

VISVESVARAYA TECHNOLOGICAL UNIVERSITY

“JnanaSangama”, Belgaum -590014, Karnataka.



LAB REPORT

on

MACHINE LEARNING (20CS6PCMAL)

Submitted by

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in partial fulfillment for the award of the degree of

BACHELOR OF ENGINEERING

in

COMPUTER SCIENCE AND ENGINEERING



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C E R T I F I C A T E

This is to certify that the Lab work entitled “**MACHINE LEARNING**” carried out by **Sudeshna Bhushan (1BM19CS189)**, who is a bonafide student of **B. M. S. College of Engineering**. It is in partial fulfillment for the award of **Bachelor of Engineering in Computer Science and Engineering** of the Visvesvaraya Technological University, Belgaum during the year 2022. The Lab report has been approved as it satisfies the academic requirements in respect of a Machine Learning - (20CS6PCMAL) work prescribed for the said degree.

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Course Outcome

CO1	Ability to apply the different learning algorithms.
CO2	Ability to analyse the learning techniques for a given dataset.
CO3	Ability to design a model using machine learning to solve a problem.
CO4	Ability to conduct practical experiments to solve problems using appropriate machine learning techniques.

1. Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples.

```
In [1]: import csv

def findS(dataset, hypothesis):
    for i in range(len(dataset)):
        if dataset[i][-1] == 'yes':
            print('The tuple', i+1, 'is a positive instance.')
            for j in range(len(hypothesis)):
                if hypothesis[j] == '0' or dataset[i][j] == hypothesis[j]:
                    hypothesis[j] = dataset[i][j]
            else:
                hypothesis[j] = '?'
            print('The hypothesis for training tuple', i+1, 'and instance', j+1, 'is:', hypothesis)
        elif dataset[i][-1] == 'no':
            print('The tuple', i+1, 'is a negative instance.')
            print('The hypothesis for training tuple', i+1, 'is:', hypothesis)
    return hypothesis

def main():
    dataset = []
    with open('FindS-CSV.csv', 'r') as csvfile:
        next(csvfile)
        for row in csv.reader(csvfile):
            dataset.append(row)
    print(dataset)
    hypothesis = ['0']*len(dataset[0])
    print('The Initial hypothesis:', hypothesis)
    hypothesis = findS(dataset, hypothesis)
    print('Final Hypothesis: ', hypothesis)

if __name__ == "__main__":
    main()

[['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes'], ['sunny', 'warm', 'high', 'strong', 'warm', 'same', 'yes'], ['rainy', 'cold', 'high', 'strong', 'warm', 'change', 'no'], ['sunny', 'warm', 'high', 'strong', 'cool', 'change', 'yes']]
The Initial hypothesis: ['0', '0', '0', '0', '0', '0', '0']
The tuple 1 is a positive instance.
The hypothesis for training tuple 1 and instance 7 is: ['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes']
The tuple 2 is a positive instance.
The hypothesis for training tuple 2 and instance 7 is: ['sunny', 'warm', '?', 'strong', 'warm', 'same', 'yes']
The tuple 3 is a negative instance.
The hypothesis for training tuple 3 is: ['sunny', 'warm', '?', 'strong', 'warm', 'same', 'yes']
The tuple 4 is a positive instance.
The hypothesis for training tuple 4 and instance 7 is: ['sunny', 'warm', '?', 'strong', '?', '?', 'yes']
Final Hypothesis: ['sunny', 'warm', '?', 'strong', '?', '?', 'yes']
```

2. 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.

