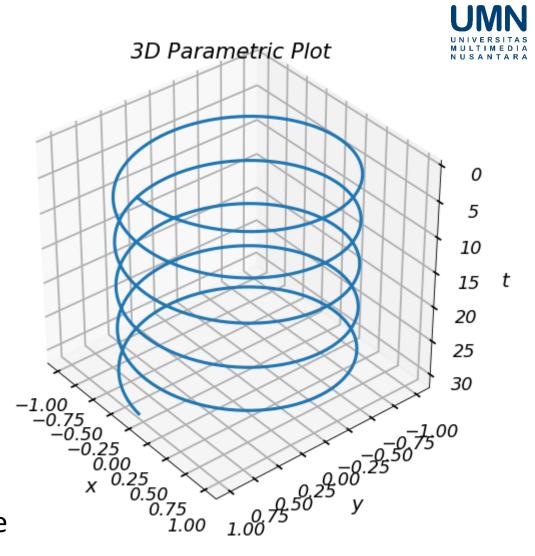
```
In [46]: M import numpy as np
    from mpl_toolkits import mplot3d
    import matplotlib.pyplot as plt
    plt.stvle.use('seaborn-poster')
In [47]: M fig = plt.figure(figsize = (10,10))
ax = plt.axes(projection='3d')
plt.show()
```

- Once we imported the mplot3d toolkit, we could create 3D axes and add data to the axes. Let's first create a 3D axes.
- The ax = plt.axes(projection='3d') created a 3D axes object, and to add data to it, we could use plot3D function. We could change the title, set the x, y, z labels for the plot as well.



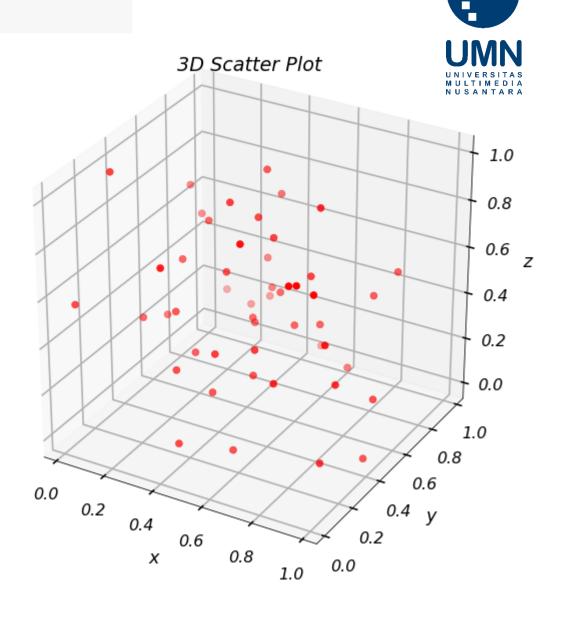
```
In [48]:
         In [49]:
         M fig = plt.figure(figsize = (8,8))
            ax = plt.axes(projection='3d')
            ax.grid()
            t = np.arange(0, 10*np.pi, np.pi/50)
            x = np.sin(t)
            y = np.cos(t)
            ax.plot3D(x, y, t)
            ax.set title('3D Parametric Plot')
            # Set axes Label
            ax.set_xlabel('x', labelpad=20)
            ax.set_ylabel('y', labelpad=20)
            ax.set zlabel('t', labelpad=20)
            plt.show()
```

 Try to rotate the above figure, and get a 3D view of the plot. You may notice that we also set the labelpad=20 to the 3-axis labels, which will make the label not overlap with the tick texts.



```
₩ # We can turn off the interactive plot using %matplotlib inline
In [50]:
             %matplotlib inline
In [51]:
          M \times = np.random.random(50)
             y = np.random.random(50)
             z = np.random.random(50)
             fig = plt.figure(figsize = (10,10))
             ax = plt.axes(projection='3d')
             ax.grid()
             ax.scatter(x, y, z, c = 'r', s = 50)
             ax.set title('3D Scatter Plot')
             # Set axes Label
             ax.set_xlabel('x', labelpad=20)
             ax.set ylabel('y', labelpad=20)
             ax.set zlabel('z', labelpad=20)
             plt.show()
```

 We could also plot 3D scatter plot using scatter function.



