Student name: Roja Kamble, 11454258

Assignment: 4

Part 1: Data Wrangling

You have to write code to answer the questions below

- Import pandas library
- Read the data stored in your local machine https://www.kaggle.com/datasets/yasserh/student-marks-dataset (https://www.kaggle.com/datasets/yasserh/student-marks-dataset)
- Name the variable df

Show information about the dataset

```
In [6]: import pandas as pd
import numpy as np

df = pd.read_csv('/Users/roja/Downloads/Student_Marks.csv')
    df
```

Out[6]:

	number_courses	time_study	Marks
0	3	4.508	19.202
1	4	0.096	7.734
2	4	3.133	13.811
3	6	7.909	53.018
4	8	7.811	55.299
95	6	3.561	19.128
96	3	0.301	5.609
97	4	7.163	41.444
98	7	0.309	12.027
99	3	6.335	32.357

100 rows × 3 columns

Show the last 10 rows

Out[7]:

	number_courses	time_study	Marks
90	7	4.182	24.394
91	8	2.730	19.564
92	4	5.027	23.916
93	8	6.471	42.426
94	8	3.919	24.451
95	6	3.561	19.128
96	3	0.301	5.609
97	4	7.163	41.444
98	7	0.309	12.027
99	3	6.335	32.357

Reset the index such that it starts from 1 (instead of 0)

Out[8]:

	number_courses	time_study	Marks
1	3	4.508	19.202
2	4	0.096	7.734
3	4	3.133	13.811
4	6	7.909	53.018
5	8	7.811	55.299
96	6	3.561	19.128
97	3	0.301	5.609
98	4	7.163	41.444
99	7	0.309	12.027
100	3	6.335	32.357

100 rows × 3 columns

Lowercase all columns

```
In [9]: # Basically column heads are already lowercase except 'Marks'column

t1 = pd.read_csv('/Users/roja/Downloads/Student_Marks.csv')
t1.head()
t1.columns = t1.columns.str.lower()
t1
```

Out[9]:

	number_courses	time_study	marks
0	3	4.508	19.202
1	4	0.096	7.734
2	4	3.133	13.811
3	6	7.909	53.018
4	8	7.811	55.299
95	6	3.561	19.128
96	3	0.301	5.609
97	4	7.163	41.444
98	7	0.309	12.027
99	3	6.335	32.357

100 rows × 3 columns

```
In [10]: # Uppercase all columns

t2 = pd.read_csv('/Users/roja/Downloads/Student_Marks.csv')
t2.head()
t2.columns = t2.columns.str.upper()
t2
```

Out[10]:

	NUMBER_COURSES	TIME_STUDY	MARKS
0	3	4.508	19.202
1	4	0.096	7.734
2	4	3.133	13.811
3	6	7.909	53.018
4	8	7.811	55.299
95	6	3.561	19.128
96	3	0.301	5.609
97	4	7.163	41.444
98	7	0.309	12.027
99	3	6.335	32.357

100 rows × 3 columns

Return a boolean value indicating whether the dataset has missing values

```
In [11]: df.isnull()
```

Out[11]:

	number_courses	time_study	Marks
1	False	False	False
2	False	False	False
3	False	False	False
4	False	False	False
5	False	False	False
96	False	False	False
97	False	False	False
98	False	False	False
99	False	False	False
100	False	False	False

100 rows × 3 columns

Show average of all columns grouped by number_course in a same DataFrame table

```
In [12]: avg = df.groupby(['number_courses']).mean()
avg
```

Out[12]:

	time_study	Marks
number_courses		
3	3.685773	18.433318
4	3.598524	19.029952
5	3.070600	17.641000
6	4.838750	29.863062
7	4.242667	29.401467
8	4.955750	33.835375

Show the sum of time_study and mean of marks grouped by each number_course in a same DataFrame table

```
In [14]: | sum = df.groupby(['number_courses'])['time_study'].sum()
         sum
Out[14]: number courses
         3
              81.087
              75.569
              30.706
              77.420
              63.640
              79.292
         Name: time study, dtype: float64
In [15]: mean = df.groupby(['number_courses'])['Marks'].mean()
         mean
Out[15]: number_courses
              18.433318
              19.029952
              17.641000
              29.863062
              29.401467
              33.835375
         Name: Marks, dtype: float64
In [16]: pip install -U prettytable
```

Requirement already satisfied: prettytable in /Users/roja/opt/anaconda3/lib/python3.9/site-packages (3.5.0)

Requirement already satisfied: wcwidth in /Users/roja/opt/anaconda3/lib/python3.9/site-packages (from prettytable) (0.2.5)

Note: you may need to restart the kernel to use updated packages.

8 79.292 8 33.835375
Name: time_study, dtype: float64 Name: Marks, dtype: float64

6 77.420

7 63.640

Write a class My_KMeans that implements k-means clustering algorithm. You are required to have the following attributes

- · Parameters:
 - Default clusters
 - max_iter
- Method:
 - fit
 - predict

Reference: https://scikit-learn.org/stable/modules/generated/sklearn.cluster.KMeans.html (https://scikit-learn.org/stable/modules/generated/sklearn.cluster.KMeans.html)