In [1]:

```
import pandas as pd
import numpy as np
import csv

data = pd.read_csv('Candidate-Elimination.csv')
d = np.array(data.iloc[:,0:-1])
print("\nInstances are:\n",d)
target = np.array(data.iloc[:, -1])
print("\nTarget Values are: ",target)

Instances are:
[['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
 ['sunny' 'warm' 'high' 'strong' 'warm' 'same']
 ['rainy' 'cold' 'high' 'strong' 'warm' 'change']
 ['sunny' 'warm' 'high' 'strong' 'cool' 'change']]

Target Values are: ['yes' 'yes' 'no' 'yes']
```

In [2]:

```
def learn(d, target):
    specific_h = d[0].copy()
    print("\nSpecific Hypothesis: ", specific_h)
    general_h = [{"?" for i in range(len(specific_h))}]
    print("\nGeneric Hypothesis: ", general_h)

    for i, h in enumerate(d):
        print("\nIteration", i+1, "is ", h)
        if target[i] == "yes":
            print("Instance is Positive ")
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    specific_h[x] = '?'
                    general_h[x][x] = '?'

        if target[i] == "no":
            print("Instance is Negative ")
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    general_h[x][x] = specific_h[x]
            else:
                general_h[x][x] = '?'

        print("Specific Hypothesis after ", i+1, "Instance is ", specific_h)
        print("Generic Hypothesis after ", i+1, "Instance is ", general_h)
        print("\n")

    indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
    for i in indices:
        general_h.remove(['?', '?', '?', '?', '?', '?'])
    return specific_h, general_h
```

In [3]:

```
specific, general = learn(d, target)

print("Final Specific Hypothesis: ", '<', ' '.join(specific), '>')
print("Final General Hypothesis: ")
for i in general:
    print('<', ' '.join(i), '>', ' ')

Specific Hypothesis: ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']

Generic Hypothesis: [['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Iteration 1 is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
Instance is Positive
Specific Hypothesis after 1 Instance is ['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
Generic Hypothesis after 1 Instance is [['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Iteration 2 is ['sunny' 'warm' 'high' 'strong' 'warm' 'same']
Instance is Positive
Specific Hypothesis after 2 Instance is ['sunny' 'warm' '?' 'strong' 'warm' 'same']
Generic Hypothesis after 2 Instance is [['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Iteration 3 is ['rainy' 'cold' 'high' 'strong' 'warm' 'change']
Instance is Negative
Specific Hypothesis after 3 Instance is ['sunny' 'warm' '?' 'strong' 'warm' 'same']
Generic Hypothesis after 3 Instance is [['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Iteration 4 is ['sunny' 'warm' 'high' 'strong' 'cool' 'change']
Instance is Positive
Specific Hypothesis after 4 Instance is ['sunny' 'warm' '?' 'strong' '?', '?']
Generic Hypothesis after 4 Instance is [['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Final Specific Hypothesis: < sunny, warm, ?, strong, ?, ? >
Final General Hypothesis:
< sunny, ?, ?, ?, ?, ? >,
< ?, warm, ?, ?, ?, ? >
```

3. 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.

```
In [4]: def print_tree(node, level):
        if node.answer != "":
            print(" *level, node.answer)
            return

        print(" *level, node.attribute)
        for value, n in node.children:
            print(" *(level+1), value)
            print_tree(n, level+2)

        def classify(node, x_test, features):
            if node.answer != "":
                print(node.answer)
                return
            pos = features.index(node.attribute)
            for value, n in node.children:
                if x_test[pos] == value:
                    classify(n, x_test, features)
```

```
In [5]: dataset, features = load_csv("id3.csv")
        model = build_tree(dataset, features)

        print("The decision tree for the dataset using ID3 algorithm is")
        print_tree(model, 0)
        testdata, features = load_csv("id3.csv")

        for xtest in testdata:
            print("The test instance:", xtest)
            print("The label for test instance:", end=" ")
            classify(model, xtest, features)
```

```
The decision tree for the dataset using ID3 algorithm is
Outlook
  overcast
    yes
  sunny
    Humidity
      high
      no
      normal
      yes
  rain
    Wind
      strong
      no
      weak
      yes
The test instance: ['sunny', 'hot', 'high', 'weak', 'no']
The label for test instance: no
The test instance: ['sunny', 'hot', 'high', 'strong', 'no']
The label for test instance: no
The test instance: ['overcast', 'hot', 'high', 'weak', 'yes']
The label for test instance: yes
The test instance: ['rain', 'mild', 'high', 'weak', 'yes']
The label for test instance: yes
The test instance: ['rain', 'cool', 'normal', 'weak', 'yes']
The label for test instance: yes
The test instance: ['rain', 'cool', 'normal', 'strong', 'no']
The label for test instance: no
The test instance: ['overcast', 'cool', 'normal', 'strong', 'yes']
The label for test instance: yes
The test instance: ['sunny', 'mild', 'high', 'weak', 'no']
The label for test instance: no
The test instance: ['sunny', 'cool', 'normal', 'weak', 'yes']
The label for test instance: yes
The test instance: ['rain', 'mild', 'normal', 'weak', 'yes']
The label for test instance: yes
The test instance: ['sunny', 'mild', 'normal', 'strong', 'yes']
The label for test instance: yes
The test instance: ['overcast', 'mild', 'high', 'strong', 'yes']
The label for test instance: yes
The test instance: ['overcast', 'hot', 'normal', 'weak', 'yes']
The label for test instance: yes
The test instance: ['rain', 'mild', 'high', 'strong', 'no']
The label for test instance: no
```