Surface Plotting



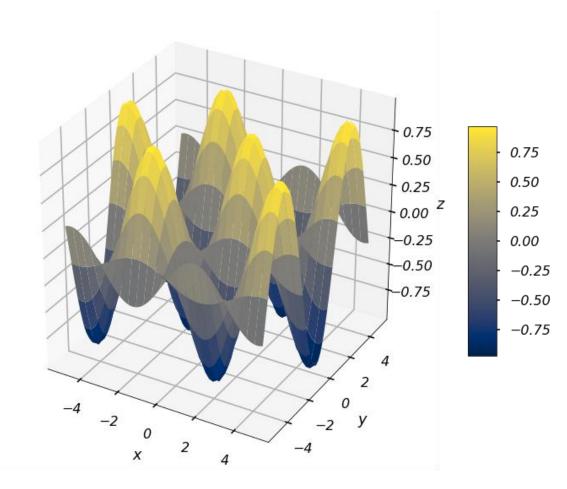
- In surface plotting, the first data structure you must create is called a mesh. Given lists/arrays of x and y values, a mesh is a listing of all the possible combinations of x and y.
- In Python, the mesh is given as two arrays X and Y, where X(i,j) and Y(i,j) define possible (x,y) pairs. A third array, Z, can then be created such that Z(i,j) = f(X(i,j),Y(i,j)).
- A mesh can be created using the np.meshgrid function in Python. The meshgrid function has the inputs x and y are lists containing the independent data set. The output variables X and Y are as described earlier.

• We could plot 3D surfaces in Python too, the function to plot the 3D surfaces is $plot_surface(X,Y,Z)$, where X and Y are the output arrays from meshgrid, and Z = f(X,Y) or Z(i,j) = f(X(i,j),Y(i,j)).

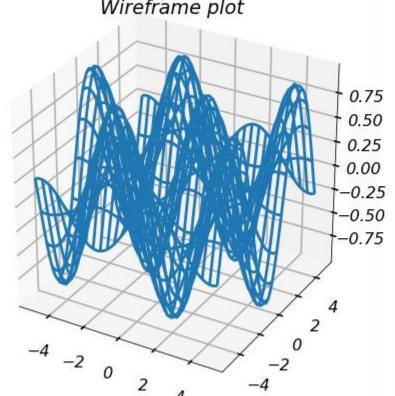


The most common surface plotting functions are surf and contour.

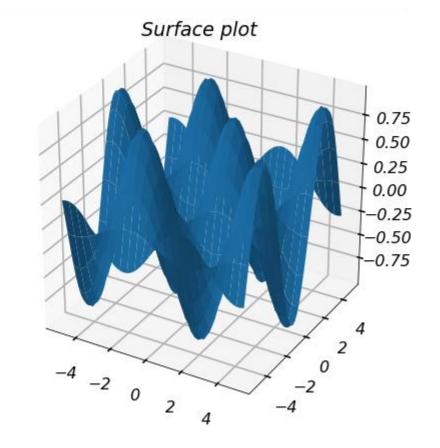
```
In [54]: M | fig = plt.figure(figsize = (12,10))
             ax = plt.axes(projection='3d')
             x = np.arange(-5, 5.1, 0.2)
             y = np.arange(-5, 5.1, 0.2)
             X, Y = np.meshgrid(x, y)
             Z = np.sin(X)*np.cos(Y)
             surf = ax.plot surface(X, Y, Z, cmap = plt.cm.cividis)
             # Set axes LabeL
             ax.set xlabel('x', labelpad=20)
             ax.set ylabel('y', labelpad=20)
             ax.set zlabel('z', labelpad=20)
             fig.colorbar(surf, shrink=0.5, aspect=8)
             plt.show()
```



 We could have subplots of different 3D plots as well. We could use the add_subplot function from the figure object we created to generate the subplots for 3D cases.







Parallel Your Python



- Now, we will introduce how to run your Python program in parallel so that we can reduce
 the time of completing the task.
- For smaller project, the benefit may not seem obvious. But for some complicated projects, the gain is significantly.
- For example, if I give you a homework that normally you need to wait for 1 hour until you can tell whether your code is correct or not. If you find something is wrong, and you change your code accordingly. Now it is time to wait another hour to find out whether your modification is proper. At the end, you made changes for 20 times, and spend more than a day to get it right. But if you can run your algorithms in a parallel way, the running time of your code may shrink to 5 min. 20 times modification only cost you less than 2 hours.
- Now you see why we bother to learn how to run our code faster using the parallelization.

Parallel Computing Basics



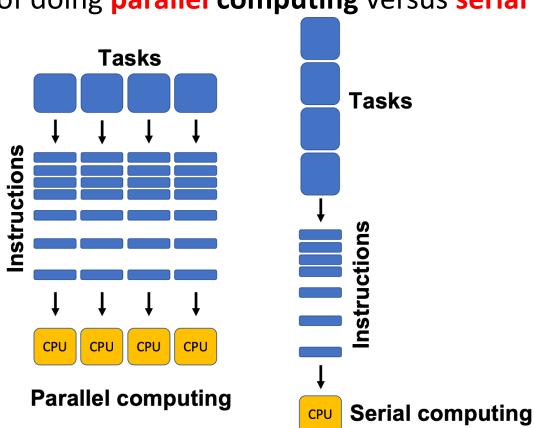
The fundamental idea of **parallel computing** is rooted in doing **multiple tasks** at the **same time** to reduce the running time of your program.

