

## **Parul** University

# FACULTY OF ENGINEERING AND TECHNOLOGY BACHELOR OF TECHNOLOGY

**MACHINE LEARNING** 

**LABORATORY** 

(303105354)

**6TH – SEMESTER** 

(ARTIFICIAL INTELLIGENCE)

**COMPUTER SCIENCE & ENGINEERING** 

**DEPARTMENT** 

LAB MANUAL



ERP NO: 2203031240160

#### **CERTIFICATE**

This Is To Certify That Mr./Ms. BHANUVARDHAN.MEDAPALLI

With Enrollment No: 2203031240160 Has Completed His/Her

Laboratory Experiments (303105354) From The Department Of

#### **COMPUTER SCIENCE (ARTIFICIAL INTELLGENCE)**

During The Academic Year 2024 - 2025



Date of Submission :		Staff Incharge:
	Head Of Department :	



SR.NO	EXPERIMENT	PAGI	E NO	DATE OF COMPLETION	SIGN	MARKS
		FROM	TO			
1.	Dealing with Data using Numpy, Pandas, Statistics libray					
2.	Data Analysis & Visualization on Diwali Sales Dataset.					
3.	Implement linear regression and logistic regression.					
4.	implement the naïve Bayesian classifier for a sample training data set stored as a (.CSV) file. Compute the accuracy of the classifier, considering a few test data sets.					
5.	Assuming a set of documents that need to be classified, use the naïve Bayesian Classifier model to perform this task.					
6.	Decision tree-based ID3 algorithm.					
7.	Write a program to implement the K-Nearest Neighbour algorithm to classify the iris data set.					
8.	Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for clustering using k-Means algorithm.					
9.	Write a program to construct a Bayesian network considering medical data. Use this model to demonstrate the diagnosis of heart patients using standard Heart Disease Data Set.					
10.	Compare the various supervised learning algorithm by using appropriate dataset. (Linear Regression, Support Vector Machine, Decision Tree).					



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11.	Compare the various Unsupervised learning algorithm by using the appropriate datasets. (K Means Clustering, K Mode, ).			
12.	Build an Artificial Neural Network by implementing the Backpropagation algorithm and test the same using appropriate data sets.			



### FACULTY OF ENGINEERING AND TECHNOLOGY MACHINE LEARNING (303105354)

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#### PRACTICAL - 1

❖ <u>AIM</u>: DEALING WITH DATA USING NUMPY, PANDAS, STATISTICS LIBRARY.

#### **PROCEDURE:**

#### **❖** NUMPY:

- CREATION OF AN ARRAY: Employ the APIs np.array() or np.zeros(), np.ones(), and np.arange() in arrays creation.
- OPERATIONS OF AN ARRAY: Carry out operations on the array elements, broadcasting and np functions such as np.sum(), np.mean()..numerical operations.
- INDEXING AND SLICING: Use indexing and slicing methods to construct and reconstruct the arrays.
- LINEAR ALGEBRA: Apply np.dot(), np.linalg.inv() and np.linalg.eig() to perform required matrix operations.
- RANDOM NUMBER GENERATION / DISTRIBUTION : Apply the method np.random or generate random numbers, obtain random samples and shuffle data.
- PERFORMANCE: Thus, for numerical computation, NumPy arrays tend to be more memory efficient and faster than using Python lists.

#### **PANDAS**:

- DATA STRUCTURES: the most primary data structure used are Series (1D) and DataFrame
   (2D)
- DATA IMPORT/EXPORT : For importing and exporting datasets, use pd.read\_csv(), for example, pd.read\_excel() and df.to\_csv() for preparing and presenting data.
- DATA MANIPULATION: Some operations are performed such as filtering, groupby, merging using merge, concatenating with concat.
- HANDLING MISSING DATA: To deal with the missing values df.isnull(), df.fillna() and df.dropna() are required.
- DATA TRANSFORMATION: Data Frames can be transformed by applying functions through apply, map, and applymap.
- TIME SERIES ANALYSIS: Use the functions exposed by the Pandas package for any kind of time and date data, including resampling and time zone.

#### **STATISTICS LIBRARY:**



- DESCRIPTIVE STATISTICS : Calculate measures of central tendency and dispersion to summarize data.
- PROBABILITY DISTRIBUTIONS: Understand and use functions for normal, binomial, and other distributions.
- HYPOTHESIS TESTING: Conduct tests like t-tests and chi-squared tests to make inferences about populations.
- CORRELATION AND REGRESSION: Use correlation coefficients and regression analysis to explore relationships between variables.
- DATA VISUALIZATION: While the Statistics library does not provide visualization, integrate with libraries like Matplotlib or Seaborn for graphical representation of statistical results.

#### **PROGRAM**:

#### **IMPORTING LIBRARIES:**

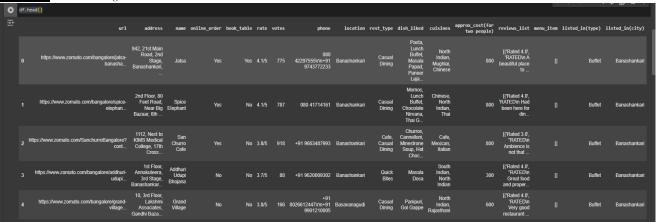
import pandas as pd
import numpy as np
import seaborn as sns
%matplotlib inline
import matplotlib.pyplot as plt
plt.rcParams['figure.figsize'] = (12, 5);
sns.set\_style('whitegrid')
import matplotlib.colors as mcolors
! pip install dexplot
import dexplot as dxp
import re
import string
import nltk
nltk.download('stopwords')
from nltk.corpus import stopwords

```
Requirement already satisfied: dexplot in /usr/local/lib/python3.10/dist-packages (0.1.2)
Requirement already satisfied: matplotlib>=3.1 in /usr/local/lib/python3.10/dist-packages (from dexplot) (3.8.0)
Requirement already satisfied: pandas>=0.24 in /usr/local/lib/python3.10/dist-packages (from dexplot) (2.2.2)
Requirement already satisfied: numpy>=1.15 in /usr/local/lib/python3.10/dist-packages (from dexplot) (1.26.4)
Requirement already satisfied: scipy>=1.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (1.3.1)
Requirement already satisfied: contourpy>=1.0.1 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (0.12.1)
Requirement already satisfied: fonttools>=0.20 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (0.12.1)
Requirement already satisfied: kiwisolver>=1.0.1 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (1.4.7)
Requirement already satisfied: packagingy=20.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (24.2)
Requirement already satisfied: pillow>=6.2.0 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (24.2)
Requirement already satisfied: python4detutil>=2.7 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (3.2.0)
Requirement already satisfied: python4detutil>=2.7 in /usr/local/lib/python3.10/dist-packages (from matplotlib>=3.1->dexplot) (28.2)
Requirement already satisfied: python4detutil>=2.7 in /usr/local/lib/python3.10/dist-packages (from pandas>=0.24->dexplot) (2024.2)
Requirement already satisfied: tzdata>=2022.7 in /usr/local/lib/python3.10/dist-packages (from pandas>=0.24->dexplot) (2024.2)
Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.10/dist-packages (from python-dateutil>=2.7->matplotlib>=3.1->dexplot) (1.17.0)
[nltk_data] Downloading package stopwords to /root/nltk_data...
[nltk_data] Package stopwords is already up-to-date!
```

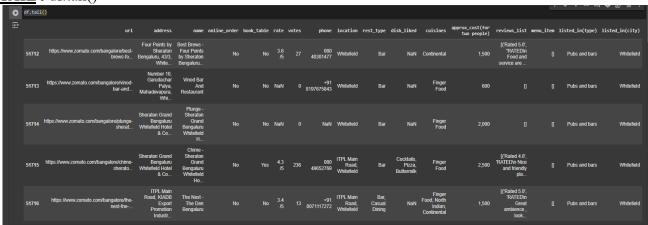
**READ CSV FILE**: df=pd.read csv('/content/zomato.csv')



**HEAD**: df.head()



#### TAIL: df.tail()



#### **INFO**: df.info()

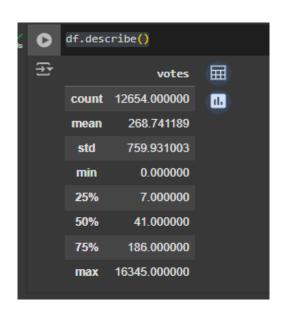
```
df.info()
0
    <class 'pandas.core.frame.DataFrame'>
    RangeIndex: 51717 entries, 0 to 51716
    Data columns (total 17 columns):
          Column
                                         Non-Null Count
                                                         Dtype
     0
          ur1
                                         51717 non-null
                                                          object
          address
                                         51717 non-null
                                                          object
                                         51717 non-null
     2
          name
                                                          object
          online_order
                                         51717 non-null
                                                          object
          book_table
     4
                                         51717 non-null
                                                          object
                                         43942 non-null
          rate
                                                          object
     6
7
          votes
                                         51717 non-null
                                                          int64
          phone
                                         50509 non-null
                                                          object
          location
                                         51696 non-null
          rest_type
                                               non-null
                                         51490
                                                          object
     10
         dish_liked
                                         23639
                                               non-null
                                                          object
         cuisines
                                         51672 non-null
                                                          object
          approx_cost(for two people)
                                         51371 non-null
                                                          object
     13
         reviews_list
                                         51717 non-null
                                                          object
         menu_item
     14
                                         51717 non-null
                                                          object
         listed_in(type)
     15
                                         51717 non-null
                                                          object
         listed_in(city)
                                                          object
     16
                                         51717 non-null
    dtypes: int64(1), object(16) memory usage: 6.7+ MB
```



**BOOLEAN VALUES**: df.isnull().sum()



DESCRIBE : df.describe()



