MACHINE LEARNING PROGREMS OUTPUT VERIFIED

PROGREAM: 01

from sklearn.datasets import make_blobs

from matplotlib import pyplot as plt

from matplotlib import style

from sklearn.datasets import make_blobs

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from matplotlib import style

style.use("fivethirtyeight")

X, y = make_blobs(n_samples = 100, centers = 3,

cluster_std = 1, n_features = 2)

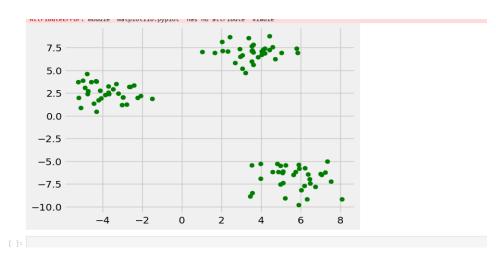
plt.scatter(X[:, 0], X[:, 1], s = 40, color = 'g')

plt.xlabel("X")

plt.ylabel("Y")

plt.show()

plt.clf()



```
from IPython.display import HTML
import pandas as pd
import numpy as np
import base64
def create_download_link(df, title = "Download CSV file", filename = "data.csv"):
  csv = df.to_csv()
  b64 = base64.b64encode(csv.encode())
  payload = b64.decode()
  html = '<a download="{filename}" href="data:text/csv;base64,{payload}" target="_blank">{title}</a>'
  html = html.format(payload=payload,title=title,filename=filename)
  return HTML(html)
df = pd.DataFrame(np.random.randn(50, 4), columns=list('ABCD'))
create_download_link(df)
     [6]: from IPython.display import HTML
         import pandas as pd
         import numpy as np
         import base64
    [14]: def create_download_link(df,title = "Download CSV file", filename = "data.csv"):
            csv = df.to_csv()
            b64 = base64.b64encode(csv.encode())
            payload = b64.decode()
            html = '<a download="{filename}" href="data:text/csv;base64,{payload}" target="_blank">{title}</a>'
            html = html.format(payload=payload,title=title,filename=filename)
            return HTML(html)
         df = pd.DataFrame(np.random.randn(50, 4), columns=list('ABCD'))
         create_download_link(df)
    [14]: Download CSV file
```

```
from pandas import read_csv

from numpy import set_printoptions

from sklearn.feature_selection import SelectKBest

from sklearn.feature_selection import chi2

path = r'D:\squad\book.csv'

names = ['preg', 'plas', 'hello']

dataframe = read_csv(path, names=names)

array = dataframe.values

print(dataframe)
```

	preg	plas	pres	skin	test	mass	pedi	age	class	
NaN	preg	plas	pres	skin	test	mass	pedi	age	class	
0.0	variance	skewness	Curtosis	Entropy	class	NaN	NaN	NaN	NaN	
1.0	3.6216	8.661	-2.8073	-0.4469	0	NaN	NaN	NaN	NaN	
2.0	4.5459	8.1674	-2.4586	-1.462	0	NaN	NaN	NaN	NaN	
3.0	3.866	-2.6383	1.9242	0.10645	0	NaN	NaN	NaN	NaN	
4.0	3.4566	9.5228	-4.0112	-3.594	0	NaN	NaN	NaN	NaN	

```
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy score
from sklearn.metrics import classification report
from sklearn.metrics import roc_auc_score
from sklearn.metrics import log loss
X \text{ actual} = [1, 1, 0, 1, 0, 0, 1, 0, 0, 0]
Y predic = [1, 0, 1, 1, 1, 0, 1, 1, 0, 0]
results = confusion matrix(X actual, Y predic)
print ('Confusion Matrix :')
print(results)
print ('Accuracy Score is',accuracy_score(X_actual, Y_predic))
print ('Classification Report : ')
print (classification report(X actual, Y predic))
print('AUC-ROC:',roc auc score(X actual, Y predic))
print('LOGLOSS Value is',log loss(X actual, Y predic))
```

```
Confusion Matrix :
[[3 3]]
[1 3]]
Accuracy Score is 0.6
Classification Report :
            precision recall f1-score support
                0.75 0.50
                                   0.60
         1
                0.50 0.75
                                   0.60
                                               4
   accuracy
                                   0.60
                                              10
macro avg 0.62 0.62
weighted avg 0.65 0.60
                                  0.60
                                              10
                                   0.60
AUC-ROC: 0.625
LOGLOSS Value is 14.41746135564686
```

```
from sklearn.datasets import load_iris

iris = load_iris()

X = iris.data

y=iris.target

from sklearn.model_selection import train_test_split

X_train,X_test,y_train, y_test=train_test_split(X,y,test_size=0.4,random_state=1)

from sklearn.naive_bayes import GaussianNB

gnb=GaussianNB()

gnb.fit(X_train, y_train)

y_pred=gnb.predict(X_test)

from sklearn import metrics

print("GaussianNaiveBayes model accuracy(in %):", metrics.accuracy_score(y_test,y_pred)*100)

GaussianNaiveBayes model accuracy(in %): 95.0
```

