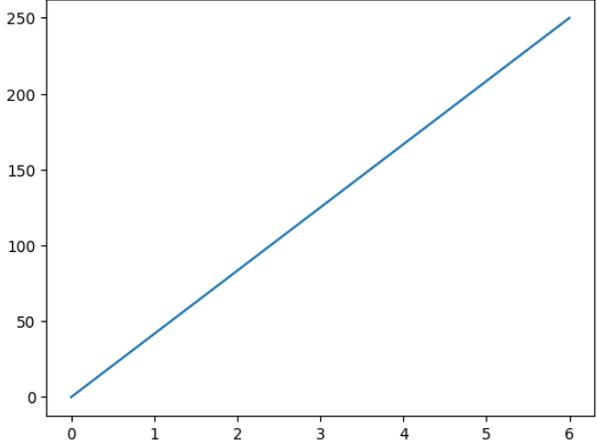




PROGRAMS

1. Demonstration of python Libraries for Machine Learning – pandas, Sklearn, numpy, matplotlib.

<pre>import matplotlib.pyplot as plt import numpy as np xpoints = np.array([0, 6]) ypoints = np.array([0, 250]) plt.plot(xpoints, ypoints) plt.show()</pre>	
<pre>import matplotlib.pyplot as plt import numpy as np xpoints = np.array([1, 2, 6, 8]) ypoints = np.array([3, 8, 1, 10]) plt.plot(xpoints, ypoints) plt.show()</pre>	<p>Machine Learning Laboratory</p>



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<pre>import numpy as np import matplotlib.pyplot as plt x = np.array([80, 85, 90, 95, 100, 105, 110, 115, 120, 125]) y = np.array([240, 250, 260, 270, 280, 290, 300, 310, 320, 330]) plt.plot(x, y) plt.title("Sports Watch Data") plt.xlabel("Average Pulse") plt.ylabel("Calorie Burnage") plt.show()</pre>	<table border="1"><caption>Sports Watch Data</caption><thead><tr><th>Average Pulse</th><th>Calorie Burnage</th></tr></thead><tbody><tr><td>80</td><td>240</td></tr><tr><td>85</td><td>250</td></tr><tr><td>90</td><td>260</td></tr><tr><td>95</td><td>270</td></tr><tr><td>100</td><td>280</td></tr><tr><td>105</td><td>290</td></tr><tr><td>110</td><td>300</td></tr><tr><td>115</td><td>310</td></tr><tr><td>120</td><td>320</td></tr><tr><td>125</td><td>330</td></tr></tbody></table>	Average Pulse	Calorie Burnage	80	240	85	250	90	260	95	270	100	280	105	290	110	300	115	310	120	320	125	330
Average Pulse	Calorie Burnage																						
80	240																						
85	250																						
90	260																						
95	270																						
100	280																						
105	290																						
110	300																						
115	310																						
120	320																						
125	330																						
<pre>import numpy as np a = np.array(42) b = np.array([1, 2, 3, 4, 5]) c = np.array([[1, 2, 3], [4, 5, 6]]) d = np.array([[[1, 2, 3], [4, 5, 6]], [[1, 2, 3], [4, 5, 6]]]) print(a.ndim) print(b.ndim) print(c.ndim) print(d.ndim)</pre>	<pre>0 1 2 3</pre>																						



2. Demonstration of Exploratory Data Analysis and Data Visualization.

<pre>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) print("X_train Shape:", X_train.shape) print("X_test Shape:", X_test.shape) print("Y_train Shape:", y_train.shape) print("Y_test Shape:", y_test.shape)</pre>	<pre>X_train Shape: (90, 4) X_test Shape: (60, 4) Y_train Shape: (90,) Y_test Shape: (60,)</pre>
<pre>from sklearn.preprocessing import LabelEncoder categorical_feature = ['cat', 'dog', 'dog', 'cat', 'bird'] encoder = LabelEncoder() encoded_feature = encoder.fit_transform(categorical_feature) print("Encoded feature:", encoded_feature)</pre>	<pre>Encoded feature: [1 2 2 1 0]</pre>



3. Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from av.CSV file.

```
import csv

with open('lab1a.csv', 'r') as f:

    reader = csv.reader(f)
    your_list = list(reader)

h = [['0', '0', '0', '0', '0', '0']]

for i in your_list:
    print(i)
    if i[-1] == "True":
        j = 0
        for x in i:
            if x != "True":
                if x != h[0][j] and h[0][j] == '0':
                    h[0][j] = x
                elif x != h[0][j] and h[0][j] != '0':
                    h[0][j] = '?'
                else:
                    pass
            j = j + 1
print("Most specific hypothesis is")
print(h)
```

OUTPUT:



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```
['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same', 'TRUE']  
['Sunny', 'Warm', 'High', 'Strong', 'Warm', 'Same', 'TRUE']  
['Rainy', 'Cold', 'High', 'Strong', 'Warm', 'Change', 'FALSE']  
['Sunny', 'Warm', 'High', 'Strong', 'Cool', 'Change', 'TRUE']  
Most specific hypothesis is  
[['Sunny', 'Warm', '?', 'Strong', '?', '?']]
```



4. For a given set of training data examples stored in a .CSV file, implement and demonstrate the **Candidate-Elimination algorithm** to output a description of the set of all hypotheses consistent with the training examples.

```
import numpy as np
import pandas as pd

# Loading Data from a CSV File
data = pd.DataFrame(data=pd.read_csv('2b.csv'))

# Separating concept features from Targetconcepts =
np.array(data.iloc[:,0:-1])

# Isolating target into a separate DataFrame#copying last
column to target array
target = np.array(data.iloc[:,-1])

def learn(concepts, target):
    #learn() function implements the learning method of the Candidateelimination algorithm.

    #Arguments:
    #concepts - a data frame with all the features
    #target - a data frame with corresponding output values

    # Initialise S0 with the first instance from concepts
    # .copy() makes sure a new list is created instead of justpointing to the same
memory location
    specific_h = concepts[0].copy() print("initialization of specific_h and
general_h")print(specific_h)
    general_h = [{"?" for i in range(len(specific_h))] for i inrange(len(specific_h))]
    print(general_h)
    # The learning iterations
    for i, h in enumerate(concepts):

        # Checking if the hypothesis has a positive target
```



```

if target[i] == "Yes":
    for x in range(len(specific_h)):

        # Change values in S & G only if values change if h[x] !=
        specific_h[x]:
            specific_h[x] = '?'
            general_h[x][x] = '?'

# Checking if the hypothesis has a positive target if target[i] == "No":
for x in range(len(specific_h)):

    # For negative hypothesis change values only in G if h[x] !=
    specific_h[x]:
        general_h[x][x] = specific_h[x] else:
            general_h[x][x] = '?'
print(" steps of Candidate Elimination Algorithm",i+1)print("Specific_h ",i+1,"\n ")
print(specific_h) print("general_h ", i+1,
"\n ")print(general_h)
# find indices where we have empty rows, meaning those that are unchanged
indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
for i in indices:
    # remove those rows from general_h
    general_h.remove(['?', '?', '?', '?', '?', '?'])

# Return final values
return
specific_h, general_h
s_final, g_final = learn(concepts, target) print("Final
Specific_h:", s_final, sep="\n")print("Final General_h:",
g_final, sep="\n")

```



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OUTPUT:

```

candidate - Jupyter Notebook - Mozilla Firefox
Home Page - Select or c... candidate - Jupyter Not...
localhost:8888/notebooks/candidate.ipynb
jupyter candidate Last Checkpoint: 5 minutes ago (unsaved changes)
Logout
File Edit View Insert Cell Kernel Widgets Help
Python 3
initialization of specific h and general h
['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
steps of Candidate Elimination Algorithm 1
Specific_h 1
['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
general_h 1
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
steps of Candidate Elimination Algorithm 2
Specific_h 2
['Sunny' 'Warm' '?' 'Strong' 'Warm' 'Same']
general_h 2
[['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
steps of Candidate Elimination Algorithm 3
Specific_h 3
['Sunny' 'Warm' '?' 'Strong' 'Warm' 'Same']
general_h 3
[['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
steps of Candidate Elimination Algorithm 4
Specific_h 4
['Sunny' 'Warm' '?' 'Strong' '?' '?']
general_h 4

```



5. Write a program to implement k -Nearest Neighbour algorithm to classify the iris data set. Print both correct and wrong predictions.