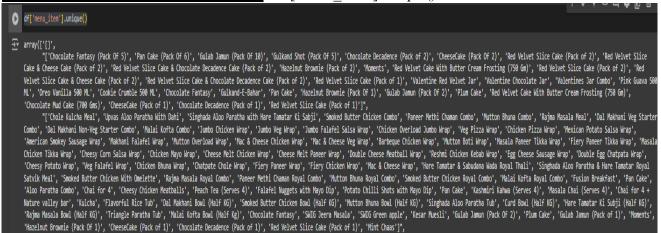
**SHAPE OF CSV FILE**: df.shape





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**DISTINICT VALUES OF COLUMNS :** df['menu item'].unique()



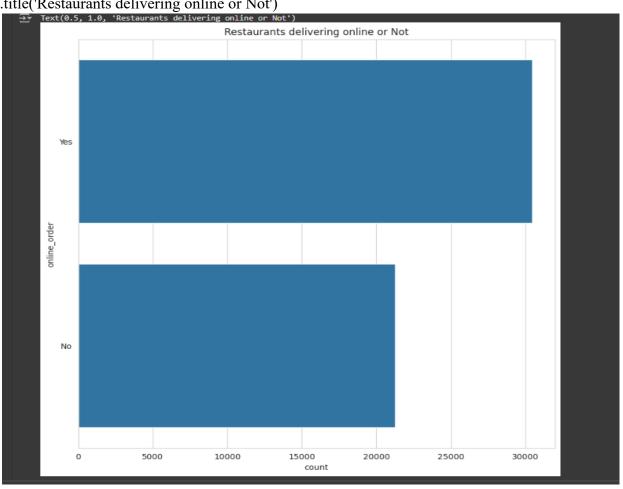
#### PLOT GRAPH OF RESTAURANTS DELIVERING ONLINE OR NOT:

# Plot of the Restaurant, whether they are accepting the online order or not sns.countplot(df['online order'])

fig = plt.gcf()

fig.set size inches(10,10)

plt.title('Restaurants delivering online or Not')



#### PLOT GRAPH OF FAMOUS RESTAURANTS IN THE CITY OF BENGALURU:

# Plotting for the Top Restuarant in the Bengaluru



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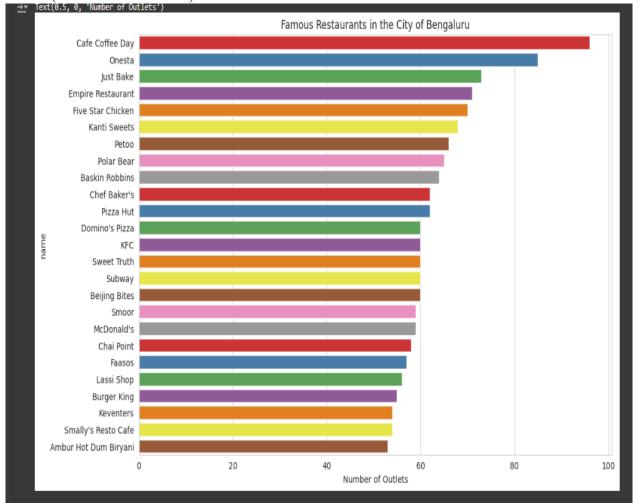
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plt.figure(figsize=(12,8))

val = df['name'].value\_counts()[:25] # Considering for the top 25 restaurants
sns.barplot(x=val, y=val.index, palette = "Set1")

plt.title("Famous Restaurants in the City of Bengaluru")

plt.xlabel("Number of Outlets")

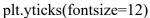


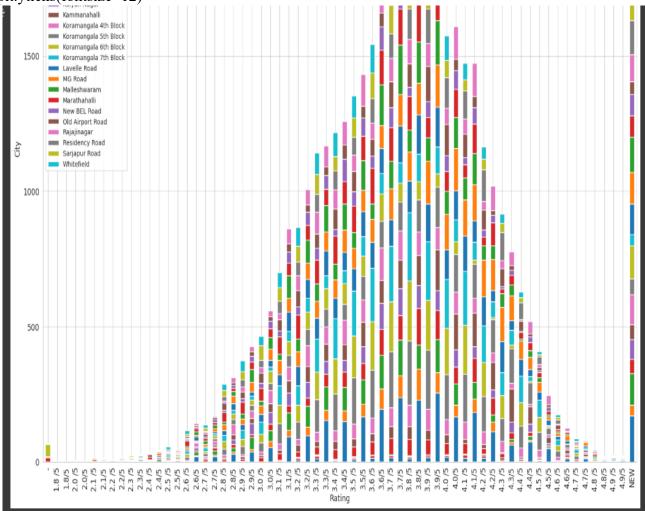


#### **PLOT GRAPH OF RATING:**

# Lets plot to see that whether there is any relation between the rating of the restaurant and the area in which they are located.

rel\_type = pd.crosstab(df['rate'], df['listed\_in(city)'])
rel\_type.plot(kind='bar', stacked=True, figsize=(20,16))
plt.title('City - Rating',fontsize=18)
plt.ylabel('City',fontsize=12)
plt.xlabel('Rating',fontsize=12)
plt.xticks(fontsize=12)







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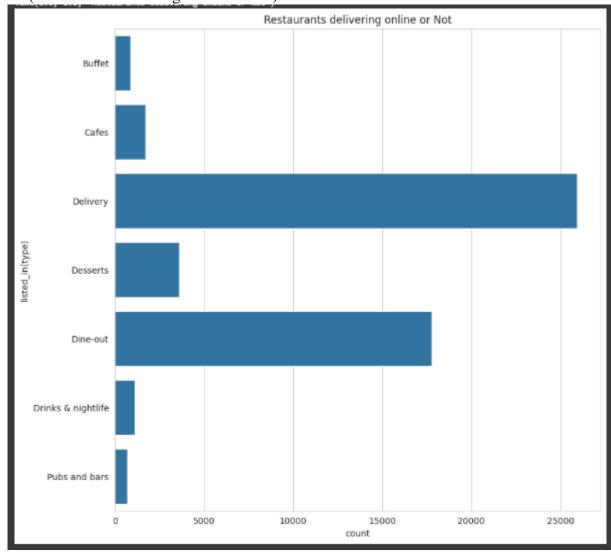
#### <u>PLOT GRAPH OF RESTAURANTS DELIVERING ONLINE OR NOT</u>:

sns.countplot(df['listed\_in(type)'])

fig = plt.gcf()

fig.set\_size\_inches(10,10)

plt.title('Restaurants delivering online or Not')





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#### PRACTICAL - 2

### ❖ <u>AIM</u>: DATA ANALYSIS AND DATA VISUALIZATION ON DIWALI SALES DATASET.

#### **PROCEDURE**:

Clearly outline the goals of your analysis. For instance, you might aim to understand sales trends during the Diwali festival, identify popular products, or analyze customer demographics.

#### 1. DATA COLLECTION:

Obtain the Diwali sales dataset, typically available in CSV format. This dataset should include relevant fields such as :

- Date: Date of sale
- Sales: Sales amount
- **Product**: Type of product sold
- User ID: Customer ID
- Cust name: Customer name
- Gender: Customer gender
- Age Group: Age category of the customer
- Marital Status: Customer's marital status
- State: State information
- Product\_Category: Category of the product sold

#### 2. DATA CLEANING:

Prepare the data for analysis by:

- Checking for missing values and handling them appropriately (e.g., imputation or removal).
- Correcting inconsistencies in data formats.
- Removing any duplicate entries.

#### 3. EXPLORATORY DATA ANALYSIS (EDA):

Conduct EDA to gain insights into the dataset:

- Use Python libraries like Pandas to load and manipulate the data.
- Generate summary statistics (mean, median, mode) to understand central tendencies.
- Identify patterns or correlations among variables using correlation matrices.

#### 4. DATA VISUALIZATION:

Create visual representations of your findings to make complex data understandable :

- Utilize libraries such as Matplotlib and Seaborn in Python for visualizations.
- Common visualizations include :
  - Bar Charts: To show sales by product category.
  - Line Graphs: To illustrate sales trends over time.
  - **Pie Charts**: To represent the proportion of sales by gender or age group.
  - **Heat Maps**: To visualize sales performance across different states or regions.



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#### 5. INTERPRETATION OF RESULTS:

Analyze the visualizations and summarize key insights:

- Identify which products had the highest sales.
- Understand customer demographics that contributed to sales.
- Recognize any seasonal trends or anomalies in sales data.

#### 6. REPORTING FINDINGS:

Prepare a report or presentation that includes:

- Visualizations that support your findings.
- A narrative that explains the insights derived from the analysis.
- Recommendations based on your analysis, such as marketing strategies or inventory management.

#### **PROGRAM**:

#### **IMPORTING LIBRARIES:**

import numpy as np import pandas as pd import matplotlib.pyplot as plt # visualizing data%matplotlib inline import seaborn as sns

#### **READING CSV FILE:**

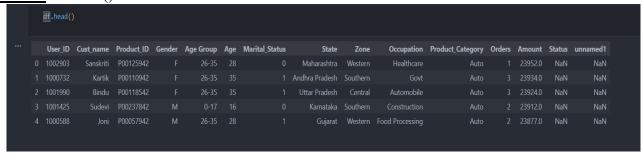
# import csv file

df = pd.read csv('Diwali Sales Data.csv', encoding= 'unicode escape')

#### **SHAPE OF CSV FILE**: df.shape



#### **HEAD**: df.head()





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#### **TO CHECK DATATYPES OF ALL COLUMN**: df.info()

D ~	C	df.info()		
	Rang	ss 'pandas.core.fr eIndex: 11251 entr columns (total 15 Column	ies, 0 to 11250	Dtype
	ø	User ID	11251 non-null	int64
		Cust name	11251 non-null	
	2	_	11251 non-null	_
		Gender	11251 non-null	•
			11251 non-null	_
	5	Age	11251 non-null	•
	6	•	11251 non-null	int64
	7	State	11251 non-null	object
	8	Zone	11251 non-null	object
	9	Occupation	11251 non-null	object
	10	Product_Category	11251 non-null	object
	11	Orders	11251 non-null	_
	12	Amount	11239 non-null	float64
	13	Status	0 non-null	float64
	14	unnamed1	0 non-null	float64
	dtyp	es: float64(3), in	t64(4), object(8	)
	memo	ry usage: 1.3+ MB		