4. 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 data sets

```
In [1]: import numpy as np
import pandas as pd

In [2]: data = pd.read_csv('/content/dataset.csv')
data.head()

Out[2]:
```

	PlayTennis	Outlook	Temperature	Humidity	Wind
0	No	Sunny	Hot	High	Weak
1	No	Sunny	Hot	High	Strong
2	Yes	Overcast	Hot	High	Weak
3	Yes	Rain	Mild	High	Weak
4	Yes	Rain	Cool	Normal	Weak

```

In [3]: y = list(data['PlayTennis'].values)
X = data.iloc[:,1:].values
print(f'Target Values: {y}')
print(f'Features: \n{X}')

Target Values: ['No', 'No', 'Yes', 'Yes', 'Yes', 'No', 'Yes', 'No', 'Yes', 'No']
Features:
[['Sunny', 'Hot', 'High', 'Weak']
 ['Sunny', 'Hot', 'High', 'Strong']
 ['Overcast', 'Hot', 'High', 'Weak']
 ['Rain', 'Mild', 'High', 'Weak']
 ['Rain', 'Cool', 'Normal', 'Weak']
 ['Rain', 'Cool', 'Normal', 'Strong']
 ['Overcast', 'Cool', 'Normal', 'Strong']
 ['Sunny', 'Mild', 'High', 'Weak']
 ['Sunny', 'Cool', 'Normal', 'Weak']
 ['Rain', 'Mild', 'Normal', 'Weak']
 ['Sunny', 'Mild', 'Normal', 'Strong']
 ['Overcast', 'Mild', 'High', 'Strong']
 ['Overcast', 'Hot', 'Normal', 'Weak']
 ['Rain', 'Mild', 'High', 'Strong']]

File display

In [4]: y_train = y[:8]
y_val = y[8:]
X_train = X[:8]
X_val = X[8:]
print(f'Number of instances in training set: {len(X_train)}')
print(f'Number of instances in testing set: {len(X_val)}')

Number of instances in training set: 8
Number of instances in testing set: 6

In [5]: class NaiveBayesClassifier:
def __init__(self, X, y):
    self.X, self.y = X, y
    self.N = len(self.X)
    self.dim = len(self.X[0])
    self.attrs = [[] for _ in range(self.dim)]
    self.output_dom = {}
    self.data = []
    for i in range(len(self.X)):
        for j in range(self.dim):
            if not self.X[i][j] in self.attrs[j]:
                self.attrs[j].append(self.X[i][j])
            if not self.y[i] in self.output_dom.keys():
                self.output_dom[self.y[i]] = 1
            else:
                self.output_dom[self.y[i]] += 1
    self.data.append((self.X[i], self.y[i]))
def classify(self, entry):
    solve = None
    max_arg = -1
    for y in self.output_dom.keys():
        prob = self.output_dom[y]/self.N
        for i in range(self.dim):
            cases = [x for x in self.data if x[0][i] == entry[i] and x[1] == y]
            n = len(cases)
            prob *= n/self.N
        if prob > max_arg:
            max_arg = prob
            solve = y
    return solve

In [6]: nbc = NaiveBayesClassifier(X_train, y_train)
total_cases = len(y_val)
good = 0
bad = 0
predictions = []
for i in range(total_cases):
    predict = nbc.classify(X_val[i])
    predictions.append(predict)
    if y_val[i] == predict:
        good += 1
    else:
        bad += 1
print('Predicted values:', predictions)
print('Actual values:', y_val)
print()
print('Total number of testing instances in the dataset:', total_cases)
print('Number of correct predictions:', good)
print('Number of wrong predictions:', bad)
print()
print('Accuracy of Bayes Classifier:', good/total_cases)

Predicted values: ['No', 'Yes', 'No', 'Yes', 'Yes', 'No']
Actual values: ['Yes', 'Yes', 'Yes', 'Yes', 'Yes', 'No']

Total number of testing instances in the dataset: 6
Number of correct predictions: 4
Number of wrong predictions: 2

Accuracy of Bayes Classifier: 0.6666666666666666
```


