

-----First Program-----

```
import csv
a = []
with open('finds.csv') as csfile:
    reader = csv.reader(csfile)
    for row in reader:
        a.append(row)
        print(row)
num_attributes = len(a[0]) - 1
print(["?" * num_attributes])
print(["0" * num_attributes])
hypothesis = a[0][:-1]
for i in range(len(a)):
    if a[i][num_attributes] == "Yes":
        for j in range(num_attributes):
            if a[i][j] != hypothesis[j]:
                hypothesis[j] = '?'
    print(i + 1, hypothesis)
print(hypothesis)
```

-----finds.csv

```
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
```

-----Second Program-----

Program 2

```
import numpy as np
import pandas as pd
# Loading Data from a CSV File
data = pd.DataFrame(data=pd.read_csv('Training_examples.csv'))
# Separating concept features from Target
concepts = np.array(data.iloc[:, 0:-1])
target = np.array(data.iloc[:, -1])
```

```

def learn(concepts, target):
    specific_h = concepts[0].copy()
    general_h = [['?' for i in range(len(specific_h))] for i in range(len(specific_h))]
    for i, h in enumerate(concepts):
        if target[i] == "Yes":
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    specific_h[x] = '?'
                    general_h[x][x] = '?'
        if target[i] == "No":
            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] = '?'
    indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
    for i in indices:
        general_h.remove(['?', '?', '?', '?', '?', '?'])
    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")

```

Training_examples.csv

Sky,Air,Humidity,Wind,Water,Forecast,EnjoySport

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

-----Program 3-----

```
import pandas as pd
```

```
import numpy as np
```

```
# Import the dataset and define the feature as well as the target datasets/columns
```

```
dataset = pd.read_csv('playtennis.csv', names=['outlook', 'temperature', 'humidity', 'wind', 'class'])
```

```
attributes = ('Outlook', 'Temperature', 'Humidity', 'Wind', 'PlayTennis')
```

```

def entropy(target_col):
    elements, counts = np.unique(target_col, return_counts=True)

    entropy = np.sum([(-counts[i] / np.sum(counts)) * np.log2(counts[i] / np.sum(counts)) for i in
range(len(elements))])

    return entropy

def InfoGain(data, split_attribute_name, target_name="class"):
    total_entropy = entropy(data[target_name])

    vals, counts = np.unique(data[split_attribute_name], return_counts=True)

    Weighted_Entropy = np.sum([(counts[i] / np.sum(counts)) *
entropy(data.where(data[split_attribute_name] == vals[i]).dropna()[target_name]) for i in
range(len(vals))])

    Information_Gain = total_entropy - Weighted_Entropy

    return Information_Gain

def ID3(data, originaldata, features, target_attribute_name="class", parent_node_class=None):
    if len(np.unique(data[target_attribute_name])) <= 1:
        return np.unique(data[target_attribute_name])[0]

    elif len(data) == 0:
        return

    np.unique(originaldata[target_attribute_name])[np.argmax(np.unique(originaldata[target_attribute_
name], return_counts=True)[1])]

    elif len(features) == 0:
        return parent_node_class

    else:
        parent_node_class =
np.unique(data[target_attribute_name])[np.argmax(np.unique(data[target_attribute_name],
return_counts=True)[1])]

        item_values = [InfoGain(data, feature, target_attribute_name) for feature in features]

        best_feature_index = np.argmax(item_values)
        best_feature = features[best_feature_index]
        tree = {best_feature: {}}

        features = [i for i in features if i != best_feature]

        for value in np.unique(data[best_feature]):
            value = value

            sub_data = data.where(data[best_feature] == value).dropna()

            subtree = ID3(sub_data, dataset, features, target_attribute_name, parent_node_class)

            tree[best_feature][value] = subtree

        return tree

```

```

def predict(query, tree, default=1):
    for key in list(query.keys()):
        if key in list(tree.keys()):
            try:
                result = tree[key][query[key]]
            except:
                return default
            result = tree[key][query[key]]
            if isinstance(result, dict):
                return predict(query, result)
            else:
                return result

def train_test_split(dataset):
    training_data = dataset.iloc[:14].reset_index(drop=True)
    return training_data

def test(data, tree):
    queries = data.iloc[:, :-1].to_dict(orient="records")
    predicted = pd.DataFrame(columns=["predicted"])
    for i in range(len(data)):
        predicted.loc[i, "predicted"] = predict(queries[i], tree, 1.0)

    print('The prediction accuracy is: ', (np.sum(predicted["predicted"] == data["class"]) / len(data)) *
    100, '%')

XX = train_test_split(dataset)
training_data = XX
tree = ID3(training_data, training_data, training_data.columns[:-1])
print('Display Tree:', tree)
print('len=', len(training_data))
test(training_data, tree)