#### **DROP UNRELATED/BLANK COLUMNS:**

df.drop(['Status', 'unnamed1'], axis=1, inplace=True)
#check for null values
pd.isnull(df).sum()

D ~	#check for null	values
	pd.isnull(df).s	um()
	User_ID	
	Cust_name	
	Product_ID	
	Gender	
	Age Group	
	Age	
	Marital_Status	
	State	
	Zone	
	Occupation	
	Product_Category	
	Orders	
	Amount	12
	dtype: int64	



#### **CHANGE DATA TYPE:**

df['Amount'] = df['Amount'].astype('int')
df['Amount'].dtypes

```
# change data type

df['Amount'] = df['Amount'].astype('int')

df['Amount'].dtypes

... dtype('int32')
```

#### **COLUMNS**: df.columns

#### $\underline{\textbf{DESCRIBE RETURNS DESCRIPTION OF THE DATA IN THE DATAFRAME}}: df. describe()$

df.	describe()				
	User_ID	Age	Marital_Status	Orders	Amount
count	1.123900e+04	11239.000000	11239.000000	11239.000000	11239.000000
mean	1.003004e+06	35.410357	0.420055	2.489634	9453.610553
std	1.716039e+03	12.753866	0.493589	1.114967	5222.355168
min	1.000001e+06	12.000000	0.000000	1.000000	188.000000
25%	1.001492e+06	27.000000	0.000000	2.000000	5443.000000
50%	1.003064e+06	33.000000	0.000000	2.000000	8109.000000
75%	1.004426e+06	43.000000	1.000000	3.000000	12675.000000
max	1.006040e+06	92.000000	1.000000	4.000000	23952.000000



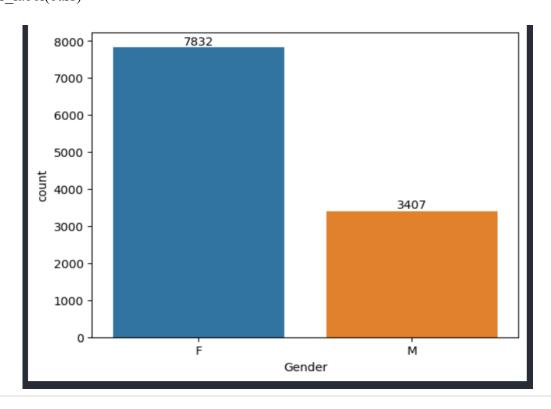
df[['Age', 'Orders', 'Amount']].describe()

D ~	# us	se describe()	for specific	columns					
	<pre>df[['Age', 'Orders', 'Amount']].describe()</pre>								
		Age	Orders	Amount					
	count	11239.000000	11239.000000	11239.000000					
	mean	35.410357	2.489634	9453.610553					
	std	12.753866	1.114967	5222.355168					
	min	12.000000	1.000000	188.000000					
	25%	27.000000	2.000000	5443.000000					
	50%	33.000000	2.000000	8109.000000					
	75%	43.000000	3.000000	12675.000000					
	max	92.000000	4.000000	23952.000000					

#### **EXPLORATORY DATA ANALYSIS:**

#### **GENDER GRAPH PLOTING:**

# plotting a bar chart for Gender and it's count
ax = sns.countplot(x = 'Gender',data = df)
for bars in ax.containers:
 ax.bar label(bars)



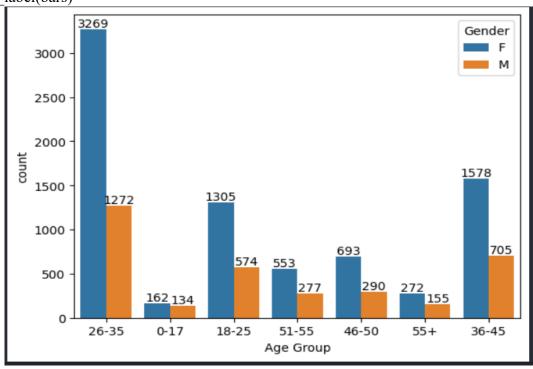


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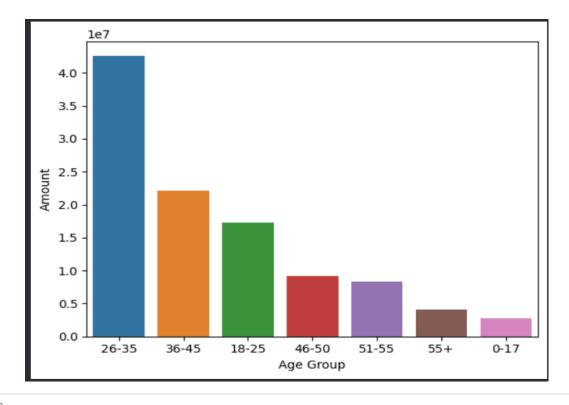
#### **AGE GRAPH PLOTING:**

ax = sns.countplot(data = df, x = 'Age Group', hue = 'Gender') for bars in ax.containers:

ax.bar label(bars)



# Total Amount vs Age Group sales\_age = df.groupby(['Age Group'], as\_index=False)['Amount'].sum().sort\_values(by='Amount', ascending=False)
sns.barplot(x = 'Age Group',y= 'Amount', data = sales\_age)



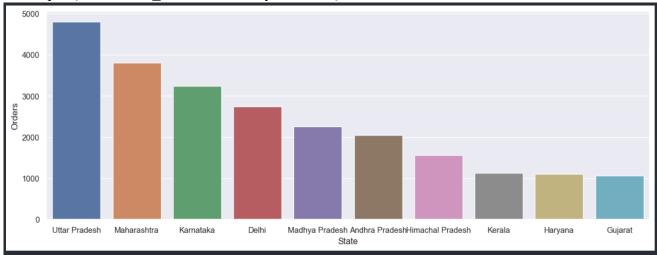


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#### **STATE GRAPH PLOTING:**

# total number of orders from top 10 states sales\_state = df.groupby(['State'], as\_index=False)['Orders'].sum().sort\_values(by='Orders', ascending=False).head(10) sns.set(rc={'figure.figsize':(15,5)})

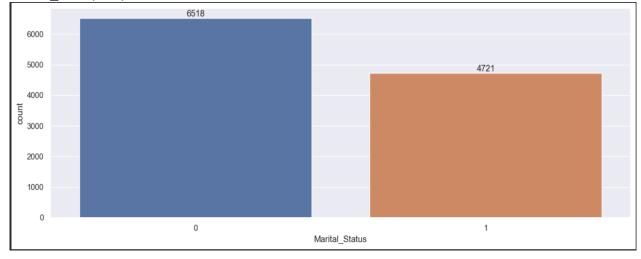
sns.barplot(data = sales state, x = 'State',y= 'Orders')



#### **MARITAL STATUS GRAPH PLOTING:**

ax = sns.countplot(data = df, x = 'Marital\_Status') sns.set(rc={'figure.figsize':(7,5)}) for bars in ax.containers:

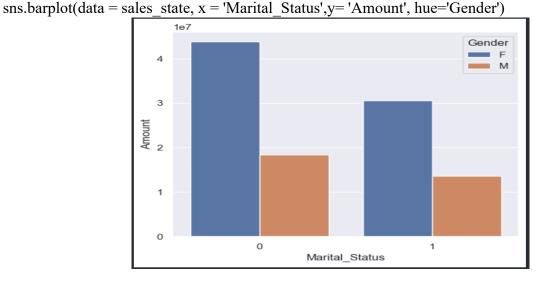
ax.bar label(bars)





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sales\_state = df.groupby(['Marital\_Status', 'Gender'],
as\_index=False)['Amount'].sum().sort\_values(by='Amount', ascending=False)
sns.set(rc={'figure.figsize':(6,5)})

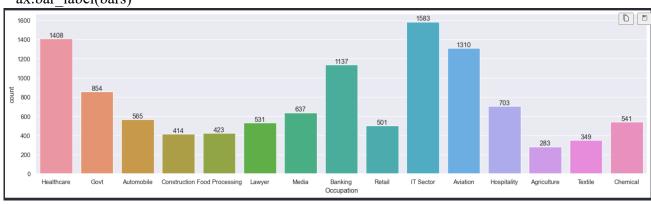


#### **OCCUPATION GRAPH PLOTING:**

sns.set(rc={'figure.figsize':(20,5)})
ax = sns.countplot(data = df, x = 'Occupation')

for bars in ax.containers:

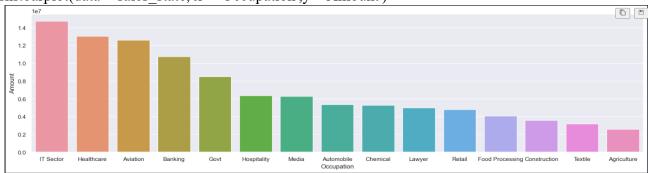
ax.bar label(bars)



Sales\_state = df.groupby(['Occupation'], as\_index=False)['Amount'].sum().sort\_values(by='Amount', ascending=False)

 $sns.set(rc = \{'figure.figsize': (20,5)\})$ 

sns.barplot(data = sales state, x = 'Occupation',y= 'Amount')



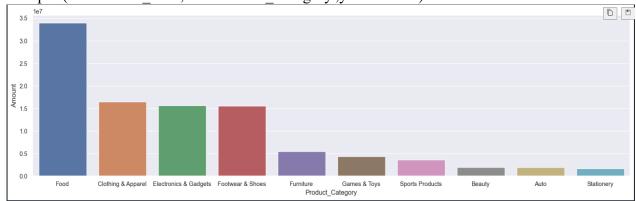


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#### PRODUCT CATEGORY GRAPH PLOTING:

 $sales\_state = df.groupby(['Product\_Category'], \\ as\_index=False)['Amount'].sum().sort\_values(by='Amount', ascending=False).head(10) \\ sns.set(rc=\{'figure.figsize':(20,5)\})$ 

sns.barplot(data = sales\_state, x = 'Product\_Category',y= 'Amount')

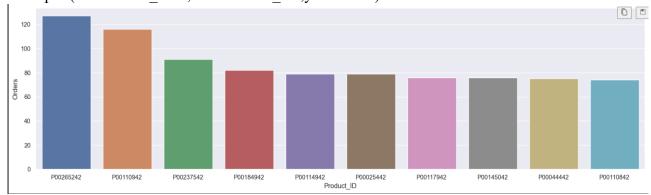


#### **PRODUCT ID GRAPH PLOTING:**

 $sales\_state = df.groupby(['Product\_ID'], as\_index=False)['Orders'].sum().sort\_values(by='Orders', ascending=False).head(10)$ 

sns.set(rc={'figure.figsize':(20,5)})

sns.barplot(data = sales state, x = 'Product ID',y='Orders')





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#### PRACTICAL - 3

❖ AIM: IMPLEMENT LINEAR REGRESSION AN LOGISTIC REGRESSION.