```
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import confusion_matrix, classification_report
from sklearn import datasets
iris=datasets.load_iris()
iris_data=iris.data
iris_labels=iris.target
print(iris_data)
x_train,X_test,Y_train,Y_test=train_test_split(iris_data,iris_labels,test_size=0.20)
classifier=KNeighborsClassifier(n_neighbors=5)
classifier.fit(x_train,Y_train)
y_prd=classifier.predict(X_test)
print(confusion_matrix(Y_test,y_prd))
print(classification_report(Y_test,y_prd))
```



OUTPUT:

```
print(confusion_matrix(Y_test,y_prd))
print(classification_report(Y_test,y_prd))
```

```
[[ 6.7 3.3 5.7 2.5]
 [ 6.7 3.  5.2 2.3]
 [ 6.3 2.5 5.  1.9]
 [ 6.5 3.  5.2 2. ]
 [ 6.2 3.4 5.4 2.3]
 [ 5.9 3.  5.1 1.8]]
```

```
[[ 8 0 0]
 [ 0 9 0]
 [ 0 1 12]]
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	8
1	0.90	1.00	0.95	9
2	1.00	0.92	0.96	13
accuracy			0.97	30
macro avg	0.97	0.97	0.97	30
weighted avg	0.97	0.97	0.97	30

```
print(iris_data)
x_train,X_test,Y_train,Y_test=train_test_split(iris_data,iris_labels,test_size=0.20)
classifier=KNeighborsClassifier(n_neighbors=5)
classifier.fit(x_train,Y_train)
y_prd=classifier.predict(X_test)
print(confusion_matrix(Y_test,y_prd))
print(classification_report(Y_test,y_prd))
```

```
[[5.1 3.5 1.4 0.2]
 [4.9 3.  1.4 0.2]
 [4.7 3.2 1.3 0.2]
 [4.6 3.1 1.5 0.2]
 [5.  3.6 1.4 0.2]
 [5.4 3.9 1.7 0.4]
 [4.6 3.4 1.4 0.3]
 [5.  3.4 1.5 0.2]
 [4.4 2.9 1.4 0.2]
 [4.9 3.1 1.5 0.1]]
```



- 6. Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use an appropriate data set for building the decision tree and apply this knowledge to classify a new sample.**

```
#Implementing Decision Tree #numpy and
pandas initialization

import numpy as np
import pandas as pd

#Loading the PlayTennis data

PlayTennis = pd.read_csv("PlayTennis.csv")
PlayTennis

#It is easy to implement Decision Tree with numerical values.
#We can convert all the non numerical values into numerical values using LabelEncoder

from sklearn.preprocessing import LabelEncoder
Le = LabelEncoder()

PlayTennis['outlook'] = Le.fit_transform(PlayTennis['outlook'])

PlayTennis['temp'] = Le.fit_transform(PlayTennis['temp'])
PlayTennis['humidity'] =
Le.fit_transform(PlayTennis['humidity'])
PlayTennis['windy'] =
Le.fit_transform(PlayTennis['windy'])
PlayTennis['play'] =
Le.fit_transform(PlayTennis['play'])

PlayTennis

#Lets split the training data and its corresponding prediction values. #y - holds all the decisions.
#X - holds the training data.

y = PlayTennis['play']
X = PlayTennis.drop(['play'],axis=1)
```



