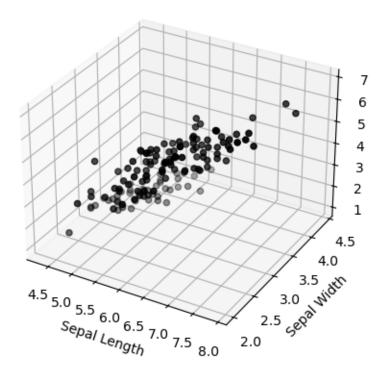
## USL KNN NB

## March 2, 2025

```
[1]: import numpy as np
     import pandas as pd
     import matplotlib.pyplot as plt
     import seaborn as sns
     from sklearn.datasets import load_iris
     from sklearn.model selection import train test split
     from sklearn.cluster import KMeans
     from sklearn.metrics import accuracy score
     from sklearn.preprocessing import StandardScaler
[2]: url = r"D:\Unsupervised Machine Learning Lab (USLL)\.Lab\Iris.csv"
     df = pd.read_csv(url)
     df.head()
[2]:
        Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm
                                                                           Species
                      5.1
                                                   1.4
                                                                 0.2 Iris-setosa
                                    3.5
                      4.9
                                    3.0
                                                   1.4
                                                                 0.2 Iris-setosa
     1
         2
     2
         3
                      4.7
                                    3.2
                                                   1.3
                                                                 0.2 Iris-setosa
     3
         4
                      4.6
                                    3.1
                                                   1.5
                                                                 0.2 Iris-setosa
                      5.0
                                                   1.4
                                                                 0.2 Iris-setosa
     4
         5
                                    3.6
[3]: df.drop('Id', axis=1, inplace=True)
     df.head()
[3]:
       SepalLengthCm SepalWidthCm PetalLengthCm
                                                    PetalWidthCm
                                                                       Species
     0
                  5.1
                                3.5
                                               1.4
                                                              0.2 Iris-setosa
                  4.9
     1
                                3.0
                                               1.4
                                                             0.2 Iris-setosa
     2
                  4.7
                                3.2
                                               1.3
                                                             0.2 Iris-setosa
                                3.1
     3
                  4.6
                                               1.5
                                                             0.2 Iris-setosa
     4
                  5.0
                                3.6
                                               1.4
                                                             0.2 Iris-setosa
[4]: X = df.iloc[:, 0]
     y = df.iloc[:, 1]
     z = df.iloc[:, 2]
[5]: df1 = pd.DataFrame(df.iloc[:, [0, 1, 2]].values)
     df1.head()
```