#### **PROCEDURE:**

#### • <u>LINEAR REGRESSION</u>:

- 1. Import Libraries:
  - Import necessary libraries such as NumPy, pandas, and scikit-learn.
- 2. Prepare the Data:
  - Load your dataset and split it into features (X) and target variable (y).
  - Optionally, split the data into training and testing sets.
- 3. Create the Model:
  - Instantiate the linear regression model.
- 4. Fit the Model:
  - Train the model using the training data.
- 5. Make Predictions:
  - Use the model to make predictions on the test set.
- 6. Evaluate the Model:
  - Assess the model's performance using metrics like Mean Squared Error (MSE) or R<sup>2</sup> score.
- 7. Visualize Results (Optional):
  - Plot the regression line against the data points.

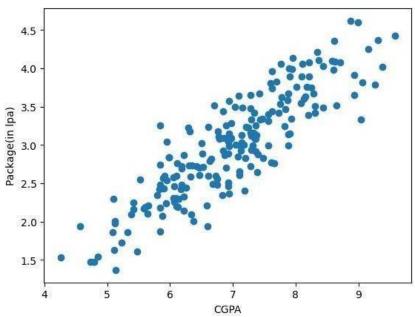
LOGISTIC REGRESSION :

- 1. Import Libraries:
  - Import necessary libraries such as NumPy, pandas, and scikit-learn.
- 2. Prepare the Data:
  - Load your dataset and split it into features (X) and target variable (y).
  - Convert the target variable to binary if necessary.
  - Optionally, split the data into training and testing sets.
- 3. Create the Model:
  - Instantiate the logistic regression model.
- 4. Fit the Model:
  - Train the model using the training data.
- 5. Make Predictions:
  - Use the model to make predictions on the test set.
- 6. Evaluate the Model:
  - Assess the model's performance using metrics like accuracy, confusion matrix, or ROC-AUC score.
- 7. Visualize Results (Optional):
  - Plot the decision boundary or ROC curve.



#### **❖** <u>PROGRAM</u>:

import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
df = pd.read\_csv('placement.csv')
df.head()
plt.scatter(df['cgpa'],df['package'])
plt.xlabel('CGPA')
plt.ylabel('Package(in lpa)')

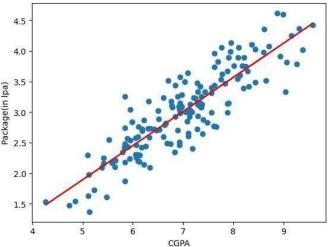


x=df.iloc[:,0:1]
y=df.iloc[:,1] y
x
from sklearn.model\_selection import train\_test\_split
x\_train,x\_test,y\_train,y\_test=train\_test\_split(x,y,test\_size=0.2,
random\_state=2)
from sklearn.linear\_model import LinearRegression
lr=LinearRegression()
lr.fit(x\_train,y\_train)

X\_test



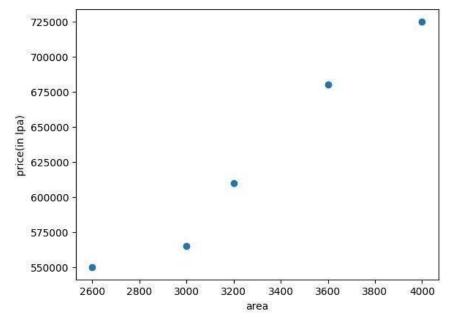
y\_test
lr.predict(x\_test.iloc[0].values.reshape(1,1))
plt.scatter(x\_train,y\_train)
plt.plot(x\_train,lr.predict(x\_train),color='red') plt.xlabel('CGPA')
plt.ylabel('Package(in lpa)')



 $m=lr.coef_a=[1,2,3,4,5,6,7,10,12,13,14,15,16,17,18,19,20,21,22,25,26,27,28,$ 29,30,31,32,34,35,36,37,38,39,40,42,44,45,46,47,48,49,50,51,52,53, 54,55,56,58,59,60,61,63,64,65,67,68,72,74] print(len(a)) b=lr.intercept  $_{\rm m} * 8.58 + b$ m\*9.5+bm\*100+b from sklearn.metrics import r2\_score, mean\_squared\_error, mean\_absolute\_error y\_pred=lr.predict(x\_test) score = r2\_score(y\_test,y\_pred) print(f'Accuracy Score : {score}') y\_pred score1=lr.score(x\_test,y\_test) print(f'Accracy Score:{score1}') MSE = mean\_squared\_error(y\_test,y\_pred) MSE MAE = mean\_absolute\_error(y\_test,y\_pred) MAE import numpy as np import pandas as pd import matplotlib.pyplot as plt df = pd.read\_csv('homeprices.csv') df.head() plt.scatter(df['area'],df['price']) plt.xlabel('area') plt.ylabel('price(in lpa)')



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x=df.iloc[:,0:1]
y=df.iloc[:,1] x
y
from sklearn.model\_selection import train\_test\_split
x\_train,x\_test,y\_train,y\_test=train\_test\_split(x,y,test\_size=0.2, random\_state=2)
from sklearn.linear\_model import LinearRegression

Logistic Linear Regression:import seaborn as sns import pandas as pd import numpy as np

lrr=LinearRegression

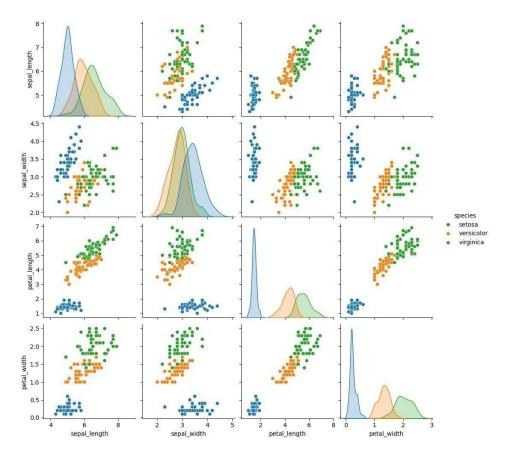
df=sns.load\_dataset('iris')
df.head()
df['species'].unique()
df.isnull().sum()

##EDA sns.pairplot(df,hue='species')



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df=df[df['species']!='setosa'] df.head()
df['species']=df['species'].map({'versicolor':0,'virginica':1}) df.head()
x=df.iloc[:,:-1]
y=df.iloc[:,1] x
y
from sklearn.model\_selection import train\_test\_split
X\_train,X\_test,y\_train,y\_test=train\_test\_split(x,y,test\_size=0.25, random\_state=42)
from sklearn.linear\_model import LogisticRegression
classifier=LogisticRegression()
classifier.fit(X\_train,y\_train)

### 

from sklearn.model\_selection import GridSearchCV parameters = {'penalty':['11','12','elasticnet'],'C':[1,2,3,4,5,6,10,20,30,40,50], 'max\_iter':[100,200,300]} classifier\_regressor = GridSearchCV(classifier, param\_grid=parameters, scoring='accuracy', cv=5) classifier\_regressor.fit(X\_train,y\_train)



/usr/local/lib/python3.10/dist-packages/sklearn/model\_selection/\_validation.py:528: FitFailedWarning: 330 fits failed out of a total of 495. The score on these train-test partitions for these parameters will be set to nan. If these failures are not expected, you can try to debug them by setting error\_score='raise'. Below are more details about the failures: 165 fits failed with the following error: Traceback (most recent call last): File "/usr/local/lib/python3.10/dist-packages/sklearn/model\_selection/\_validation.py", line 866, in \_fit\_and\_score estimator.fit(X\_train, y\_train, \*\*fit\_params) File "/usr/local/lib/python3.10/dist-packages/sklearn/base.py", line 1389, in wrapper return fit\_method(estimator, \*args, \*\*kwargs)
File "/usr/local/lib/python3.10/dist-packages/sklearn/linear\_model/\_logistic.py", line 1193, in fit solver = \_check\_solver(self.solver, self.penalty, self.dual) File "/usr/local/lib/python3.10/dist-packages/sklearn/linear\_model/\_logistic.py", line 63, in \_check\_solver raise ValueError( ValueError: Solver lbfgs supports only 'l2' or None penalties, got l1 penalty. 165 fits failed with the following error: Traceback (most recent call last): File "/usr/local/lib/python3.10/dist-packages/sklearn/model\_selection/\_validation.py", line 866, in \_fit\_and\_score estimator.fit(X\_train, y\_train, \*\*fit\_params) File "/usr/local/lib/python3.10/dist-packages/sklearn/base.py", line 1389, in wrapper nan 0.96 nan nan 0.96 nan nan 0.96 nan 0.96 nan nan nan 0.96 warnings.warn(

Output is truncated. View as a <u>scrollable element</u> or open in a <u>text editor</u>. Adjust cell output <u>settings</u>...