6 29.863062

7 29.401467

Using a pre-built library yields no credit. You have to write everything from scratch

```
In [18]: import math
         import numpy as np
         def euclidean(point, data):
             return np.sqrt(np.sum((point - data)**2, axis=1))
         class My KMeans:
             1= []
             def init (self, n clusters = 3, max iter = 100):
                 self.n clusters = n clusters
                 self.max iter = max iter
                 self.centroids =0
             def fit(self, X train):
                 min , max = np.min(X train, axis=0), np.max(X train, axis=0)
                 self.centroids = [np.random.uniform(min , max ) for in range(self.n clusters)]
                 iteration = 0
                 prev centroids = None
                 while np.not equal(self.centroids, prev centroids).any() and iteration < self.max iter:</pre>
                     sorted points = [[] for in range(self.n clusters)]
                     for x in X train:
                         dists = euclidean(x, self.centroids)
                         centroid idx = np.argmin(dists)
                         sorted points[centroid idx].append(x)
                     prev centroids = self.centroids
                     self.centroids = [np.mean(cluster, axis=0) for cluster in sorted points]
                     for i, centroid in enumerate(self.centroids):
                         if np.isnan(centroid).any():
                             self.centroids[i] = prev centroids[i]
                     iteration += 1
             def predict(self, X):
                 centroids = []
                 centroid idxs = []
                 for x in X:
```

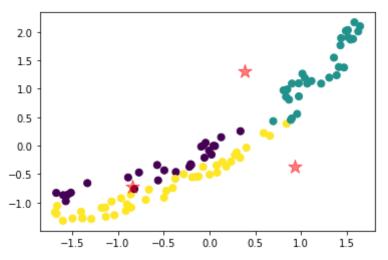
```
dists = euclidean(x, self.centroids)
  centroid_idx = np.argmin(dists)
  t = self.centroids[centroid_idx]
  centroids.append(t)
  centroid_idxs.append(centroid_idx)

cluster_centers_ = centroid_idxs
  self.l = centroid_idxs
  return centroid_idxs
```

Run the code

```
In [19]: learn.preprocessing import StandardScaler
        matplotlib.pyplot as plt
         seaborn as sns
         ardize the data
         StandardScaler().fit_transform(df)
        ocal implementation of kmeans
         = My KMeans(3,100)
        fit(X std)
        s = kmeans.predict(X std)
         , 1]: X_std dataset with all rows and the second column (time_study in this case)
         , 2]: X_std dataset with all rows and the third column (marks in this case)
         cluster elements
        tter(x=X_std[:, 1], y=X_std[:, 2], c=y_kmeans, s=50, cmap='viridis')
         cluster centers
        tter([x for x, _, _ in kmeans.centroids], [y for _, _, y in kmeans.centroids], marker='*', color='r', s=2
```

Out[19]: <matplotlib.collections.PathCollection at 0x7f792a020490>



Your analysis about the plot

According to my inspection of the previous plot, both the points and clusters, as well as the centroid that was created, are not precise. Even if the model appears to be functioning OK, a few corrections should still need to be made. However, the model that is now being created may be applied since the outcomes are adequate.

Type *Markdown* and LaTeX: α^2

(Bonus 20pts) Modify your algorithm to k-means++ with a smart initialization, and repeat the task above

```
In [20]: def get kmeans pp centroids(X1, k = 5):
             centroids = []
             for i in range(k):
                 centroids.append([X1[10,0],X1[10,1],X1[10,2]])
             i = 1
             d = []
             while i <= k:
                 r = centroids[i-2 if i-2 >= 0 else 0]
                 maxd = 0
                 for j in X1:
                     d = np.sqrt((j[1] - r[1])**2 + (j[2] - r[2])**2)
                      if d > maxd:
                          centroids[i-1] = [j[0], j[1], j[2]]
                         maxd = d
                 i+=1
             min , max = np.min(X1, axis=0), np.max(X1, axis=0)
             iteration = 0
             prev centroids = None
             while np.not equal(centroids, prev centroids).any() and iteration < k:</pre>
                 sorted points = [[] for in range(k)]
                 for x in X1:
                      dists = euclidean(x, centroids)
                      centroid idx = np.argmin(dists)
                     sorted points[centroid idx].append(x)
                 prev centroids = centroids
                 centroids = [np.mean(cluster, axis=0) for cluster in sorted points]
                 for i, centroid in enumerate(centroids):
                     if np.isnan(centroid).any():
                         centroids[i] = prev centroids[i]
                  iteration += 1
             return centroids
         def predict(X,c):
             centroids = []
             centroid idxs = []
             for x in X:
                 dists = euclidean(x,c)
                 centroid idx = np.argmin(dists)
```

```
t = c[centroid_idx]
  centroids.append(t)
  centroid_idxs.append(centroid_idx)
  return centroid_idxs
```

```
In [21]: centroids = get_kmeans_pp_centroids(X_std, k = 3)
    y_kmeans = predict(X_std,centroids)

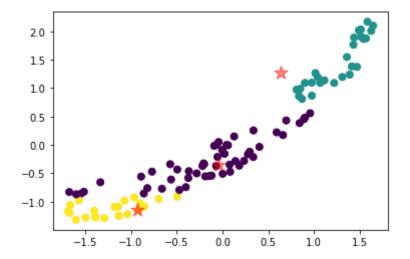
# Plot cluster elements
    plt.scatter(x=X_std[:, 1], y=X_std[:, 2], c=y_kmeans, s=50, cmap='viridis')

# Plot cluster centers
    plt.scatter([x for x, _, _ in centroids], [y for _, _, y in centroids], marker='*', color='r', s=200, alputation of the color of the
```

/Users/roja/opt/anaconda3/lib/python3.9/site-packages/numpy/core/fromnumeric.py:3419: RuntimeWarning: Mean of empty slice.

return _methods._mean(a, axis=axis, dtype=dtype,
/Users/roja/opt/anaconda3/lib/python3.9/site-packages/numpy/core/_methods.py:188: RuntimeWarning: inva
lid value encountered in double_scalars
ret = ret.dtype.type(ret / rcount)

Out[21]: <matplotlib.collections.PathCollection at 0x7f792a1ea280>



```
In [ ]:
```