

AMITY UNIVERSITY NOIDA AMITY SCHOOL OF ENGINEERING AND TECHNOLOGY

ARTIFICIAL INTELLIGENCE AND MACHINE LEARNING PRACTICAL FILE

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PRACTICAL - 1

AIM:

- 1) To read the Iris Dataset.
- 2) To find any missing values.
- 3) To find the average value of each column.
- 4) To find the maximum value in each column.
- 5) To find the minimum value in each column.

LIBRARIES IMPORTED: Pandas

INPUT:

1) Import pandas as pd

>>>

>>> #Correct file path to the CSV file

>>> file_path = r'C:\Users\Peter|Downloads\Iris.csv'

>>>

>>> #Read the entire CSV file into a DataFrame

>>> iris_df = pd.read_csv (file_path)

>>>

>>> #Display the entire dataset (if its not too large)

>>> print(iris_df)

		_ /				
	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica
[150	rows	x 6 columns]				

```
2) import pandas as pd
>>>
>>> #File path to the CSV file
>>> file_path = r'C:\Users\Peter\Downloads\Iris.csv'
>>>
>>> #Read the CSV file into a DataFrame
>>> iris_df = pd.read_csv(file path)
>>>
>>> #Check for missing values in each column
>>> missing_values = iris_df.isna().sum()
>>>
>>> #Display missing values count for each column
>>> print(missing_values)
```

```
Id 0
SepalLengthCm 0
SepalWidthCm 0
PetalLengthCm 0
PetalWidthCm 0
Species 0
dtype: int64
>>>>
```

```
3) # Select only numeric columns from the dataset
>>> numeric_columns = iris_df.select_dtypes(include=['float64', 'int64'])
>>>
>>> # Calculate the mean (average) of each numeric column
>>> average_values = numeric_columns.mean()
>>>
```

>>> # Display the average for each numeric column

>>> print(average_values)

OUTPUT:

```
Id 75.500000

SepalLengthCm 5.843333

SepalWidthCm 3.054000

PetalLengthCm 3.758667

PetalWidthCm 1.198667

dtype: float64

>>>>
```

4) # Select only numeric columns from the dataset

>>> numeric_columns = iris_df.select_dtypes(include=['float64', 'int64'])

>>>

>>> # Find the maximum value of each numeric column

>>> max_values = numeric_columns.max()

>>>

>>> # Display the maximum value for each numeric column

>>> print(max_values)

OUTPUT:

```
Id 150.0
SepalLengthCm 7.9
SepalWidthCm 4.4
PetalLengthCm 6.9
PetalWidthCm 2.5
dtype: float64
```

5) # Select only numeric columns from the dataset

```
>>> numeric_columns = iris_df.select_dtypes(include=['float64', 'int64'])
>>>
>>> # Find the minimum value of each numeric column
>>> min_values = numeric_columns.min()
>>>
>>> # Display the minimum value for each numeric column
>>> print(min_values)
```

```
Id 1.0
SepalLengthCm 4.3
SepalWidthCm 2.0
PetalLengthCm 1.0
PetalWidthCm 0.1
dtype: float64
>>>>
```

PRACTICAL - 2

AIM:

Implementing k-nearest neighbour algorithm in python.

INPUT:

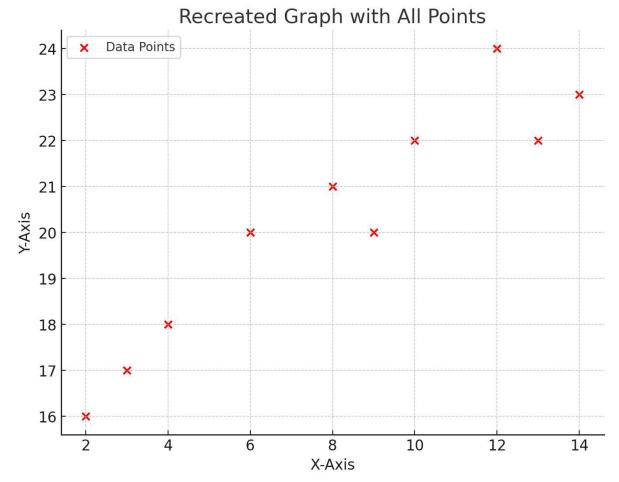
Import matplotlib.pyplot as plt

From sklearn.neighbors import KNeighborsClassifier as knc

```
#dataset
x = [4, 5, 10, 4, 3, 11, 14, 8, 10, 12]
y = [21, 19, 24, 17, 16, 25, 24, 22, 21, 21]
classes = [0, 0, 1, 0, 0, 1, 1, 0, 1, 1]
# plt.plot(x, y, '*')
# plt.show()
printf('Plotting the given DataSet')
plt.scatter(x = x, y = y, c = classes, marker = '*')
plt.show()
data = list(zip(x, y))
printf('Making the Array')
print(data)
k = knc(n_neighbors = 3)
k.fit(data, classes)
new_x = 8
new_y = 21
np = [(new_x, new_y)]
class
prediction = k.predict(np)
printf('Predicting the new point class')
print(prediction)
```

print('Plotting full data')
plt.scanner(x = x + [new_x], y = y + [new_y], c = classes
[prediction[0]], marker = '*')
plt.show()
OUTPUT:

0011 01.