print(classification report(y test,y pred))

precision	recall f1	-score	support		
0 0.9	4 0.94	0.94	16		
1 0.8	9 0.89	0.89	9		
accuracy		0.92	25		
macro avg	0.91	).91 C	).91	25	
weighted avg	0.92	0.92	0.92	25	

from sklearn.metrics import confusion\_matrix cm =
confusion\_matrix(y\_test,y\_pred)
print(cm)

[[15 1] [1 8]]



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import matplotlib.pyplot as plt

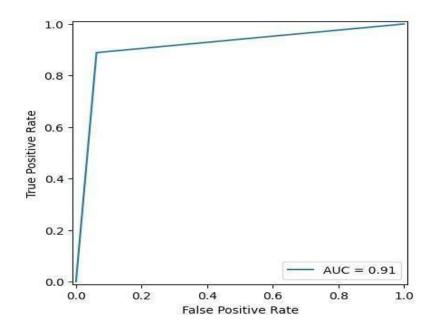
from sklearn.metrics import roc\_curve, roc\_auc\_score, RocCurveDisplay fpr, tpr,

 $thresholds = roc\_curve(y\_test, \, y\_pred)$ 

auc\_score = roc\_auc\_score(y\_test, y\_pred)

RocCurveDisplay(fpr=fpr, tpr=tpr, roc\_auc=auc\_score).plot()

plt.show()





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#### PRACTICAL - 4

❖ <u>AIM</u>: IMPLEMENT THE NAIVE BAYESIAN CLASSIFIER FOR A SAMPLE TRAINING DATA SET STORED AS A (.CSV) FILE. COMPUTE THE ACCURACY OF THE CLASSIFIER, CONSIDERING A FEW TEST DATA SETS.

#### **PROCEDURE:**

To implement a naïve Bayesian classifier using a sample training dataset in CSV format, follow these general steps:

- 1. Load the CSV data into your programming environment.
- 2. Preprocess the data (handle missing values, encode categorical variables).
- 3. Split the data into training and testing sets.
- 4. Train the Naïve Bayes model on the training data.
- 5. Make predictions on the test data.
- 6. Compute the accuracy by comparing predicted labels with actual labels.

#### • PREPROCESS THE DATA:

- Encode categorical variables using **LabelEncoder** or one-hot encoding.
- Convert all attributes to floating-point numbers for numerical processing.

#### • SPLIT THE DATA:

• Divide the dataset into training and testing sets using a specified ratio (e.g., 70% training, 30% testing).

#### • TRAIN THE NAÏVE BAYES MODEL:

• Calculate the mean and standard deviation for each attribute in each class from the training data.

#### **■** MAKE PREDICTIONS:

- Use the trained model to predict the class labels for the test data.
- Load and preprocess the dataset.
- Split the data into training and testing sets.
- Train the Naïve Bayes model and make predictions.
- Calculate and display the accuracy of the model.

#### **PROGRAM:**

import numpy as np import pandas as pd

import matplotlib.pyplot as plt

from sklearn.model\_selection import train\_test\_split from sklearn.naive\_bayes import GaussianNB

np.set printoptions(suppress=True, precision=6)

df = pd.read csv("titanic-data.csv")

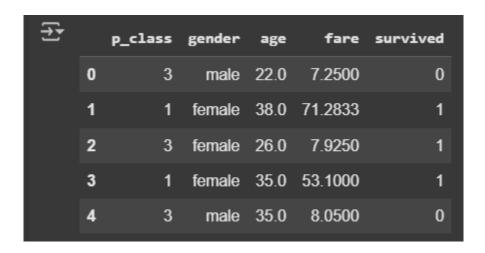
df.head()





df.drop(["passenger\_id", "name", "sib\_sp", "parch", "ticket", "cabin", "embarked"], axis=1, inplace=True)

df.head()



target = df["survived"]

inputs = df.drop("survived", axis=1)

display(target.head())

display(inputs.head())





dummies = pd.get dummies(inputs["gender"])

display(dummies.head())
print(dummies.dtypes)



inputs = pd.concat([inputs, dummies], axis=1)
inputs.head()

<del>∑</del> *		p_class	gender	age	fare	survived
	0	3	male	22.0	7.2500	0
	1	1	female	38.0	71.2833	1
	2	3	female	26.0	7.9250	1
	3	1	female	35.0	53.1000	1
	4	3	male	35.0	8.0500	0

inputs.drop(["gender"], axis=1, inplace=True)
inputs.head()

_						
₹		p_class	age	fare	female	male
	0	3	22.0	7.2500	False	True
	1	1	38.0	71.2833	True	False
	2	3	26.0	7.9250	True	False
	3	1	35.0	53.1000	True	False
	4	3	35.0	8.0500	False	True

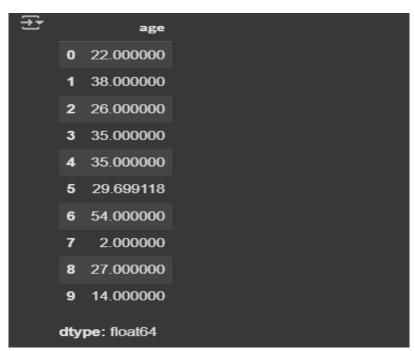


inputs.age[:10]



inputs["age"] = inputs["age"].fillna(inputs["age"].mean())

inputs.age[:10]



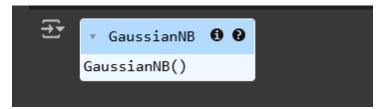
X\_train, X\_test, y\_train, y\_test = train\_test\_split(inputs, target, test\_size=0.2) print(len(X\_train), len(X\_test), len(inputs))

print(len(X\_train) / len(inputs))
print(len(X\_test) / len(inputs))

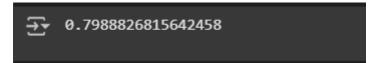
712 179 891 0.7991021324354658 0.20089786756453423



model = GaussianNB() model.fit(X train, y train)



model.score(X\_test, y\_test)



pred = np.array(model.predict(X\_test))
pred\_probability = np.array(model.predict\_proba(X\_test))
print(pred[:5])
for i in range(1, 6):
 print(pred\_probability[i][0],end=", ")

(0 0 1 0 1) 0.9895202768716742, 0.003675565181087268, 0.9908563111883015, 0.036794007058115134, 0.9787398295999188,



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#### PRACTICAL - 5

❖ <u>AIM</u>: ASSUMING A SET OF DOCUMENTS THAT NEED TO BE CLASSIFIED, USE THE NAIVE BAYESIAN CLASSIFIER MODEL TO PERFORM THIS TASK.

#### **PROCEDURE:**

#### DATA COLLECTION :

• Gather Documents: Collect a labeled dataset of documents. Each document should be associated with a class label (e.g., categories like "sports," "politics," "technology," etc.).

#### • PREPROCESSING:

- Text Cleaning: Remove any irrelevant information, such as HTML tags, special characters, and numbers.
- Tokenization: Split the text into individual words or tokens.
- Lowercasing: Convert all text to lowercase to ensure uniformity.
- Stop Word Removal: Remove common words (e.g., "and," "the," "is") that do not contribute to the meaning.
- Stemming/Lemmatization: Reduce words to their base or root form (e.g., "running" to "run").

#### FEATURE EXTRACTION :

- Create a Vocabulary: Build a list of unique words (features) from the training documents.
- Vectorization: Convert documents into numerical format using techniques like:
  - Bag of Words (BoW): Count the frequency of each word in the document.
  - Term Frequency-Inverse Document Frequency (TF-IDF): Weigh the frequency of words based on their importance across all documents.

#### • SPLITTING THE DATASET:

• Train-Test Split: Divide the dataset into a training set and a test set (e.g., 80% training, 20% testing).

#### **❖** PROGRAM :

import pandas as pd from sklearn.model\_selection import train\_test\_split from sklearn.feature\_extraction.text import CountVectorizer from sklearn.naive\_bayes import MultinomialNB from sklearn.metrics import accuracy\_score, classification\_report

msg=pd.read\_csv('document.csv', names=['message','label'])
print("Total Instances of dataset")
msg['labelnum']=msg['label'].map({'pos':1,'neg':0})
X=msg['message']
y=msg['label'] print(msg)



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Tota	al Instances of dataset		
	message	label	labelnum
0	I love this sandwich	pos	1
1	This is an amazing place	pos	1
2	I feel very good about these beers	pos	1
3	This is my best work	pos	1
4	What an awesome view	pos	1
5	I do not like this restaurant	neg	0
6	I am tired of this stuff	neg	0
7	I can't deal with this	neg	0
8	He is my sworn enemy	neg	0
9	My boss is horrible	neg	0
10	This is an awesome place	pos	1
11	I do not like the taste of this juice	neg	0
12	I love to dance	pos	1
13	I am sick and tired of this place	neg	0
14	What a great holiday	pos	1
15	That is a bad locality to stay	neg	0
16	We will have good fun tomorrow	pos	1
17	I went to my enemy's house today	neg	0

Xtrain, Xtest, ytrain, ytest = train\_test\_split(X, y)

vectorizer = CountVectorizer()

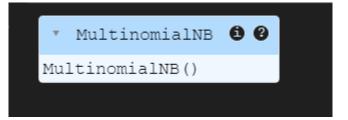
Xtrain\_dm = vectorizer.fit\_transform(Xtrain)

Xtest\_dm = vectorizer.transform(Xtest)

Xtrain\_dm

<13x39 sparse matrix of type '<class 'numpy.int64'>'
 with 65 stored elements in Compressed Sparse Row format>

clf=MultinomialNB()
clf.fit(Xtrain\_dm, ytrain)





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from sklearn.metrics import accuracy\_score, confusion\_matrix pred =
clf.predict(Xtest\_dm)
print('Accuracy: ', accuracy\_score(ytest, pred)) print('Confusion
Matrix:\n', confusion\_matrix(ytest, pred))

Accuracy: 0.6
Confusion Matrix:
[[1 2]
[0 2]]

pred

```
array(['pos', 'pos', 'pos', 'neg'], dtype='<U3')
```

```
user_input = input("Enter a message to predict its sentiment: ") user_input_dm =
vectorizer.transform([user_input])
user_pred = clf.predict(user_input_dm) sentiment =
'pos' if user_pred[0] == 1 else 'neg'
print(f"The sentiment of your message is: {sentiment}")
```

Enter a message to predict its sentiment: men don't celebrate valentine day The sentiment of your message is: neg



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## PRACTICAL - 6

❖ <u>AIM</u>: DECISION TREE BASED ID3 ALGORITHM.