5. Write a program to construct a Bayesian network considering training data. Use this model to make predictions.

```
In [7]: import numpy as np
import pandas as pd
import csv
```

```
In [8]: from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
```

```
In [9]: heartDisease = pd.read_csv('heart.csv')
heartDisease = heartDisease.replace('?', np.nan)

print('Sample instances from the dataset are given below')
print(heartDisease.head())
```

```
Sample instances from the dataset are given below
   age  sex  cp  trestbps  chol  fbs  restecg  thalach  exang  oldpeak  slope  \
0   63    1    1     145    233     1         2     150      0        2.3      3
1   67    1    4     160    286     0         2     108      1        1.5      2
2   67    1    4     120    229     0         2     129      1        2.6      2
3   37    1    3     130    250     0         0     187      0        3.5      3
4   41    0    2     130    204     0         2     172      0        1.4      1

   ca  thal  heartdisease
0   0     6           0
1   3     3           2
2   2     7           1
3   0     3           0
4   0     3           0
```

```
In [10]: print('\n Attributes and datatypes')
print(heartDisease.dtypes)
```

```
Attributes and datatypes
age                int64
sex                int64
cp                int64
trestbps           int64
chol               int64
fbs               int64
restecg            int64
thalach            int64
exang              int64
oldpeak            float64
slope              int64
ca                 object
thal               object
heartdisease       int64
dtype: object
```

```
In [11]: model = BayesianModel([('age', 'heartdisease'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'heartdisease'), ('heartdisease', 'restecg')])
print('\n Learning CPD using Maximum likelihood estimators')
model.fit(heartDisease, estimator=MaximumLikelihoodEstimator)

print('\n Inferencing with Bayesian Network:')
```

```
Learning CPD using Maximum likelihood estimators

Inferencing with Bayesian Network:
```

```
In [12]: HeartDiseasetest_infer = VariableElimination(model)

print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)
```

```
Finding Elimination Order: : 100% | 5/5 [00:00<00:00, 2500.78it/s]
Eliminating: chol: 100% | 5/5 [00:00<00:00, 185.63it/s]
1. Probability of HeartDisease given evidence= restecg
+-----+-----+
| heartdisease | phi(heartdisease) |
+-----+-----+
| heartdisease(0) | 0.1012 |
+-----+-----+
| heartdisease(1) | 0.0000 |
+-----+-----+
| heartdisease(2) | 0.2392 |
+-----+-----+
| heartdisease(3) | 0.2015 |
+-----+-----+
| heartdisease(4) | 0.4581 |
+-----+-----+
```

```
In [13]: print('\n 2. Probability of HeartDisease given evidence= cp ')
q2=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

