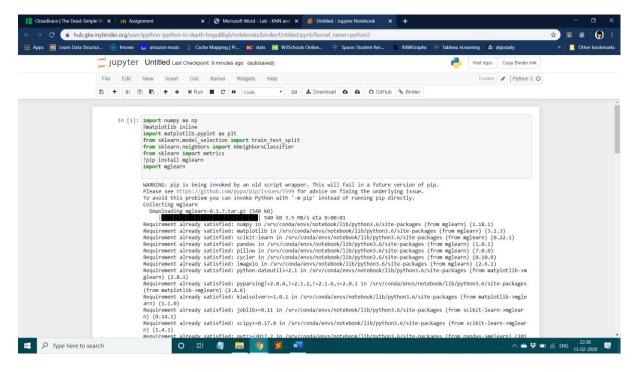
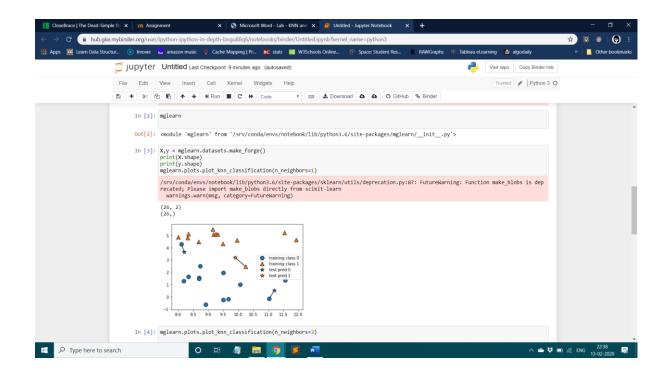
Lab 6

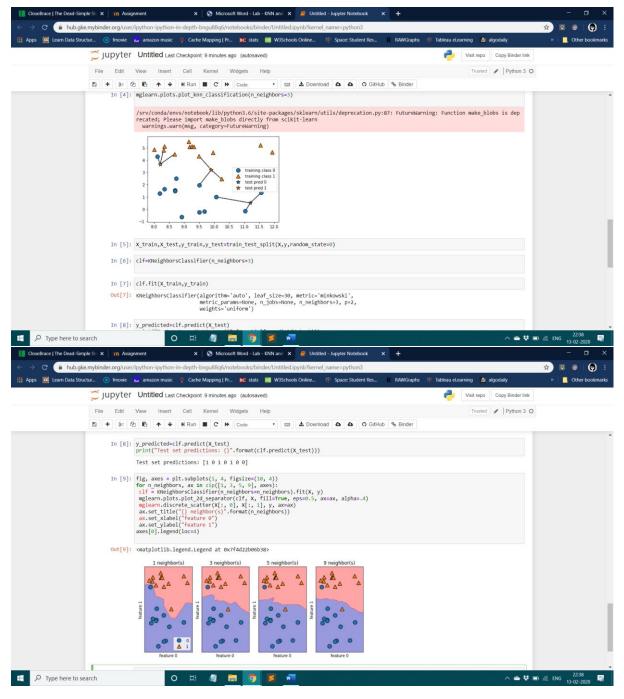
R Harini

18BCE1010

KNN Classifier







df=pd.read_csv('cancer.csv')

df

df= df.drop(columns=['id','Unnamed: 32'])

df.head()

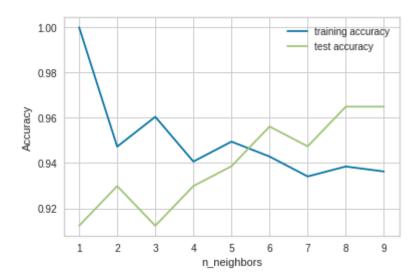
from sklearn.model_selection import train_test_split

array=df.values

df.head()

X=array[:,1:33]

```
y=array[:,[0]]
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)
X_train.shape,X_test.shape
clf=KNeighborsClassifier(n_neighbors=3)
clf.fit(X_train,y_train)
acc_train = clf.score(X_train, y_train)
print('Train set accuracy: ', acc_train)
acc_train = clf.score(X_test, y_test)
print('Train set accuracy: ', acc_train)
training_accuracy =[]
testing_accuracy = []
neighbors_settings = range(1, 10)
for n_neighbors in neighbors_settings:
  clf = KNeighborsClassifier(n_neighbors=n_neighbors)
  clf.fit(X_train, y_train)
  training_accuracy.append(clf.score(X_train, y_train))
  testing_accuracy.append(clf.score(X_test, y_test))
plt.plot(neighbors_settings, training_accuracy, label="training accuracy")
plt.plot(neighbors_settings, testing_accuracy, label="test accuracy")
plt.ylabel("Accuracy")
plt.xlabel("n_neighbors")
plt.legend()
plt.show()
```



KMeans Clustering

```
from sklearn.cluster import KMeans
import numpy as np

df=pd.read_csv('protein.csv')

df

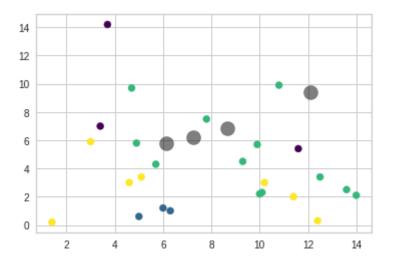
X = df.iloc[:,1:].values
kmeans = KMeans(n_clusters=4)

y_kmeans = kmeans.fit_predict(X)

print(y_kmeans5)
kmeans.cluster_centers_
plt.scatter(X[:, 1], X[:,4], c=y_kmeans, s=50, cmap='viridis')

centers = kmeans.cluster_centers_
```

plt.scatter(centers[:, 0], centers[:, 1], c='black', s=200, alpha=0.5);



from sklearn.metrics import pairwise_distances_argmin

centers = new_centers

return centers, labels

```
Error =[]
for i in range(1, 11):
    kmeans = KMeans(n_clusters = i).fit(X)
    kmeans.fit(X)
    Error.append(kmeans.inertia_)
import matplotlib.pyplot as plt
plt.plot(range(1, 11), Error)
plt.title('Elbow method')
plt.xlabel('No of clusters')
plt.ylabel('Error')
plt.show()
```