## **PROCEDURE:**

- Start with the Entire Dataset :Begin with the entire dataset as the root node of the tree.
- Check for Stopping Conditions:
  - If all instances in the dataset belong to the same class, create a leaf node with that class label.
  - If there are no remaining attributes to split on, create a leaf node with the majority class label of the instances in the dataset.
- Calculate Information Gain :

For each attribute in the dataset, calculate the Information Gain (IG) based on the entropy of the dataset before and after the split.

- Entropy is calculated using the formula:
  - [\text{Entropy}(S) = -\sum\_{i=1}^{c} p\_i \log\_2(p\_i)] where  $(p_i)$  is the proportion of instances belonging to class (i) in the dataset (S), and (c) is the number of classes.
- Information Gain is calculated as :
  - $[IG(S, A) = \text{Entropy}(S) \sum_{v \in \{V \in \{V \in \{V \in \{S_v\}\} \} } | frac\{|S_v|\} \{|S|\} \}$  \text{Entropy}(S\_v) ] where (S\_v) is the subset of (S) for which attribute (A) has value (v).
- Select the Best Attribute :
  - Choose the attribute with the highest Information Gain to split the dataset. This attribute will become the decision node in the tree.
- Create Subsets:
  - Split the dataset into subsets based on the values of the selected attribute.
- Recursively Build the Tree :
  - For each subset, repeat steps 2 to 5:
  - Use the subset as the new dataset.
  - Remove the selected attribute from consideration for further splits.
  - Continue until stopping conditions are met.
- Construct the Tree:
  - As the recursion unwinds, construct the decision tree by linking nodes and leaf nodes based on the splits made.
- Output the Decision Tree:
  - Once the recursion is complete, the final decision tree can be outputted, which can then be used for classification of new instances.

### $\bullet$ PROGRAM:

STEP 1: IMPORT NECESSARY LIBRARIES

import math

import pandas as pd

from graphviz import Digraph

from IPython.display import Image



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#### STEP 2: FUNCTION TO CALCULATE ENTROPY

```
def calculate_entropy(dataset):
    class_counts = dataset.iloc[:, -1].value_counts()
    total_instances = len(dataset)
    entropy = 0
    for count in class_counts:
        probability = count / total_instances
        entropy -= probability * math.log2(probability)
    return entropy
```

#### STEP 3: FUNCTION TO CALCULATE INFORMATION GAIN

```
def calculate_information_gain(dataset, attribute):
    total_entropy = calculate_entropy(dataset)
    attribute_values = dataset[attribute].unique()
    weighted_entropy = 0
    for value in attribute_values:
        subset = dataset[dataset[attribute] == value]
        weighted_entropy += (len(subset) / len(dataset)) * calculate_entropy(subset)
    return total_entropy - weighted_entropy
```

#### STEP 4: FUNCTION TO BUILD THE DECISION TREE

```
def build_decision_tree(dataset, attributes, parent_node=None, graph=None): # Build the decision tree
if graph is None:
    graph = Digraph('DecisionTree')
if len(dataset.iloc[:, -1].unique()) == 1: # Make Attributes of in image
    leaf = dataset.iloc[0, -1]
    graph.node(parent node, label=leaf, shape='ellipse', style='filled', color='lightblue')
```

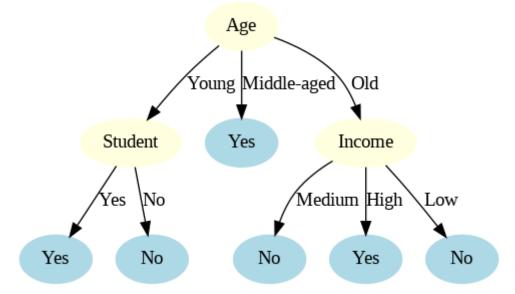


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```
return leaf
  if len(attributes) == 0:
     majority class = dataset.iloc[:, -1].mode()[0]
     graph.node(parent node, label=majority class, shape='ellipse', style='filled', color='lightgreen')
     return majority class
  best attribute = max(attributes, key=lambda attr: calculate information gain(dataset, attr))
  graph.node(parent node, label=best attribute, shape='ellipse', style='filled', color='lightyellow')# This
Main Entity
  attribute values = dataset[best attribute].unique()
  for value in attribute values:
     subset = dataset[dataset[best attribute] == value].drop(columns=[best attribute])
     child node = f'{parent node} {value}'
     graph.edge(parent node, child node, label=value)
     build decision tree(subset, [attr for attr in attributes if attr != best attribute], child node, graph)
  return graph
STEP 5: EXAMPLE USAGE
data = {
  'Age': ['Young', 'Young', 'Middle-aged', 'Old', 'Old', 'Old'],
  'Income': ['High', 'High', 'Low', 'Medium', 'High', 'Low'],
  'Student': ['Yes', 'No', 'No', 'Yes', 'Yes', 'No'],
  'Buy Computer': ['Yes', 'No', 'Yes', 'No', 'Yes', 'No']
}
dataset = pd.DataFrame(data)
attributes = ['Age', 'Income', 'Student']
STEP 6: BUILD AND VISUALIZE THE DECISION TREE
graph = build decision tree(dataset, attributes, 'Root')
graph.render('decision tree', format='png')
Image(filename='decision tree.png')
```



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## PRACTICAL - 7

❖ <u>AIM</u>: WRITE A PROGRAM TO IMPLEMENT THE K-NEAREST NEIGHBOUR ALGORITHM TO CLASSIFY THE IRIS DATA SET.

### **PROCEDURE:**

- IMPLEMENT K-NEAREST NEIGHBORS (KNN) FOR THE IRIS DATASET :
  - 1. Import Required Libraries
    Import necessary Python libraries such as numpy, pandas, matplotlib, seaborn, and sklearn.
  - 2. Load the Iris Dataset
    Use sklearn.datasets.load iris() to load the dataset and convert it into a DataFrame.
  - 3. Preprocess the Data
    - Split the dataset into features (X) and target labels (y).
    - Normalize or standardize data if necessary.
    - Split the data into training and testing sets using train test split().
  - 4. Implement K-Nearest Neighbors (KNN) Algorithm
    - Use KNeighborsClassifier from sklearn.neighbors.
    - Choose an appropriate value for K (e.g., K=5).
    - Fit the model with training data.
  - 5. Make Predictions & Evaluate Model
    - Predict the class labels for test data.
    - Compute accuracy using accuracy score().
    - Display confusion matrix and classification report.



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## **PROGRAM**:

Step 1: IMPORTING NECESSARY LIBRARIES

import pandas as pd import matplotlib.pyplot as plt import seaborn as sns from sklearn.datasets import load\_iris from sklearn.model\_selection import train\_test\_split from sklearn.preprocessing import StandardScaler from sklearn.neighbors import KNeighborsClassifier from sklearn matrics import accuracy, score, confu

from sklearn.metrics import accuracy\_score, confusion\_matrix, classification\_report

Step 2: LOAD THE IRIS DATASET

iris = load\_iris()

X = iris.data # Features

y = iris.target # Labels (species)

Step 3: DATA PREPROCESSING

scaler = StandardScaler()
X\_scaled = scaler.fit\_transform(X)

Step 4: SPLIT THE DATA INTO TRAINING AND TESTING SETS (80% TRAIN, 20% TEST)

X train, X test, y train, y test = train test split(X scaled, y, test size=0.2, random state=42)

Step 5: TRAIN THE KNN MODEL

knn = KNeighborsClassifier(n\_neighbors=5) # You can change the number of neighbors knn.fit(X train, y train)



Step 6: MAKE PREDICTIONS

y pred = knn.predict(X test)

Step 7: EVALUATE THE MODEL



accuracy = accuracy\_score(y\_test, y\_pred)

print(f'Accuracy: {accuracy \* 100:.2f}%') Output: Accuracy : 100.00%

#### Step 8: CONFUSION MATRIX

print("\nConfusion Matrix:")

print(confusion\_matrix(y\_test, y\_pred))

Confusion Matrix:
[[10 0 0]
 [ 0 9 0]
 [ 0 0 11]]

#### Step 9: CLASSIFICATION REPORT

print("\nClassification Report:")

print(classification\_report(y\_test, y\_pred))

Classification Report: precision recall f1-score support 0 1.00 1.00 1.00 10 1.00 1.00 1.00 1 9 1.00 1.00 1.00 11 1.00 30 accuracy 1.00 1.00 1.00 30 macro avg 1.00 1.00 30 weighted avg 1.00

#### Step 10: VISUALIZE THE CONFUSION MATRIX

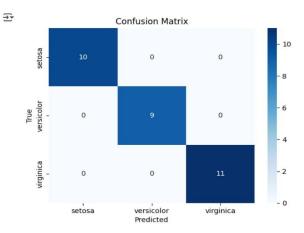
cm = confusion matrix(y test, y pred)

sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=iris.target\_names, yticklabels=iris.target\_names) plt.xlabel('Predicted')

plt.ylabel('True')

plt.title('Confusion Matrix')

plt.show()





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## **PRACTICAL - 8**

❖ <u>AIM</u>: APPLY EM ALGORITHM TO CLUSTER A SET OF DATA STORED IN A (.CSV) FILE. USE THE SAME DATA SET FOR CLUSTERING USING K-MEANS ALGORITHM.

## **PROCEDURE:**

To apply the Expectation-Maximization (EM) Algorithm and K-Means Algorithm to cluster a given dataset stored in a .csv file and compare their results.

## **PROGRAM:**

Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for clustering using k-Means algorithm.

#### **Step 1: Import Necessary Libraries**

# Importing required libraries

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

from sklearn.preprocessing import StandardScaler

from sklearn.cluster import KMeans

from sklearn.mixture import GaussianMixture

from sklearn.metrics import silhouette score

- Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for clustering using k-Means algorithm.
- Step 1: Import Necessary Libraries
- # Importing required libraries import pandas as pd import numpy as np import matplotlib.pyplot as plt from sklearn.preprocessing import StandardScaler from sklearn.cluster import KMeans from sklearn.mixture import GaussianMixture from sklearn.metrics import silhouette\_score



#### **Step 2: Load the Dataset from CSV**

Make sure you upload your .CSV file to Google Colab before running the following code. If your file is named data.csv, you can load it as follows.

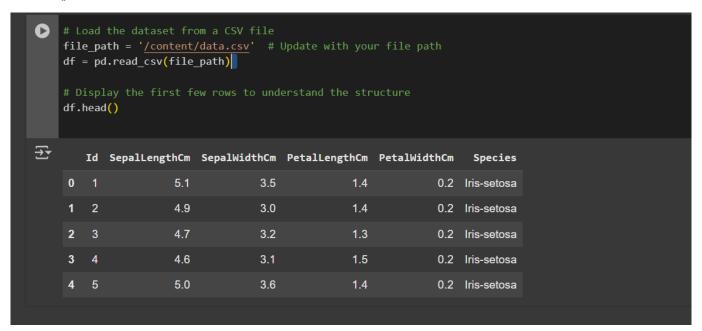
# Load the dataset from a CSV file

file path = '/content/data.csv' # Update with your file path

df = pd.read csv(file path)

# Display the first few rows to understand the structure

df.head()



#### **Step 3: Data Preprocessing.**

We will standardize the data (i.e., mean = 0, standard deviation = 1), as both K-Means and GMM benefit from scaling for better performance.

# Dropping non-numeric columns if any (adjust according to your data)

df = df.select dtypes(include=[np.number])

# Standardize the data (important for K-Means and GMM)

scaler = StandardScaler()

X scaled = scaler.fit transform(df)

# Check the scaled data

print(X\_scaled[:5]) # First 5 rows of scaled data