```
Finding Elimination Order: : 100% | 5/5 [00:00<00:00, 2507.06it/s]
Eliminating: restecg: 100% | 5/5 [00:00<00:00, 179.06it/s]
2. Probability of HeartDisease given evidence= cp
+-----+-----+
| heartdisease | phi(heartdisease) |
+-----+-----+
| heartdisease(0) | 0.3610 |
+-----+-----+
| heartdisease(1) | 0.2159 |
+-----+-----+
| heartdisease(2) | 0.1373 |
+-----+-----+
| heartdisease(3) | 0.1537 |
+-----+-----+
| heartdisease(4) | 0.1321 |
+-----+-----+
```


6. Apply k-Means algorithm to cluster a set of data stored in a .CSV

```
In [2]: from sklearn.preprocessing import StandardScaler
        from sklearn.cluster import KMeans
        import pandas as pd
        import numpy as np
        from itertools import cycle, islice

In [3]: from google.colab import drive
        drive.mount("/content/drive")

Mounted at /content/drive

In [4]: data = pd.read_csv('/content/drive/MyDrive/minute_weather.csv')

In [5]: data.shape

Out[5]: (1587257, 13)

In [6]: data.head()

Out[6]:
```

	rowID	hpwren_timestamp	air_pressure	air_temp	avg_wind_direction	avg_wind_speed	max_wind_direction	max_wind_speed	min_wind_direction	min_wind_speed
0	0	2011-09-10 00:00:49	912.3	64.76	97.0	1.2	106.0	1.6	85.0	1.0
1	1	2011-09-10 00:01:49	912.3	63.86	161.0	0.8	215.0	1.5	43.0	0.2
2	2	2011-09-10 00:02:49	912.3	64.22	77.0	0.7	143.0	1.2	324.0	0.3
3	3	2011-09-10 00:03:49	912.3	64.40	89.0	1.2	112.0	1.6	12.0	0.7
4	4	2011-09-10 00:04:49	912.3	64.40	185.0	0.4	260.0	1.0	100.0	0.1

```
In [7]: #data sampling
        sampled_df = data[(data['rowID'] % 10) == 0]
        sampled_df.shape

Out[7]: (158726, 13)

In [8]: sampled_df.describe().transpose()

Out[8]:
```

	count	mean	std	min	25%	50%	75%	max
rowID	158726.0	793625.000000	458203.937509	0.00	396812.5	793625.00	1190437.50	1587250.00
air_pressure	158726.0	916.830161	3.051717	905.00	914.8	916.70	918.70	929.50
air_temp	158726.0	61.851589	11.833569	31.64	52.7	62.24	70.88	99.50
avg_wind_direction	158680.0	162.156100	95.278201	0.00	62.0	182.00	217.00	359.00
avg_wind_speed	158680.0	2.775215	2.057624	0.00	1.3	2.20	3.80	31.90
max_wind_direction	158680.0	163.462144	92.452139	0.00	68.0	187.00	223.00	359.00
max_wind_speed	158680.0	3.400558	2.418802	0.10	1.6	2.70	4.60	36.00
min_wind_direction	158680.0	166.774017	97.441109	0.00	76.0	180.00	212.00	359.00
min_wind_speed	158680.0	2.134664	1.742113	0.00	0.8	1.60	3.00	31.60
rain_accumulation	158725.0	0.000318	0.011236	0.00	0.0	0.00	0.00	3.12
rain_duration	158725.0	0.409627	8.665523	0.00	0.0	0.00	0.00	2960.00
relative_humidity	158726.0	47.609470	26.214409	0.90	24.7	44.70	68.00	93.00

```
In [9]: sampled_df[sampled_df['rain_accumulation'] == 0].shape

Out[9]: (157812, 13)

In [10]: sampled_df[sampled_df['rain_duration'] == 0].shape

Out[10]: (157237, 13)

In [11]: del sampled_df['rain_accumulation']
        del sampled_df['rain_duration']