```

.csv

0,0,0,0,0

1,2,1,1,1

1,0,1,0,1

0,0,0,1,0

0,1,0,0,0

2,1,0,1,0

1,0,0,0,1

0,2,1,0,1

0,1,1,1,1

2,1,0,0,1

2,1,1,0,1

1,1,1,1,1

2,2,1,0,1

0,1,1,1,1

2,2,1,1,0

1,1,0,1,1

-----# Program 4-----

```
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 = 7000
learning_rate = 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))
wo = np.random.uniform(size=(hiddenlayer_neurons, output_neurons))
bo = np.random.uniform(size=(1, output_neurons))

for i in range(epoch):
    net_h = np.dot(X, wh) + bh
    sigma_h = sigmoid(net_h)
    net_o = np.dot(sigma_h, wo) + bo
    output = sigmoid(net_o)
    deltaK = (y - output) * derivatives_sigmoid(output)
    deltaH = deltaK.dot(wo.T) * derivatives_sigmoid(sigma_h)
    wo = wo + sigma_h.T.dot(deltaK) * learning_rate
    wh = wh + X.T.dot(deltaH) * learning_rate

print("Input: \n" + str(X))
print("Actual Output: \n" + str(y))
print("Predicted Output: \n", output)
```

-----Program 5-----

```
import pandas as pd

from sklearn.model_selection import train_test_split

from sklearn.naive_bayes import MultinomialNB

from sklearn import metrics

data = pd.read_csv('dataset5.csv')

X = data.iloc[:, :-1]

y = data.iloc[:, -1]

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

model = MultinomialNB()

model.fit(X_train, y_train)

y_pred = model.predict(X_test)

accuracy = metrics.accuracy_score(y_test, y_pred)

precision = metrics.precision_score(y_test, y_pred, average='weighted', zero_division=1)

recall = metrics.recall_score(y_test, y_pred, average='weighted')

print(f"Accuracy: {accuracy:.2f}")

print(f"Precision: {precision:.2f}")

print(f"Recall: {recall:.2f}")
```

-----dataset5.csv-----

feature1,feature2,feature3,feature4,target

```
5.1,3.5,1.4,0.2,class1
4.9,3.0,1.4,0.2,class1
4.7,3.2,1.3,0.2,class1
4.6,3.1,1.5,0.2,class1
5.0,3.6,1.4,0.2,class1
5.4,3.9,1.7,0.4,class2
4.6,3.4,1.4,0.3,class2
5.0,3.4,1.5,0.2,class2
4.4,2.9,1.4,0.2,class2
4.9,3.1,1.5,0.1,class2
5.4,3.7,1.5,0.2,class3
4.8,3.4,1.6,0.2,class3
4.8,3.0,1.4,0.1,class3
4.3,3.0,1.1,0.1,class3
```

5.8,4.0,1.2,0.2,class3

-----#Program 6-----

```
import pandas as pd

from pgmpy.estimators import MaximumLikelihoodEstimator

from pgmpy.models import BayesianModel

from pgmpy.inference import VariableElimination

data = pd.read_csv("ds4.csv")

heart_disease = pd.DataFrame(data)

print(heart_disease)

model = BayesianModel([

    ('age', 'Lifestyle'),

    ('Gender', 'Lifestyle'),

    ('Family', 'heartdisease'),

    ('diet', 'cholesterol'),

    ('Lifestyle', 'diet'),

    ('cholesterol', 'heartdisease'),

    ('diet', 'cholesterol')])

model.fit(heart_disease, estimator=MaximumLikelihoodEstimator)

HeartDisease_infer = VariableElimination(model)

print('For Age enter SuperSeniorCitizen:0, SeniorCitizen:1, MiddleAged:2, Youth:3, Teen:4')

print('For Gender enter Male:0, Female:1')

print('For Family History enter Yes:1, No:0')

print('For Diet enter High:0, Medium:1')

print('for LifeStyle enter Athlete:0, Active:1, Moderate:2, Sedentary:3')

print('for Cholesterol enter High:0, BorderLine:1, Normal:2')

q = HeartDisease_infer.query(variables=['heartdisease'], evidence={

    'age': int(input('Enter Age: ')),

    'Gender': int(input('Enter Gender: ')),

    'Family': int(input('Enter Family History: ')),

    'diet': int(input('Enter Diet: ')),

    'Lifestyle': int(input('Enter Lifestyle: ')),

    'cholesterol': int(input('Enter Cholesterol: '))})

print(q)

---ds4.csv

age,Gender,Family,diet,Lifestyle,cholesterol,heartdisease
```

0,0,1,1,3,0,1
0,1,1,1,3,0,1
1,0,0,0,2,1,1
4,0,1,1,3,2,0
3,1,1,0,0,2,0
2,0,1,1,1,0,1
4,0,1,0,2,0,1
0,0,1,1,3,0,1
3,1,1,0,0,2,0
1,1,0,0,0,2,1
4,1,0,1,2,0,1
4,0,1,1,3,2,0
2,1,0,0,0,0,0
2,0,1,1,1,0,1
3,1,1,0,0,1,0
0,0,1,0,0,2,1
1,1,0,1,2,1,1
3,1,1,1,0,1,0
4,0,1,1,3,2,0

-----#program 7

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

# Step 1: Load data from a CSV file
def load_data(csv_file):
    data = pd.read_csv(csv_file)
    return data

# Step 2: Apply K-Means clustering
def apply_kmeans(data, n_clusters):
    # K-Means algorithm
    kmeans = KMeans(n_clusters=n_clusters)
    kmeans.fit(data)