```
# Dropping non-numeric columns if any (adjust according to your data)

df = df.select_dtypes(include=[np.number])

# Standardize the data (important for K-Means and GMM)

scaler = StandardScaler()

X_scaled = scaler.fit_transform(df)

# Check the scaled data

print(X_scaled[:5]) # First 5 rows of scaled data

[-1.72054204 -0.90068117 1.03205722 -1.3412724 -1.31297673]

[-1.69744751 -1.14301691 -0.1249576 -1.3412724 -1.31297673]

[-1.67435299 -1.38535265 0.33784833 -1.39813811 -1.31297673]

[-1.65125846 -1.50652052 0.10644536 -1.2844067 -1.31297673]

[-1.62816394 -1.02184904 1.26346019 -1.3412724 -1.31297673]
```

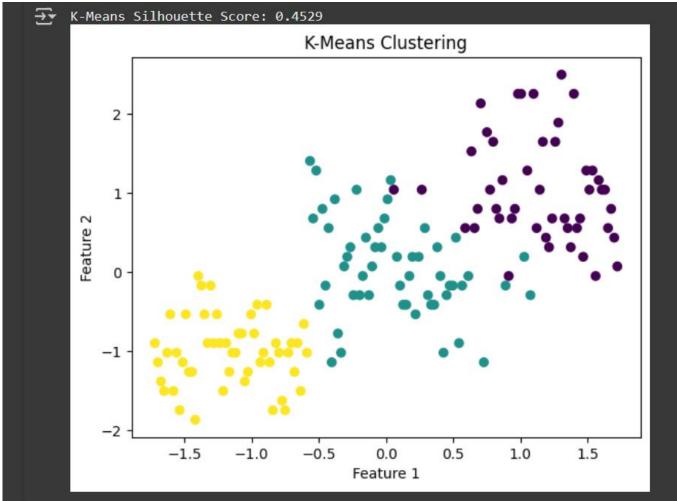
#### **Step 4: Apply K-Means Clustering**

Now, we'll use K-Means to cluster the data. First, we will try clustering for a given number of clusters, say k=3. You can adjust this based on your dataset.

```
# Apply K-Means clustering
kmeans = KMeans(n_clusters=3, random_state=42)
kmeans.fit(X_scaled)
# Assign labels to each point
kmeans_labels = kmeans.predict(X_scaled)
# Evaluate the clustering with silhouette score
silhouette_kmeans = silhouette_score(X_scaled, kmeans_labels)
print(f'K-Means Silhouette Score: {silhouette_kmeans:.4f}")
# Plot the clusters
plt.scatter(X_scaled[:, 0], X_scaled[:, 1], c=kmeans_labels, cmap='viridis')
plt.title('K-Means Clustering')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.show()
```



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#### **Step 5: Apply Expectation-Maximization (GMM)**

The Expectation-Maximization algorithm is typically implemented using Gaussian Mixture Models in sklearn. Here, we will cluster the data using a GMM.

```
# Apply Gaussian Mixture Model (GMM) for clustering
gmm = GaussianMixture(n_components=3, random_state=42)
gmm.fit(X_scaled)
# Assign labels to each point
gmm_labels = gmm.predict(X_scaled)
# Evaluate the clustering with silhouette score
silhouette_gmm = silhouette_score(X_scaled, gmm_labels)
print(f'GMM Silhouette Score: {silhouette_gmm:.4f}")
# Plot the clusters
plt.scatter(X_scaled[:, 0], X_scaled[:, 1], c=gmm_labels, cmap='viridis')
```



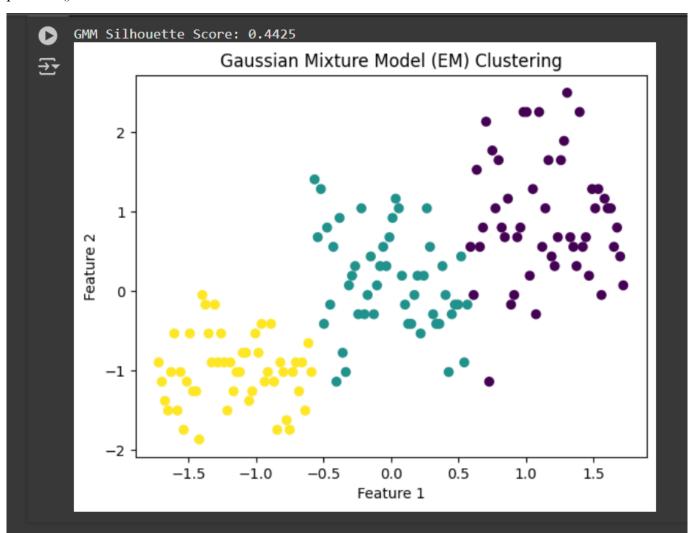
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plt.title('Gaussian Mixture Model (EM) Clustering')

plt.xlabel('Feature 1')

plt.ylabel('Feature 2')

plt.show()



#### **Step 6: Compare Results**

Now, let's compare the clustering results visually and numerically using the silhouette scores for each method.

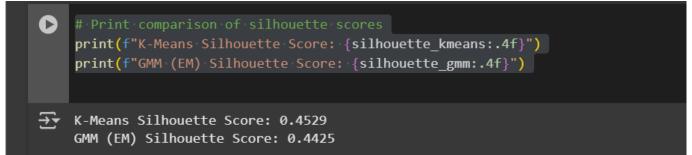
# Print comparison of silhouette scores

print(f"K-Means Silhouette Score: {silhouette kmeans:.4f}")

print(f"GMM (EM) Silhouette Score: {silhouette\_gmm:.4f}")



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## PRACTICAL - 9

❖ <u>AIM</u>: WRITE A PROGRAM TO CONSTRUCT A BAYESIAN NETWORK CONSIDERING MEDICAL DATA. USE THIS MODEL TO DEMONSTRATE THE DIAGNOSIS OF HEART PATIENTS USING STANDARD HEART DISEASE DATA SET.

### **PROCEDURE**:

To construct a Bayesian Network using medical data and use this model to diagnose heart disease based on the Standard Heart Disease Dataset.

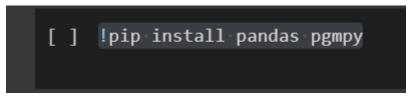
## **PROGRAM**:

Write a program to construct a Bayesian network considering medical data. Use this model to demonstrate the diagnosis of heart patients using standard Heart Disease Data Set.

#### **Step 1: Install Required Libraries**

In a new Google Colab notebook, the first thing you'll need to do is install the necessary libraries (pandas and pgmpy). Add the following code in a cell:

!pip install pandas pgmpy



#### **Step 2: Import Required Libraries**

import pandas as pd

import matplotlib.pyplot as plt

import networkx as nx

from pgmpy.models import BayesianModel

from pgmpy.estimators import MaximumLikelihoodEstimator

from pgmpy.inference import VariableElimination

#### **Step 3: Upload the Dataset**

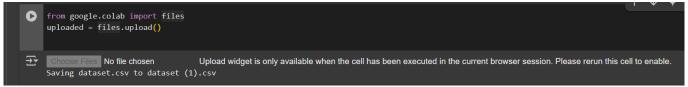
To upload your dataset into Colab, run the following code to prompt the file upload:

from google.colab import files

uploaded = files.upload()



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#### Step 4: Load the Data

Load the uploaded dataset into a pandas DataFrame:

data = pd.read csv("/content/dataset.csv") # Ensure the path matches your file's name heart disease = pd.DataFrame(data)

```
data = pd.read_csv("/content/dataset.csv") # Ensure the path matches your file's name
heart disease = pd.DataFrame(data)
```

#### **Step 5: Define and Fit the Bayesian Network**

Define the structure of the Bayesian Network and fit it with the data using MaximumLikelihoodEstimator:

# Define the Bayesian Network structure