In [12]: rows_before = sampled_df.shape[0]
        sampled_df = sampled_df.dropna()
        rows_after = sampled_df.shape[0]
```

file.

In [13]:

Dut[13]: 46

In [14]: sampled_df.columns

In [15]: features = ['air_pressure', 'air_temp', 'avg_wind_direction', 'avg_wind_speed', 'max_wind_direction', 'max_wind_speed', 'relative_humidity']

In [17]:

In [18]: select df

Dut[18]:

air_pressure	air_temp	avg_wind_direction	avg_wind_speed	max_wind_direction	max_wind_speed	relative_humidity
--------------	----------	--------------------	----------------	--------------------	----------------	-------------------

10	912.3	62.24	144.0	1.2	167.0	1.8	38.6
20	912.2	63.32	100.0	2.0	122.0	2.6	68.2

US

ad10

d
7
8

1587210 915.9 75.56 2aD.D

1.3

1S0BB0 s x 7 columns

In [19]:

Dut[19]: array([[[-1.48456281, 0.24544455, -0.68385323, ..., -0.62153592, ..., -1.48456281, 0.03247142, -0.19055941, ..., 0.03826701, -0.66171726, -0.34710804], [-1.51733167, 0.12374562, -0.65236639, ..., -0.44847286, ...

[-0.30488381, 1.15619654, 1.9dF5SJ2d, ..., 2.0393087, -0.01538018], -0.67791749, -0.63899948, -1.67791749, -0.62037434, -0.69711347]]])

mnNm1 - c.*it[z]

[-0.7245782, 0.51194369, 0.17191124, -0.58229578, 0.34128394,

0.67243274, -0.15J29I94].

1.98322667, 0.53830956,

-0.63899948, -1.67791749,

```

[-0.16372869, 0.86324348, -1.31172732,
-0.6047306 , -0.64119682],
[ 0.25182364, -0.99684608, 0.45839645, -0.58942801, -1.16773268,
-6.1248 +112i, 1.1*621623],
[ 0.23422959, 0.32038874, 1.88815273, -0.65179307,

[ 0.48752537, 0.48036551, 0.28249096,
-0.54507948, -0.76332259],
[-.59D42211, -1.20432314, 0.37675641,

[ 1.36796382, -0.08175169, -1.20532878, -0.05333267, -1.073853 ,

```

•.Z51t2ES6.

7. Apply EM algorithm to cluster a set of data stored in a .CSV file. Compare the results of k-Means algorithm and EM algorithm.

```
In [1]: 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
```

```
In [2]: 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']
```

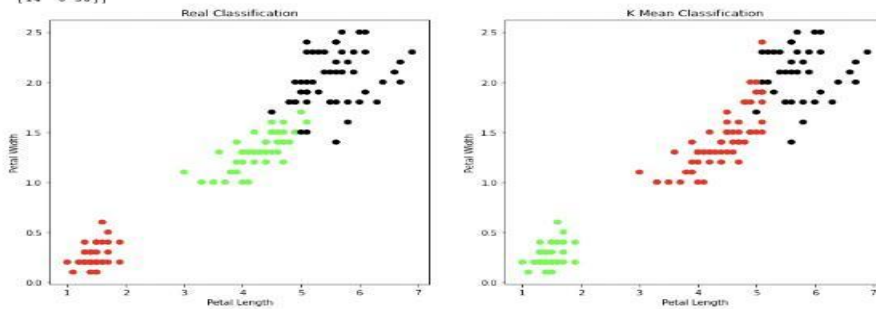
```
In [3]: model = KMeans(n_clusters=3)
model.fit(X)

plt.figure(figsize=(14,7))
colormap = np.array(['red', 'lime', 'black'])

# Plot the Original Classifications
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')

# 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 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 matrix of K-Mean: ', sm.confusion_matrix(y, model.labels_))
```

```
The accuracy score of K-Mean: 0.24
The Confusion matrix of K-Mean: [[ 0 50  0]
 [48  2]
 [14  36]]
```



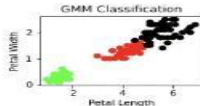
```
In [4]: from sklearn import preprocessing
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
```

```
In [5]: from sklearn.mixture import GaussianMixture
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)

y_gmm = gmm.predict(xs)
#y_cluster_gmm

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')
```

```
Out[5]: Text(0, 0.5, 'Petal Width')
```



```
In [6]: print('The accuracy score of EM: ', sm.accuracy_score(y, y_gmm))
print('The Confusion matrix of EM: ', sm.confusion_matrix(y, y_gmm))
```