EXPERIMENT: 3

AIM: Implement K mean clustering algorithm.

LIBRARIES IMPORTED: Numpy, Sklearn

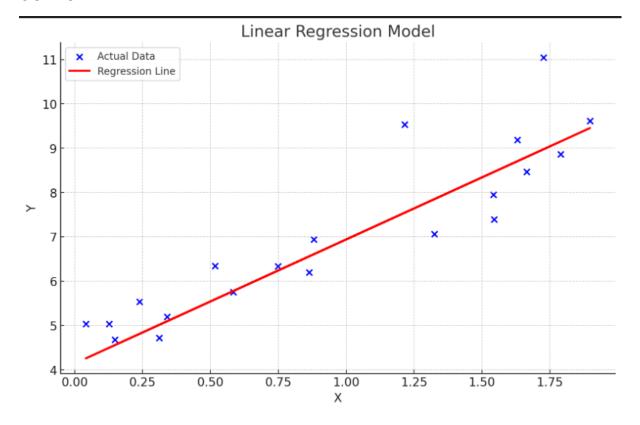
INPUT:

```
import numpy as np
from sklearn.cluster import KMeans
#data points using numpy
data = np.array([
    [2, 1], [3, 4], [2, 4], [8, 8], [4, 1], [4, 3], [5, 6],
    [7, 8], [6, 5], [6,2], [8,1], [6,1], [5,2], [4,2], [2,5]
    1)
# Number of clusters (k)
k = 4
# Apply K-Means clustering
kmeans = KMeans(n clusters=k, random state=42, n init=10)
kmeans.fit(data)
centroids = kmeans.cluster centers
labels = kmeans.labels_
print("Final Centroids:") #printing centroids
print(centroids)
print("\nClusters Formed:")
for i in range(k):
    # Grouping data points by clusters
    print(f"Cluster {i}: {data[labels == i]}")
```

OUTPUT SCREEN:

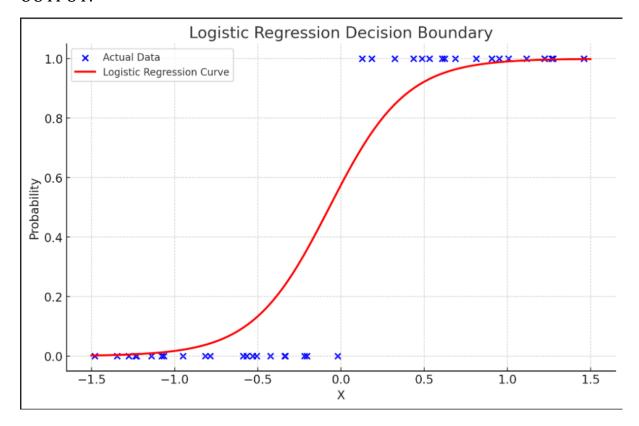
```
Final Centroids:
[[3.8
              1.8
              6.75
 [6.5]
 [2.33333333 4.33333333]
 [6.66666667 1.333333333]]
Clusters Formed:
Cluster 0: [[2 1]
 [4\ 1]
 [4 3]
 [5 2]
 [4 2]]
Cluster 1: [[8 8]
 [5 6]
 [7 8]
 [6 5]]
Cluster 2: [[3 4]
 [2 4]
 [2 5]]
Cluster 3: [[6 2]
 [8 1]
 [6 1]]
>>>
```

```
AIM: Linear Regression
LIBRARIES IMPORTED: Numpy, Sklearn
INPUT:
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model selection import train test split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean squared errornp.random.seed(42)
x = 2 * np.random.rand(100, 1)
y = 4 + 3 * x + np.random.randn(100, 1)
x_{train}, x_{test}, y_{train}, y_{test} = train_{test_split}(x, y, test_{size} = 0.2, y_{train})
random_state=42)
model = LinearRegression()
model.fit(x_train, y_train)
y_pred = model.predict(x_test)
mse = mean_squared_error(y_test, y_pred)
print(f"Mean Squared Error: {mse}")
print(f"Intercept: {model.intercept_}, Coefficient: {model.coef_}")
plt.scatter(x_test, y_test, color='blue', label='Actual Data')
plt.plot(x_test, y_pred, color='red', linewidth=2, label='Regression Line')
plt.xlabel("X")
plt.ylabel("Y")
plt.legend()
plt.show()
```



```
AIM: Logistic Regression
LIBRARIES IMPORTED: Numpy, Sklearn
INPUT:
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model selection import train test split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy score, confusion matrix,
classification_report
np.random.seed(42)
x = 3 * np.random.rand(200, 1) - 1.5 # Random values between -1.5 and
1.5
y = (x > 0).astype(int).ravel() # Label: 1 if x > 0, else 0
x_{train}, x_{test}, y_{train}, y_{test} = train_{test_split}(x, y, test_{size}=0.2,
random state=42)
model = LogisticRegression()
model.fit(x_train, y_train)y_pred = model.predict(x_test)accuracy =
accuracy_score(y_test, y_pred)
conf_matrix = confusion_matrix(y_test, y_pred)
report = classification_report(y_test, y_pred)
print(f"Accuracy: {accuracy:.2f}")
print("Confusion Matrix:")
print(conf_matrix)
print("Classification Report:")
print(report)
x_{values} = np.linspace(-1.5, 1.5, 100).reshape(-1, 1)
```

```
y_prob = model.predict_proba(x_values)[:, 1]
plt.scatter(x_test, y_test, color='blue', label='Actual Data')
plt.plot(x_values, y_prob, color='red', linewidth=2, label='Logistic Regression Curve')
plt.xlabel("X")
plt.ylabel("Probability")
plt.legend()
plt.title("Logistic Regression Decision Boundary")
plt.show()
```