    # Step 3: Get the cluster labels
    labels = kmeans.labels_
```



```

return labels, kmeans.cluster_centers_

# Step 4: Visualize the clusters

def visualize_clusters(data, labels, cluster_centers):

    plt.scatter(data.iloc[:, 0], data.iloc[:, 1], c=labels, cmap='rainbow')

    # plt.scatter(cluster_centers[:, 0], cluster_centers[:, 1], s=300, c='black', marker='X') # Cluster
centers

    plt.title("K-Means Clustering")

    plt.xlabel("Feature 1")

    plt.ylabel("Feature 2")

    plt.show()

if __name__ == "__main__":

    # Load data from CSV (Assume the CSV file has two numeric columns)

    data = load_data('your_data.csv')

    # Specify the number of clusters

    n_clusters = 3

    # Apply K-Means algorithm

    labels, cluster_centers = apply_kmeans(data, n_clusters)

    # Visualize the clusters

    visualize_clusters(data, labels, cluster_centers)

```

-----your_data.csv

Feature1,Feature2

5.1,3.5	4.7,2.8	7.9,7.4	8.1,7.8
4.9,3.0	6.3,3.6	8.2,7.8	7.6,7.1
6.2,3.4	4.9,2.7	8.3,7.6	8.4,7.3
5.9,3.0	5.6,3.4	8.5,7.9	8.2,7.9
5.0,3.4	6.4,3.8	8.0,7.2	7.5,7.4
5.2,3.6	5.5,3.3	7.7,7.4	8.3,7.6
5.3,3.2	5.1,2.9	8.1,7.5	8.5,7.8
4.8,2.9	4.6,2.8	8.4,7.7	8.7,7.5
6.1,3.3	5.0,3.1	8.2,7.6	2.0,1.5
5.5,3.1	5.3,3.5	7.9,7.3	1.9,1.4
5.7,3.8	5.6,3.9	7.8,7.2	2.1,1.6
6.0,3.2	8.0,7.5	8.3,7.5	1.8,1.2
5.8,3.5	7.8,7.3	8.6,7.9	2.2,1.7
5.4,3.7	8.1,7.6	8.0,7.7	

1. Import Data

```
from sklearn.datasets import load_iris
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier

# Load Iris dataset
iris = load_iris()

# Create a DataFrame from the iris dataset
iris_df = pd.DataFrame(data=iris.data, columns=iris.feature_names)
iris_df['species'] = iris.target

# Save the dataset to a CSV file
iris_df.to_csv('iris_data', index=False)

# Print some details
print("Feature Names:", iris.feature_names)
print("Iris Data:")
print(iris.data)
print("Target Names:", iris.target_names)
print("Target:", iris.target)
```

2. Split the data into Training and Test sets

```
X_train, X_test, y_train, y_test = train_test_split(iris.data, iris.target, test_size=0.25)
```

3. Build the KNN model

```
clf = KNeighborsClassifier()
clf.fit(X_train, y_train)
```

4. Calculate accuracy on test data

```
accuracy = clf.score(X_test, y_test)
print("Accuracy =", accuracy)
```

5. Calculate predictions with the test data

```
predictions = clf.predict(X_test)
print("Predicted Data:")
print(predictions)
```

Display the test labels

```
print("Test Data:")
print(y_test)
```

6. Identify misclassifications

```

misclassifications = predictions - y_test
print("Result of misclassifications:")
print(misclassifications)
# Calculate total misclassified samples
total_misclassified = sum(abs(misclassifications))
print('Total number of samples misclassified =', total_misclassified)
-----9
from math import ceil
import numpy as np
from scipy import linalg
def lowess(x, y, f, iterations):
    n = len(x)
    r = int(ceil(f * n))
    h = [np.sort(np.abs(x - x[i]))[r] for i in range(n)]
    w = np.clip(np.abs((x[:, None] - x[None, :]) / h), 0.0, 1.0)
    w = (1 - w ** 3) ** 3
    yest = np.zeros(n)
    delta = np.ones(n)
    for iteration in range(iterations):
        for i in range(n):
            weights = delta * w[:, i]
            b = np.array([np.sum(weights * y), np.sum(weights * y * x)])
            A = np.array([[np.sum(weights), np.sum(weights * x)], [np.sum(weights * x), np.sum(weights *
x * x)]])
            beta = linalg.solve(A, b)
            yest[i] = beta[0] + beta[1] * x[i]
        residuals = y - yest
        s = np.median(np.abs(residuals))
        delta = np.clip(residuals / (6.0 * s), -1, 1)
        delta = (1 - delta ** 2) ** 2
    return yest
import math
n = 100
x = np.linspace(0, 2 * math.pi, n)
y = np.sin(x) + 0.3 * np.random.randn(n)

```

```
f = 0.25
```

```
iterations = 3
```

```
yest = lowess(x, y, f, iterations)
```

```
import matplotlib.pyplot as plt
```

```
plt.plot(x, y, "r.")
```

```
plt.plot(x, yest, "b-")
```