```
The accuracy score of EM: 0.3333333333333333
The Confusion matrix of EM: [[ 0 50  0]
 [45  5]
 [ 0  50]]
```

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

```
from sklearn import datasets
```

```
print('eral ?<r.g:h'. ' :<pn1 wiz:h', 'gctnl lcr.gtu', 'retzi  
width') print(x)  
print('~ia*s: o lri: 6s:o< z 1 iris ver:i-odour, z ml Virjir.icz  
print(y)
```

```
>>cpal-long:h repel-wrath pct f-leng:b ps:*1-width
```

```
[4.9 3. 1.4 0.2]
```

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[4.6 3.1 1.5 0.2]
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[5. 3.6 1.4 0.2]
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[5. 3.4 1.5 0.2]
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[5.1 3.5 1.4 0.3]
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[5.1 3.7 1.5 0.4]
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[4.6 3.6 1. 0.2]
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[5.1 3.3 1.7 0.5]
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[5.2 3.4 1.4 0.2]
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[5.2 4.1 1.5 0.1]
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```
[4.9 3.1 1.5 0.2]
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[5. 3.2 1.2 0.2]
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[4.4 3. 1.3 0.2]
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[5. 3.5 1.3 0.3]
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[4.4 3.2 1.3 0.2]
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[5.1 3.8 1.9 0.4]
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[4.6 3.2 1.4 0.2]
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[6.5 2.8 4.6 1.5]
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[4.9 2.4 3.3 1. ]
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[6.2 2.2 4.5 1.5]
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[6.6 3. 4.4 1.4]
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[6. 2.9 4.5 1.5]
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[5.4 3. 4.5 1.5]
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[5.6 3. 4.1 1.3]
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[5.6 3. 4.1 1.3]
[5.5 2.5 4. 1.3]
[5.5 2.6 4.4 1.2]

[5.8 2.6 4. 1.2]
[5. 2.3 3.3 1.]
[5.6 2.7 4.2 1.3]
[5.7 3. 4.2 1.2]
[5.7 2.9 4.2 1.3]
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[5.7 2.5 4.1 1.3]
[5.7 2.4 4.1 1.3]
[6. 3 3.6 2.8]
[5.4 2.7 3.1 1.9]
[7. 1 3. 5.9 2.1]
[6. 3 2.9 3.6 1.8]
[6. 5 3. 5.5 2.2]

[3.9 2.5 4.5 1.7]
[7. 3 2.9 6.2 1.8]
[6. 7 2.5 3.1 1.8]
[7. 2.3 6.6 1.2.8]
[6.8 3.2 3.1 1.2.8]
[6.4 2.7 3.2 1.9]
[6.8 3. 5.5 2.1]
[5.1 2.5 3. 3.]
[5.B 2.B 3.1 3.4]
[6. a 3.2 3.2 2.3]
[6.5 3. 5.5 1.8]
[7.1 3.4 6.7 2.2]
[7.1 2.6 6.9 3.3]
[6. 2.2 3. 1.8]
[6.9 3.2 3.7 3.3]
[5.6 2.4 3.9 3.]
[7.1 2.4 6.7 2.]
[6.3 2.7 3.9 1.8]
[6.7 3.3 5.7 2.1]
[7.1 3.2 6. 1.8]
[6.1 2.B 4.4 1.8]

[6.4 2.8 5.6 2.1]
[7.2 3. 5.8 1.6]
[7. 2.4 6.1 1.9]

[6. 2.4 3.5 2.2]
[6.3 2.4 3.1 1.8]
[6 1 2.6 3.6 1.4]
[7.7 3. 6.1 2.3]
[6.3 3.4 5.6 2.4]
[6.4 3.1 5.5 1.8]

[6.9 3.1 5.4 2.1]

[0. 9 3.1 3.1 2.3]
[5.B 2.7 3.1 1.9]
[6 # 3.2 3.9 2.3]
[6.7 3.3 5.7 2.5]
[0. 7 3. 3.2 2.3]
[0. 3 2.5 3. 1.9]
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[5.93.3.11.8]
class: 0-Iris-Setosa, 1-
[0 0 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0 0
1 1 1 1 1 1 1 1 1 1
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2 2]

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Iris-Versicolour, 2- Iris-Virginica
0 0 0 0 0 0 0 0 0 0 0 0 0 0
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1 1 1 1 1 1 1 1 1 1 1 1 2 2 2
2 2 2 2 2 2 2 2 2 2 2 2 2 2

x_train, *_teat, y_train, y_tozt = train_test_split(n.y,test_zizo*0.1]

y = red-oLazsiKier.prediot<n_tozt*

t[16 a d j

		1. aa 1. aa 0. 8a		1. ba			
	a	1. aa		b. 9a	14		16
	1	0.82		d.e9	15		
	2	1. aa					45
accuracy		6. 92		0.9J			
wrSg%tod avg	0.95			4t		0.93	

9. Implement the Linear Regression algorithm in order to fit data points. Select the appropriate data set for your experiment and draw graphs.

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd

In [3]: dataset = pd.read_csv('salary_dataset.csv')
X = dataset.iloc[:, 1:-1].values
y = dataset.iloc[:, 1].values

In [4]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=1/3, random_state=0)

In [5]: # Fitting Simple Linear Regression to the Training set
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
```