PROGREAM: 06

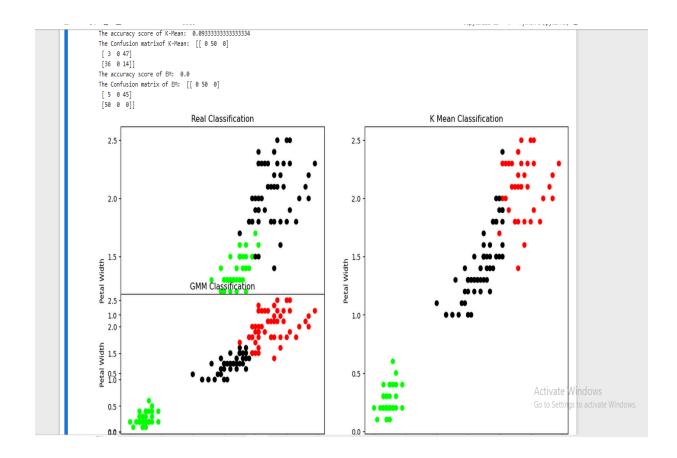
PROGREAM: 07

import matplotlib.pyplot as plt

from sklearn import datasets

```
from sklearn.cluster import KMeans
import sklearn.metrics as sm
import pandas as pd
import numpy as np
iris = datasets.load_iris()
X = pd.DataFrame(iris.data)
X.columns = ['Sepal_Length', 'Sepal_Width', 'Petal_Length', 'Petal_Width']
y = pd.DataFrame(iris.target)
y.columns = ['Targets']
model = KMeans(n_clusters=3)
model.fit(X)
plt.figure(figsize=(14,7))
colormap = np.array(['red', 'lime', 'black'])
plt.subplot(1, 2, 1)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y.Targets], s=40)
plt.title('Real Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
plt.subplot(1, 2, 2)
plt.scatter(X.Petal Length, X.Petal Width, c=colormap[model.labels ], s=40)
plt.title('K Mean Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('The accuracy score of K-Mean: ',sm.accuracy_score(y, model.labels_))
```

```
print('The Confusion matrixof K-Mean: ',sm.confusion_matrix(y, model.labels_))
from sklearn import preprocessing
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
from sklearn.mixture import GaussianMixture
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)
y_gmm = gmm.predict(xs)
plt.subplot(2, 2, 3)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y_gmm], s=40)
plt.title('GMM Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('The accuracy score of EM: ',sm.accuracy_score(y, y_gmm))
print('The Confusion matrix of EM: ',sm.confusion_matrix(y, y_gmm))
```



PROGREAM: 09

 $from \ sklearn. datasets \ import \ make_classification$

from sklearn import tree

 $from \ sklearn.model_selection \ import \ train_test_split$

X, t = make_classification(100, 5, n_classes=2, shuffle=True, random_state=10)

```
X_train, X_test, t_train, t_test = train_test_split(
X, t, test_size=0.3, shuffle=True, random_state=1)
model = tree.DecisionTreeClassifier()
model = model.fit(X train, t train)
predicted_value = model.predict(X_test)
print(predicted_value)
tree.plot_tree(model)
zeroes = 0
ones = 0
for i in range(0, len(t train)):
  if t train[i] == 0:
    zeroes += 1
  else:
    ones += 1
    print(zeroes)
    print(ones)
val = 1 - ((zeroes/70)*(zeroes/70) + (ones/70)*(ones/70))
print("Gini :", val)
match = 0
UnMatch = 0
for i in range(30):
  if predicted_value[i] == t_test[i]:
    match += 1
```