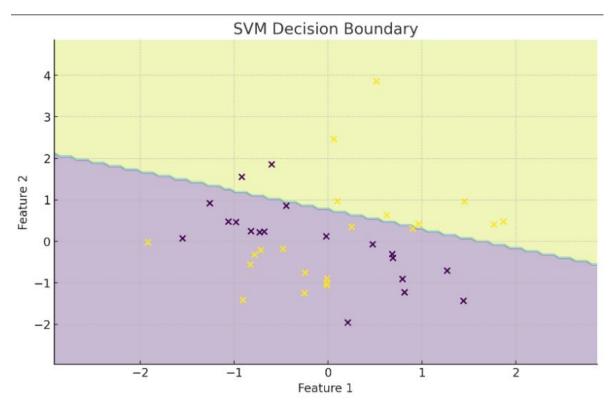


	precision	recall	f1-score	support
0	1.00	0.95	0.98	21
1	0.95	1.00	0.97	19
accuracy			0.97	40
macro avg	0.97	0.98	0.97	40

```
AIM: Support Vector Machine
LIBRARIES IMPORT: Numpy, Sklearn
INPUT:
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model selection import train test split
from sklearn.svm import SVC
from sklearn.metrics import accuracy score, confusion matrix,
classification_report
np.random.seed(42)
x = np.random.randn(200, 2) # 2D data for visualization
y = (x[:, 0] * x[:, 1] > 0).astype(int) # Label: 1 if x1*x2 > 0, else 0x_{train},
x_{test}, y_{test} = train_test_split(x, y, test_size=0.2,
random_state=42)model = SVC(kernel='linear')
model.fit(x_train, y_train)
y_pred = model.predict(x_test)
accuracy = accuracy_score(y_test, y_pred)
conf_matrix = confusion_matrix(y_test, y_pred)
report = classification_report(y_test, y_pred)
print(f"Accuracy: {accuracy:.2f}")
print("Confusion Matrix:")
print(conf_matrix)
print("Classification Report:")
print(report)
def plot_decision_boundary(model, 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
xx, yy = np.meshgrid(np.linspace(x_min, x_max, 100), np.linspace(y_min, y_max, 100))
Z = model.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.contourf(xx, yy, Z, alpha=0.3)
plt.scatter(X[:, 0], X[:, 1], c=y, edgecolors='k')
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.title("SVM Decision Boundary")
plt.show()
```

plot_decision_boundary(model, x_test, y_test)



	precision	recall	f1-score	support
0	0. 57	0.89	0.69	19
1	0.80	0.38	0.52	21
accuracy			0.62	40
macro avg	0.68	0.64	0.61	40

AIM: Wine Quality Prediction

LIBRARIES: Pandas, Numpy

INPUT:

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

import seaborn as sb

from sklearn.model_selection import train_test_split

from sklearn.preprocessing import MinMaxScaler

from sklearn import metrics

from sklearn.svm import SVC

from xgboost import XGBClassifier

from sklearn.linear_model import LogisticRegression

import warnings

warnings.filterwarnings('ignore')

0 1 2 3 4	type fix white white white white white	ed acidity 7.0 6.3 8.1 7.2 7.2	volatile	acidity 0.27 0.30 0.28 0.23	0.34 0.46 0.32		sugar 20.7 1.6 6.9 8.5 8.5	\
0 1 2 3 4	chlorides 0.045 0.049 0.050 0.058 0.058	free sulfu	45.0 45.0 14.0 30.0 47.0 47.0	total	sulfur dioxid 170. 132. 97. 186. 186.	0 1.0010 0 0.9940 0 0.9951 0 0.9956	3.00 3.30 3.26 3.19	\
0 1 2 3 4	sulphates 0.45 0.49 0.44 0.40	alcohol 0 8.8 9.5 10.1 9.9	uality 6 6 6 6					