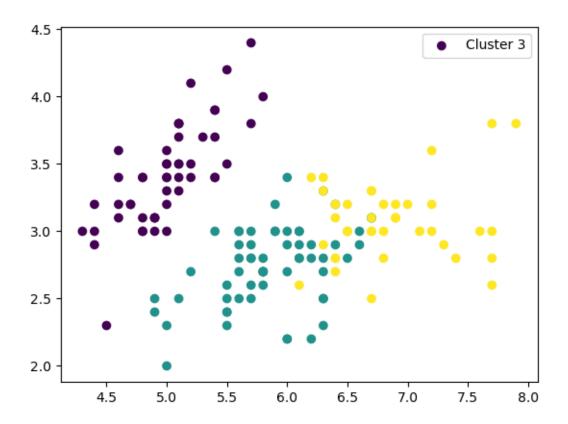
```
[5]: 0 1 2
0 5.1 3.5 1.4
1 4.9 3.0 1.4
2 4.7 3.2 1.3
3 4.6 3.1 1.5
4 5.0 3.6 1.4
```

```
fig = plt.figure()
ax = fig.add_subplot(111, projection='3d')
ax.scatter(X, y, z, c='black')
ax.set_xlabel('Sepal Length')
ax.set_ylabel('Sepal Width')
ax.set_zlabel('Petal Length')
plt.show()
```



```
[ ]: kmeans = KMeans(n_clusters=3)
[8]: # Fit the KMeans model
kmeans.fit(df1)
# Predict the clusters
```

```
predictions = kmeans.predict(df1)
     # Display the predictions
     print(predictions)
    2 1]
    c:\Users\Neil\anaconda3\Lib\site-
    packages\joblib\externals\loky\backend\context.py:136: UserWarning: Could not
    find the number of physical cores for the following reason:
    [WinError 2] The system cannot find the file specified
    Returning the number of logical cores instead. You can silence this warning by
    setting LOKY_MAX_CPU_COUNT to the number of cores you want to use.
      warnings.warn(
      File "c:\Users\Neil\anaconda3\Lib\site-
    packages\joblib\externals\loky\backend\context.py", line 257, in
    count physical cores
       cpu info = subprocess.run(
                ____
      File "c:\Users\Neil\anaconda3\Lib\subprocess.py", line 548, in run
       with Popen(*popenargs, **kwargs) as process:
      File "c:\Users\Neil\anaconda3\Lib\subprocess.py", line 1026, in __init__
       self._execute_child(args, executable, preexec_fn, close_fds,
      File "c:\Users\Neil\anaconda3\Lib\subprocess.py", line 1538, in _execute_child
       hp, ht, pid, tid = _winapi.CreateProcess(executable, args,
    c:\Users\Neil\anaconda3\Lib\site-packages\sklearn\cluster\ kmeans.py:1429:
    UserWarning: KMeans is known to have a memory leak on Windows with MKL, when
    there are less chunks than available threads. You can avoid it by setting the
    environment variable OMP_NUM_THREADS=1.
      warnings.warn(
[23]: plt.scatter(df1[0], df1[1], c=predictions, cmap='viridis')
     plt.legend({'Cluster 1'})
     plt.legend({'Cluster 2'})
     plt.legend({'Cluster 3'})
     plt.show()
```



```
[]: import numpy as np
     import pandas as pd
     import matplotlib.pyplot as plt
     from sklearn.cluster import KMeans
     from sklearn.datasets import load_iris
     from mpl_toolkits.mplot3d import Axes3D
     # Load the iris dataset
     iris = load_iris()
     df = pd.DataFrame(iris.data, columns=iris.feature_names)
     # Select three features for 3D visualization
     features = ['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)']
     data = df[features]
     fig = plt.figure(figsize=(10, 7))
     ax = fig.add_subplot(111, projection='3d')
     ax.scatter(data.iloc[:, 0], data.iloc[:, 1], data.iloc[:, 2], cmap='viridis',__
      →marker='o', s=50)
     ax.set xlabel(features[0])
     ax.set_ylabel(features[1])
```

```
ax.set_zlabel(features[2])
ax.set_title('K-Means Clustering on Iris Dataset (3D)')
ax.legend()
# Perform K-Means clustering
kmeans = KMeans(n_clusters=3, random_state=42)
kmeans.fit(data)
labels = kmeans.labels
centroids = kmeans.cluster_centers_
# Plot the clusters in 3D
fig = plt.figure(figsize=(10, 7))
ax = fig.add_subplot(111, projection='3d')
ax.scatter(data.iloc[:, 0], data.iloc[:, 1], data.iloc[:, 2], c=labels,_u
 ⇔cmap='viridis', marker='o', s=50)
# Plot centroids
ax.scatter(centroids[:, 0], centroids[:, 1], centroids[:, 2], c='red', __
  ⇔marker='X', s=200, label='Centroids')
# Labels and title
ax.set_xlabel(features[0])
ax.set_ylabel(features[1])
ax.set_zlabel(features[2])
ax.set_title('K-Means Clustering on Iris Dataset (3D)')
ax.legend()
plt.show()
C:\Users\Neil\AppData\Local\Temp\ipykernel_42600\2503888613.py:18: UserWarning:
No data for colormapping provided via 'c'. Parameters 'cmap' will be ignored
  ax.scatter(data.iloc[:, 0], data.iloc[:, 1], data.iloc[:, 2], cmap='viridis',
marker='o', s=50)
C:\Users\Neil\AppData\Local\Temp\ipykernel_42600\2503888613.py:23: UserWarning:
No artists with labels found to put in legend. Note that artists whose label
```

c:\Users\Neil\anaconda3\Lib\site-packages\sklearn\cluster\\_kmeans.py:1429:

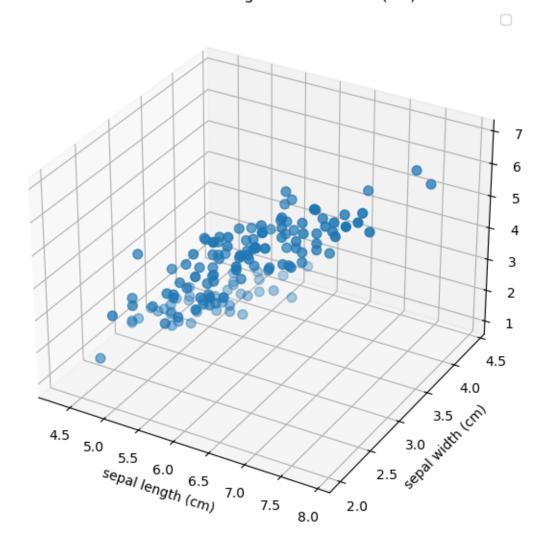
start with an underscore are ignored when legend() is called with no argument.

environment variable OMP\_NUM\_THREADS=1.

warnings.warn(

ax.legend()

## K-Means Clustering on Iris Dataset (3D)



## K-Means Clustering on Iris Dataset (3D)

