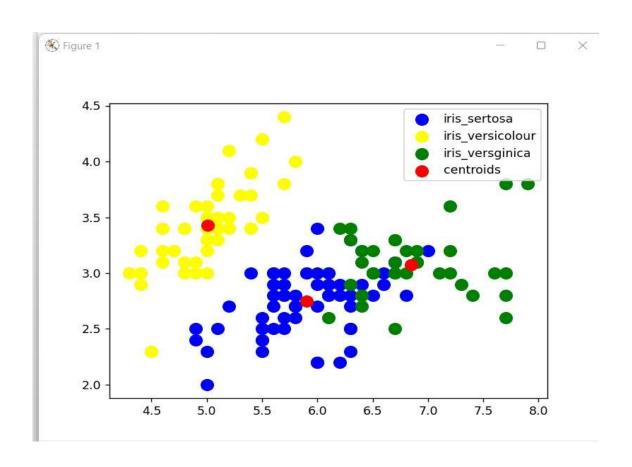
# Assignment No. 1

# Code:import numpy as np import matplotlib.pyplot as plt import pandas as pd from sklearn.cluster import KMeans X = [[0.1,0.6],[0.15,0.71],[0.08,0.9],[0.16,0.85],[0.2,0.3],[0.25,0.5],[0.24,0.1],[0.3,0.2]]centroids=np.array([[0.1,0.6],[0.25,0.5]]) kmeans=KMeans(n\_clusters=2,init=centroids) kmeans.fit(X) print("Labels after trainig:",kmeans.labels\_) #Q1 print("P6 belongs to ",kmeans.labels\_[5],"cluster") #Q2 print("Population around centroid 2(P6) is ",np.count\_nonzero(kmeans.labels\_==1)) #Q3 print("New values of centroids are:",kmeans.cluster\_centers\_) from sklearn import datasets iris=datasets.load\_iris()

iris\_x=iris.data

```
wcss=list()
for i in range(1,11):
  kmeans=KMeans(n_clusters=i,init='k-means++')
  kmeans.fit(iris_x)
  wcss.append(kmeans.inertia_)
plt.plot(range(1,11),wcss)
plt.xlabel('No. of clusters')
plt.ylabel('WCSS')
plt.show()
kmeans=KMeans(n clusters=3,init='k-means++')
kmeans.fit(iris_x)
y_means=kmeans.predict(iris_x)
plt.scatter(iris_x[y_means==0,0],iris_x[y_means==0,1],c='blue',s=100,label='iris_sertosa')
plt.scatter(iris_x[y_means==1,0],iris_x[y_means==1,1],c='yellow',s=100,label='iris_versicolo
ur')
plt.scatter(iris_x[y_means==2,0],iris_x[y_means==2,1],c='green',s=100,label='iris_versginica'
plt.scatter(kmeans.cluster_centers_[:,0],kmeans.cluster_centers_[:,1],c='red',s=100,label='c
entroids')
plt.legend()
plt.show()
```

## Output:-



# Assignment No. 3

```
Code:-
from sklearn.linear_model import LinearRegression
import matplotlib.pyplot as plt
import pandas as pd
from sklearn.datasets import load_diabetes
from sklearn.linear_model import LinearRegression, Ridge
from sklearn.model_selection import cross_val_score
diabetes = load_diabetes()
Ir = LinearRegression(normalize = True)
lr_scores = cross_val_score(lr, diabetes.data, diabetes.target,cv=10)
print("Scores obtained by Linear Regression: ",Ir scores)
print("\nMean: ",lr_scores.mean())
Output:-
Scores obtained by Linear Regression: [0.55614411 0.23056092 0.35357777 0.62190498
0.26587602 0.61819338
0.41815916 0.43515232 0.43436983 0.68568514]
Mean: 0.4619623619583371
## Ridge
rg = Ridge(0.005, normalize=True)
rg_scores = cross_val_score(rg, diabetes.data, diabetes.target, cv=10)
```