The following figure illustrates the simple idea of doing parallel computing versus serial

computing that we used so far.

 Most of the modern computers are using the multi-core design, which means on a single computing component, there are multiple independent processing units, the so called cores, that are available to do different tasks.

https://pythonnumericalmethods.berkeley.edu/no tebooks/chapter13.01-Parallel-Computing-Basics.html



Process and Thread



- In Python, there are two basic approaches to conduct parallel computing, that is using the multiprocessing or threading library.
- A process is an instance of a program (such as Python interpreter, Jupyter notebook etc.). A process is created by the operating system to run program, and each process has its own memory block. A thread is a sub-process that reside within the process. Each process can have multiple threads, that these threads will share the same memory block within the process.
- Therefore, for multiple threads in a process, due to the shared memory space, the
 variables or objects are all shared. If you change one variable in one thread, it will
 change for all the other threads. But things are different in different processes, change
 one variable in one process will not change the one in other processes.
- Process and thread both have advantages or disadvantages, and can be used in different tasks to maximize the benefits.

Python's GIL Problem



- Python has something called Global Interpreter Lock (GIL) which allow only one native
 thread to run at a time, it prevents multiple threads from running simultaneously.
- This is because Python was designed before the multi-core processor on the personal computers (this shows you how old the language is).
- Even though there are workarounds in Python to do multi-threading programming, we
 will only cover the multiprocessing library in the next section, which we will use most of
 the time for taking advantage of multi-core parallel computing.
- Of course, there are disadvantages of using parallel computing. Such as, more complicated code and overheads when spawn new processes.
- Thus, if your task is **small**, using parallel computing will **take longer** time, since it takes time for the system to **initialize new process** and **maintain** them.





- In Python, there are also other 3rd party packages that can make the parallel computing easier, especially for some daily tasks.
- joblib is one of them, it provides an easy simple way to do parallel computing (it has many other usages as well).
- First you need to install it by running: | | !pip install joblib
- The Parallel is a helper class that essentially provides a convenient interface for the multiprocessing module. The delayed is used to capture the arguments of the target function, in this case, the random_square.

```
from joblib import Parallel, delayed
import numpy as np

def random_square(seed):
    np.random.seed(seed)
    random_num = np.random.randint(0, 10)
    return random_num**2
```

```
In [2]: M import time

t0 = time.time()

results = Parallel(n_jobs=2)\
          (delayed(random_square)(i) for i in range(1000000))

t1 = time.time()
print(f'Execution time {t1 - t0} s')
```

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```

Execution time 49,224443435668945 s

```
    We can turn on the verbose
argument to output the status
messages.
```

 There are multiple backends in joblib, which means using different ways to do the parallel computing.

```
In [3]: M t0 = time.time()
            results = Parallel(n jobs=-1, verbose=1)\
                (delayed(random square)(i) for i in range(1000000))
            t1 = time.time()
            print(f'Execution time {t1 - t0} s')
            [Parallel(n jobs=-1)]: Using backend LokyBackend with 4 concurrent workers.
            [Parallel(n jobs=-1)]: Done 24640 tasks
                                                           elapsed:
                                                                       1.25
            [Parallel(n jobs=-1)]: Done 205240 tasks
                                                            elapsed:
                                                                      8.95
            [Parallel(n jobs=-1)]: Done 506240 tasks
                                                            elapsed:
                                                                      22.95
            [Parallel(n jobs=-1)]: Done 927640 tasks
                                                          elapsed:
                                                                      42.85
            Execution time 46.2181658744812 s
```

[Parallel(n jobs=-1)]: Done 1000000 out of 1000000 | elapsed: 46.1s finis





- Basics of Linear Algebra
- Linear Transformations
- Systems of Linear Equations
- Solutions to Systems of Linear Equations
- Solve Systems of Linear Equations in Python
- Matrix Inversion





- Kong, Qingkai; Siauw, Timmy, and Bayen, Alexandre. 2020. Python Programming and Numerical Methods: A Guide for Engineers and Scientists. Academic Press.
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- Other online and offline references



Menjadi Program Studi Strata Satu Informatika **unggulan** yang menghasilkan lulusan **berwawasan internasional** yang **kompeten** di bidang Ilmu Komputer (*Computer Science*), **berjiwa wirausaha** dan **berbudi pekerti luhur**.





- I. Menyelenggarakan pembelajaran dengan teknologi dan kurikulum terbaik serta didukung tenaga pengajar profesional.
- 2. Melaksanakan kegiatan penelitian di bidang Informatika untuk memajukan ilmu dan teknologi Informatika.
- 3. Melaksanakan kegiatan pengabdian kepada masyarakat berbasis ilmu dan teknologi Informatika dalam rangka mengamalkan ilmu dan teknologi Informatika.