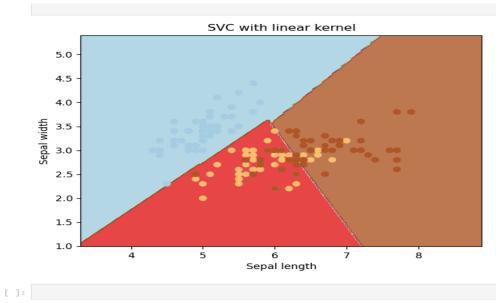
```
else:
    UnMatch += 1
    accuracy = match/30
    print("Accuracy is: ", accuracy)
```

```
import numpy as np
X = np.array(([2, 9], [1, 5], [3, 6]), dtype=float)
y = np.array(([92], [86], [89]), dtype=float)
X = X/np.amax(X,axis=0)
y = y/100
def sigmoid (x):
    return 1/(1 + np.exp(-x))
```

```
def derivatives sigmoid(x):
  return x * (1 - x)
epoch=5
lr=0.1
inputlayer neurons = 2
hiddenlayer neurons = 3
output neurons = 1
wh=np.random.uniform(size=(inputlayer_neurons,hiddenlayer_neurons))
bh=np.random.uniform(size=(1,hiddenlayer_neurons))
wout=np.random.uniform(size=(hiddenlayer neurons,output neurons))
bout=np.random.uniform(size=(1,output neurons))
for i in range(epoch):
  hinp1=np.dot(X,wh)
  hinp=hinp1 + bh
hlayer act = sigmoid(hinp)
outinp1=np.dot(hlayer act,wout)
outinp= outinp1+bout
output = sigmoid(outinp)
EO = y-output
outgrad = derivatives_sigmoid(output)
d output = EO * outgrad
EH = d output.dot(wout.T)
hiddengrad = derivatives_sigmoid(hlayer_act)
```

```
d_hiddenlayer = EH * hiddengrad
wout += hlayer_act.T.dot(d_output) *Ir
wh += X.T.dot(d_hiddenlayer) *Ir
print ("------Epoch-", i+1, "Starts ------")
print("Input: \n" + str(X))
print("Actual Output: \n" + str(y))
print("Predicted Output: \n" ,output)
print ("-------Epoch-", i+1, "Ends ------ \n")
print("Input: \n" + str(X))
print("Actual Output: \n" + str(y))
print("Actual Output: \n" + str(y))
print("Predicted Output: \n" ,output)
```

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn import svm, datasets
iris = datasets.load iris()
X = iris.data[:, :2]
y = iris.target
C = 1.0
svc = svm.SVC(kernel ='linear', C = 1).fit(X, y)
x \min_{x} \max = X[:, 0].\min() - 1, X[:, 0].\max() + 1
y min, y max = X[:, 1].min() - 1, X[:, 1].max() + 1
h = (x_max / x_min)/100
xx, yy = np.meshgrid(np.arange(x_min, x_max, h),
np.arange(y_min, y_max, h))
plt.subplot(1, 1, 1)
Z = svc.predict(np.c [xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.contourf(xx, yy, Z, cmap = plt.cm.Paired, alpha = 0.8)
plt.scatter(X[:, 0], X[:, 1], c = y, cmap = plt.cm.Paired)
plt.xlabel('Sepal length')
plt.ylabel('Sepal width')
plt.xlim(xx.min(), xx.max())
plt.title('SVC with linear kernel')
```



import pandas as pd

import numpy as np

from sklearn.linear_model import LogisticRegression

from sklearn.model_selection import train_test_split

import seaborn as sns

import matplotlib.pyplot as plt

from sklearn import metrics

Data= pd.read_csv(r"C:\divya ml\diabetes.csv")

Data.head().transpose()

Data.describe ()

[5]:		0.38076	0.05068	0.061696	0.021872	0.021872.1	-1.044223
	count	1.0	1.0	1.0	1.0	1.0	1.0
	mean	441.0	441.0	441.0	441.0	441.0	441.0
	std	NaN	NaN	NaN	NaN	NaN	NaN
	min	441.0	441.0	441.0	441.0	441.0	441.0
	25%	441.0	441.0	441.0	441.0	441.0	441.0
	50%	441.0	441.0	441.0	441.0	441.0	441.0
	75 %	441.0	441.0	441.0	441.0	441.0	441.0
	max	441.0	441.0	441.0	441.0	441.0	441.0