print("Scores obtained by Ridge Regression: ",rg\_scores)

```
print("\nMean: ",rg_scores.mean())
Output:-
Scores obtained by Ridge Regression: [0.55014745 0.24000807 0.36373382 0.61657429
0.2695777 0.62172635
0.42346862 0.42813504 0.43117332 0.68303604]
Mean: 0.462758069707298
## RidgeCV for set of alpha values, alpha_ to extract alpha value
from sklearn.linear_model import RidgeCV
rg = RidgeCV(alphas=(1.0, 0.1, 0.01, 0.005, 0.0025, 0.001, 0.00025),normalize=True)
rg.fit(diabetes.data, diabetes.target)
print(rg.alpha )
rg_cv_scores = cross_val_score(rg, diabetes.data, diabetes.target, cv=10)
print("Scores obtained by RidgeCV Regression: ",rg_cv_scores)
print("\nMean: ",rg_scores.mean())
Output:-
Scores obtained by RidgeCV Regression: [0.52803256 0.23657595 0.3565488 0.607513
0.2695777 0.62047382
0.42149214 0.43894932 0.43138195 0.6642474 ]
Mean: 0.462758069707298
## Lasso, LassoCV
from sklearn.linear model import Lasso, LassoCV
Is = Lasso(alpha=0.005, normalize=True)
ls_scores = cross_val_score(ls, diabetes.data, diabetes.target, cv=10)
```

print("Scores obtained by Lasso Regression: ",ls\_scores)

print("\nMean: ",ls\_scores.mean())

#### Output:-

Scores obtained by Lasso Regression: [0.55078146 0.23993097 0.36453647 0.61454396 0.26894036 0.62269952

0.42465645 0.42726522 0.43075777 0.68414508]

Mean: 0.46282572553668555

from sklearn.linear\_model import LassoCV

ls\_cv = LassoCV(alphas=(1.0,0.1,0.01,0.005,0.0025,0.001,0.00025),normalize=True)

ls\_cv.fit(diabetes.data, diabetes.target)

ls\_cv.alpha\_

ls\_cv\_scores = cross\_val\_score(ls\_cv, diabetes.data, diabetes.target,cv=10)

print("Scores obtained by LassoCV Regression: ",ls\_cv\_scores)

print("\nMean: ",ls\_cv\_scores.mean())

Output:-

Scores obtained by LassoCV Regression: [0.51796189 0.23824666 0.35415718 0.59752149 0.27503201 0.62269952

0.41851368 0.42047653 0.42562829 0.68231416]

Mean: 0.45525514142746404

## ElasticNet, ElasticNetCV
from sklearn.linear\_model import ElasticNet, ElasticNetCV
en = ElasticNet(alpha=0.001,l1\_ratio=0.8, normalize=True)
en\_scores = cross\_val\_score(en, diabetes.data, diabetes.target, cv=10)
print("Scores obtained by ElasticNet Regression: ",en\_scores)
print("\nMean: ",en\_scores.mean())

```
Output:-
```

Scores obtained by ElasticNet Regression: [0.53103739 0.24682675 0.38160097 0.60832995 0.2830996 0.62083992

0.43113636 0.43484866 0.43055758 0.6676087 ]

Mean: 0.46358858847836454

```
encv = ElasticNetCV(alphas=(0.1, 0.01,0.005, 0.0025, 0.001),l1_ratio=(0.1,0.25,0.5,0.75,
0.8), normalize=True)
encv.fit(diabetes.data, diabetes.target)
print(encv.alpha_)
print(encv.l1_ratio_)
encv_scores = cross_val_score(encv, diabetes.data, diabetes.target,cv=10)
print("Scores obtained by ElasticNetCV Regression: ",encv_scores)
print("\nMean: ",encv_scores.mean())

Output:-
```

Scores obtained by ElasticNetCV Regression: [0.52796681 0.24682675 0.38160097 0.60743123 0.28526958 0.61931497

0.43113636 0.43907824 0.43137097 0.66429212]

Mean: 0.46342880159848276

## Comparative analysis

#Linear Regression Score

print(Ir\_scores.mean())

#Ridge Score

print(rg\_scores.mean())

#Lasso Score

print(Is\_scores.mean())

#ElasticNet Score

print(en\_scores.mean())

Output:- 0.4619623619583371

#### 0.462758069707298

## 0.46282572553668555

## 0.46358858847836454

import matplotlib.pyplot as plt

import numpy as np

objects = ('Linear','Ridge','Lasso','ElasticNet')

y\_pos = np.arange(len(objects))

**p** =

[0.4619623619583371, 0.4627580697072979, 0.4628257255366856, 0.46358858847836454]

plt.bar(y\_pos, p, align='center',alpha=0.5, color=['hotpink', 'yellow', 'lime','cyan'])

plt.xticks(y\_pos, objects)

plt.ylabel('Accuracy')

plt.title('Linear Models')

ElasticNetCV(alphas=(0.1, 0.01, 0.005, 0.0025, 0.001),

l1\_ratio=(0.1, 0.25, 0.5, 0.75, 0.8), normalize=True)

plt.show()

## Output:-