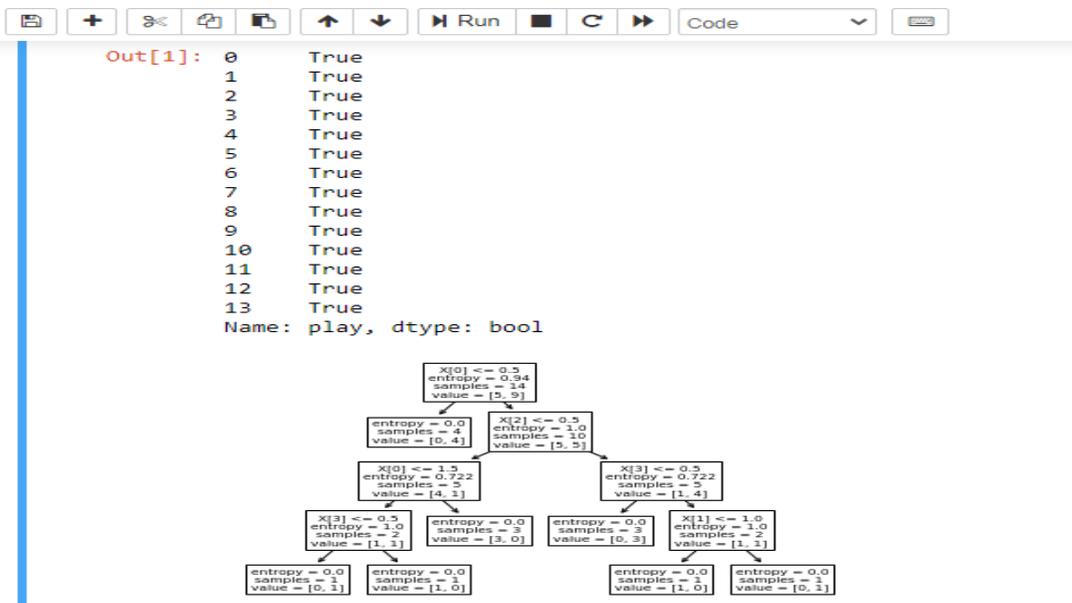
```
# Fitting the model from
sklearn import tree
clf = tree.DecisionTreeClassifier(criterion = 'entropy')clf = clf.fit(X, y)
```

```
# We can visualize the tree using tree.plot_treetree.plot_tree(clf)
```

```
#In Graph
#X[0] -> Outlook #X[1] ->
Temperature#X[2] ->
Humidity #X[3] -> Wind
```

```
# The predictions are stored in X_predX_pred =
clf.predict(X)
# verifying if the model has predicted it all right.X_pred == y
```

OUTPUT:





7. Implement a Clustering algorithm using K-means clustering for the dataset.

```
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.cluster import KMeans
import pandas as pd
import numpy as np

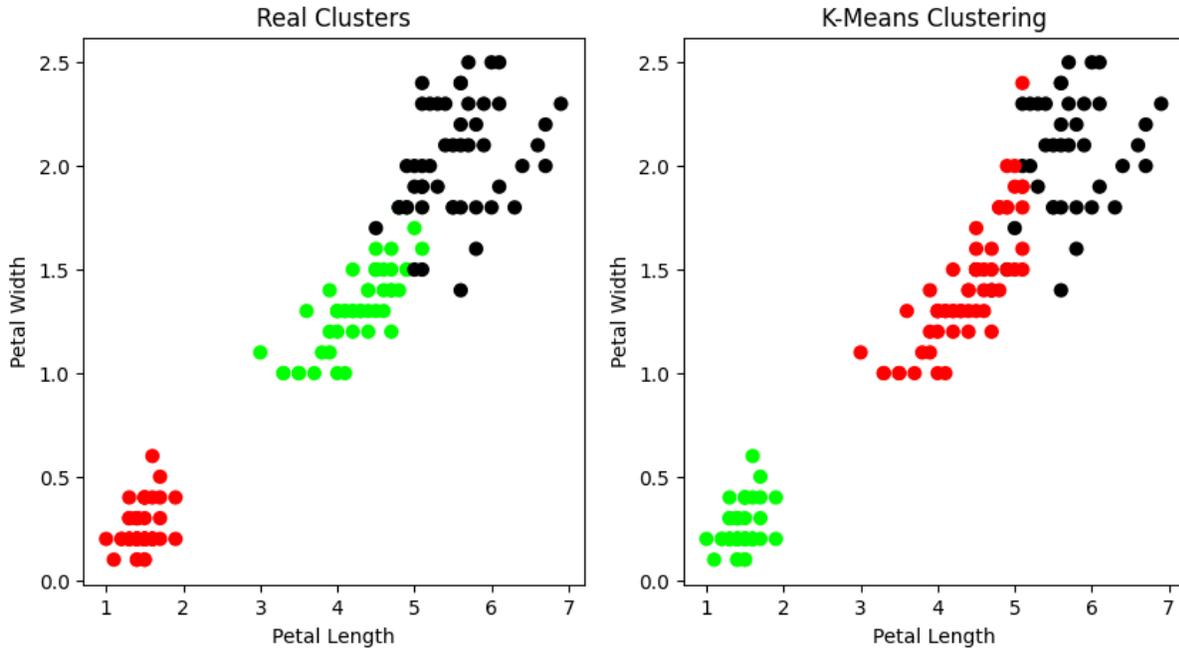
# import some data to play with
iris = datasets.load_iris()
X = pd.DataFrame(iris.data)
X.columns = ['Sepal_Length','Sepal_Width','Petal_Length','Petal_Width']
y = pd.DataFrame(iris.target)
y.columns = ['Targets']

# Build the K Means Model
model = KMeans(n_clusters=3)
model.fit(X) # model.labels_ : Gives cluster no for which samples belongs to

## Visualise the clustering results
plt.figure(figsize=(10,5))
colormap = np.array(['red', 'lime', 'black'])
# Plot the Original Classifications using Petal features
plt.subplot(1, 2, 1)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y.Targets], s=40)
plt.title('Real Clusters')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
# Plot the Models Classifications
plt.subplot(1, 2, 2)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[model.labels_], s=40)
plt.title('K-Means Clustering')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
plt.show()
```