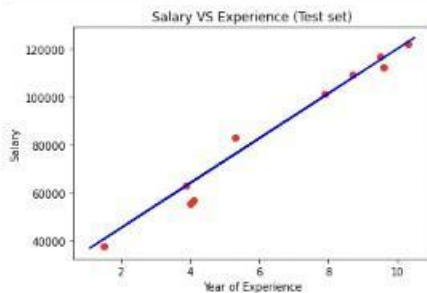
Out[5]: LinearRegression()

```
In [6]: # Predicting the Test set results
y_pred = regressor.predict(X_test)
```

```
In [7]: # Visualizing the Training set results
viz_train = plt
viz_train.scatter(X_train, y_train, color='red')
viz_train.plot(X_train, regressor.predict(X_train), color='blue')
viz_train.title('Salary VS Experience (Training set)')
viz_train.xlabel('Year of Experience')
viz_train.ylabel('Salary')
viz_train.show()
```



```
In [8]: # Visualizing the Test set results
viz_test = plt
viz_test.scatter(X_test, y_test, color='red')
viz_test.plot(X_train, regressor.predict(X_train), color='blue')
viz_test.title('Salary VS Experience (Test set)')
viz_test.xlabel('Year of Experience')
viz_test.ylabel('Salary')
viz_test.show()
```



10. Implement the non-parametric Locally Weighted Regression algorithm in order to fit data points. Select the appropriate data set for your experiment and draw graphs.

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
```

```
In [3]: dataset = pd.read_csv('salary_dataset.csv')
X = dataset.iloc[:, 1:-1].values
y = dataset.iloc[:, 1].values
```

```
In [4]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=1/3, random_state=0)
```

```
In [5]: # Fitting Simple Linear Regression to the Training set
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
```

```
Out[5]: LinearRegression()
```

```
In [6]: # Predicting the Test set results
y_pred = regressor.predict(X_test)
```

```
In [7]: # Visualizing the Training set results
viz_train = plt
viz_train.scatter(X_train, y_train, color='red')
viz_train.plot(X_train, regressor.predict(X_train), color='blue')
viz_train.title('Salary VS Experience (Training set)')
viz_train.xlabel('Year of Experience')
viz_train.ylabel('Salary')
viz_train.show()
```



```
In [8]: # Visualizing the Test set results
viz_test = plt
viz_test.scatter(X_test, y_test, color='red')
viz_test.plot(X_train, regressor.predict(X_train), color='blue')
viz_test.title('Salary VS Experience (Test set)')
viz_test.xlabel('Year of Experience')
viz_test.ylabel('Salary')
viz_test.show()
```