OUTPUT:





8. Build an Artificial Neural Network by implementing the Backpropagation algorithm and test these using appropriate data sets.

```
import numpy as np
import matplotlib as m

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=7000
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)*lr
wh+=X.T.dot(d_hiddenlayer)*lr
print("Input:\n"+str(X)) print("Actual
Output:\n"+str(y)) print("Predicted
Output:\n",output)
```

OUTPUT:

The screenshot shows a Jupyter Notebook interface in Mozilla Firefox. The browser address bar shows 'localhost:8888/notebooks/Untitled2.ipynb?kernel...'. The notebook title is 'Untitled2' and it indicates 'Last Checkpoint: a minute ago (unsaved changes)'. The menu bar includes File, Edit, View, Insert, Cell, Kernel, Widgets, and Help. The toolbar contains icons for file operations and execution. The code cell contains the following Python code:

```
d_output=d_output
EH=d_output.dot(wout.T)
hiddengrad=derivatives_sigmoid(hlayer_act)
d_hiddenlayer=EH*hiddengrad
wout+=hlayer_act.T.dot(d_output)*lr
wh+=X.T.dot(d_hiddenlayer)*lr
print("Input:\n"+str(X))
print("Actual Output:\n"+str(y))
print("Predicted Output:\n",output)
```

The output of the code is displayed below the cell:

```
Input:
[[0.66666667 1.          ]
 [0.33333333 0.55555556]
 [1.          0.66666667]]
Actual Output:
[[0.92]
 [0.86]
 [0.89]]
Predicted Output:
[[0.89687109]
 [0.8806334 ]
 [0.89206686]]
```



9. Write a program to implement the naïve Bayesian classifier for a sample training data set stored as a .CSV file. Compute the accuracy of the classifier, considering few test datasets.

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn import metrics

df = pd.read_csv("pima_indian.csv")
feature_col_names = ['num_preg', 'glucose_conc', 'diastolic_bp', 'thickness', 'insulin', 'bmi', 'diab_pred', 'age']
predicted_class_names = ['diabetes']

X = df[feature_col_names].values # these are factors for the prediction
y = df[predicted_class_names].values # this is what we want to predict

#splitting the dataset into train and test data

xtrain,xtest,ytrain,ytest=train_test_split(X,y,test_size=0.33)

print ('\n the total number of Training Data :',ytrain.shape)
print ('\n the total number of Test Data :',ytest.shape)

# Training Naive Bayes (NB) classifier on training data.

clf = GaussianNB().fit(xtrain,ytrain.ravel())
predicted = clf.predict(xtest)
predictTestData= clf.predict([[6,148,72,35,0,33.6,0.627,50]])

#printing Confusion matrix, accuracy, Precision and Recall

print('\n Confusion matrix')
print(metrics.confusion_matrix(ytest,predicted))

print('\n Accuracy of the classifier is',metrics.accuracy_score(ytest,predicted))
```



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```
print('\n The value of Precision', metrics.precision_score(ytest,predicted))  
print('\n The value of Recall', metrics.recall_score(ytest,predicted))  
print("Predicted Value for individual Test Data:", predictTestData)
```

OUTPUT:

```
the total number of Training Data : (514, 1)  
the total number of Test Data : (254, 1)  
  
Confusion matrix  
[[133  32]  
 [ 32  57]]  
  
Accuracy of the classifier is 0.7480314960629921  
  
The value of Precision 0.6404494382022472  
  
The value of Recall 0.6404494382022472  
Predicted Value for individual Test Data: [1]
```