```
model = BayesianModel([
  ('age', 'Lifestyle'),
  ('Gender', 'Lifestyle'),
  ('Family', 'heartdisease'),
  ('diet', 'cholestrol'),
  ('Lifestyle', 'diet'),
  ('cholestrol', 'heartdisease'),
  ('diet', 'cholestrol')
1)
# Fit the model using MaximumLikelihoodEstimator
model.fit(heart disease, estimator=MaximumLikelihoodEstimator)
```



```
# Define the Bayesian Network structure
model = BayesianModel([
    ('age', 'Lifestyle'),
    ('Gender', 'Lifestyle'),
    ('Family', 'heartdisease'),
    ('diet', 'cholestrol'),
    ('Lifestyle', 'diet'),
    ('cholestrol', 'heartdisease'),
    ('diet', 'cholestrol')
])

# Fit the model using MaximumLikelihoodEstimator
model.fit(heart_disease, estimator=MaximumLikelihoodEstimator)

**WARNING:pgmpy:BayesianModel has been renamed to BayesianNetwork. Please use BayesianNetwork class, BayesianModel will be removed in future.
```

#### **Step 6: Perform Inference and Predict Heart Disease**

```
Take user input for the various factors, perform inference, and display the prediction:
HeartDisease infer = VariableElimination(model)
# Instructions for user input
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 }')
try:
  age = int(input('Enter age (0: SuperSeniorCitizen, 1: SeniorCitizen, 2: MiddleAged, 3: Youth, 4: Teen):
'))
  gender = int(input('Enter Gender (0: Male, 1: Female): '))
  family history = int(input('Enter Family History (1: Yes, 0: No): '))
  diet = int(input('Enter Diet (0: High, 1: Medium): '))
  lifestyle = int(input('Enter Lifestyle (0: Athlete, 1: Active, 2: Moderate, 3: Sedentary): '))
  cholestrol = int(input('Enter Cholestrol (0: High, 1: BorderLine, 2: Normal): '))
  # Query the model for heart disease prediction
  q = HeartDisease infer.query(variables=['heartdisease'], evidence={
     'age': age,
     'Gender': gender,
     'Family': family history,
```



'diet': diet, 'Lifestyle': lifestyle, 'cholestrol': cholestrol # Display the heart disease prediction print("Heart Disease Prediction:") for state, prob in zip(q.values, q.state names['heartdisease']): print(f"{state}: {prob:.4f}") # Plot the inference results as a bar chart plt.bar(q.state names['heartdisease'], q.values) plt.xlabel('Heart Disease Status') plt.ylabel('Probability') plt.title('Heart Disease Prediction Probability')

except ValueError:

plt.show()

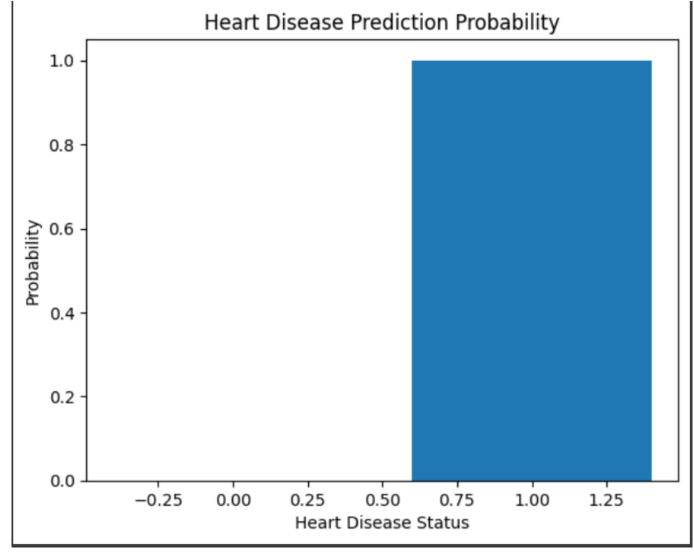
})

print("Invalid input. Please enter valid integers for the options.")

```
For age Enter { SuperSeniorCitizen:0, SeniorCitizen:1, MiddleAged:2, Youth:3, Teen:4 } For Gender Enter { Male:0, Female:1 }
For Family History Enter { yes:1, No:0 }
For diet Enter { High:0, Medium:1 }
For diet Enter { High:0, Medium:1 }
For lifeStyle Enter { Athlete:0, Active:1, Moderate:2, Sedentary:3 }
For cholesterol Enter { High:0, BorderLine:1, Normal:2 }
Enter age (0: SuperSeniorCitizen, 1: SeniorCitizen, 2: MiddleAged, 3: Youth, 4: Teen): 4
Enter Gender (0: Male, 1: Female): 0
Enter Family History (1: Yes, 0: No): 1
Enter Diet (0: High, 1: Medium): 0
Enter Lifestyle (0: Athlete, 1: Active, 2: Moderate, 3: Sedentary): 1
Enter Cholestrol (0: High, 1: BorderLine, 2: Normal): 0
WARNING:pgmpy:BayesianModel has been renamed to BayesianNetwork. Please use BayesianNetwork class, BayesianModel will be removed in future.
WARNING:pgmpy:BayesianModel has been renamed to BayesianNetwork. Please use BayesianNetwork class, BayesianModel will be removed in future.
Heart Disease Prediction:
0.0: 0.0000
```



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## PRACTICAL - 10

❖ <u>AIM</u>: COMPARE THE VARIOUS SUPERVISED LEARNING ALGORITHM BY USING APPROPRIATE DATASET.(LINEAR REGRESSION, SUPPORT VECTOR MACHINE, DECISION TREE).

## **PROCEDURE:**

- 1. IMPORT REQUIRED LIBRARIES:
  - Load essential libraries such as NumPy, Pandas, Matplotlib, Seaborn, and Scikit-learn.
- 2. SELECT AN APPROPRIATE DATASET:
  - Choose a dataset suitable for regression (e.g., California Housing Dataset) or classification (e.g., Iris Dataset).
- 3. PREPROCESS THE DATA:
  - Handle missing values, normalize/standardize the data, and split it into training and testing sets (typically 80% train, 20% test).
- 4. TRAIN THE MODELS:
  - Train Linear Regression for regression tasks.
  - Train Support Vector Machine (SVM) using a linear kernel.
  - Train Decision Tree with an appropriate depth.
- 5. EVALUATE MODEL PERFORMANCE:
  - Use appropriate metrics like Mean Squared Error (MSE) for regression models and Accuracy Score for classification models.
- 6. COMPARE THE MODELS:
  - Visualize the results using bar plots to compare the errors or accuracies of each algorithm.
- 7. DRAW CONCLUSIONS:
  - Identify which algorithm performs best based on the dataset characteristics and chosen evaluation metrics.



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#### **PROGRAM**:

# Importing required libraries
import numpy as np
import pandas as pd
from sklearn import datasets
from sklearn.model\_selection import train\_test\_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear\_model import LogisticRegression
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy\_score
import matplotlib.pyplot as plt

# Load the dataset
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']

# Split the dataset into training and testing sets (80% training, 20% testing)

X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.2, random\_state=42)

scaler = StandardScaler()

X\_train\_scaled = scaler.fit\_transform(X\_train)

X\_test\_scaled = scaler.transform(X\_test)

# Initialize the models (Logistic Regression, SVM, Decision Tree) lr = LogisticRegression(max\_iter=200) svm = SVC(kernel='linear') # Linear kernel for simplicity dt = DecisionTreeClassifier(random\_state=42)

# Train the models
lr.fit(X\_train\_scaled, y\_train)
svm.fit(X\_train\_scaled, y\_train)
dt.fit(X\_train, y\_train) # Decision Tree is not sensitive to feature scaling

```
/usr/local/lib/python3.11/dist-packages/sklearn/utils/validation.py:1408: DataC y = column_or_1d(y, warn=True)
/usr/local/lib/python3.11/dist-packages/sklearn/utils/validation.py:1408: DataC y = column_or_1d(y, warn=True)

DecisionTreeClassifier

DecisionTreeClassifier(random_state=42)
```



**#Predictions** 

y\_pred\_lr = lr.predict(X\_test\_scaled)
y\_pred\_svm = svm.predict(X\_test\_scaled)
y\_pred\_dt = dt.predict(X\_test)

# Evaluate the models using accuracy score acc\_lr = accuracy\_score(y\_test, y\_pred\_lr) acc\_svm = accuracy\_score(y\_test, y\_pred\_svm) acc\_dt = accuracy\_score(y\_test, y\_pred\_dt)

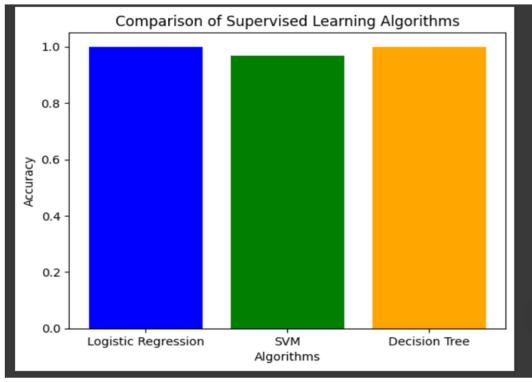
# Print the accuracy scores for each model print(f"Logistic Regression Accuracy: {acc\_lr:.4f}") print(f"Support Vector Machine Accuracy: {acc\_svm:.4f}") print(f"Decision Tree Accuracy: {acc\_dt:.4f}")

Logistic Regression Accuracy: 1.0000
Support Vector Machine Accuracy: 0.9667
Decision Tree Accuracy: 1.0000

# Visualize the accuracy comparison
algorithms = ['Logistic Regression', 'SVM', 'Decision Tree']
accuracies = [acc\_lr, acc\_svm, acc\_dt]

plt.bar(algorithms, accuracies, color=['blue', 'green', 'orange'])
plt.xlabel('Algorithms')
plt.ylabel('Accuracy')

 $plt.title ('Comparison\ of\ Supervised\ Learning\ Algorithms')\ plt.show()$ 





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## PRACTICAL - 11

❖ <u>AIM</u>: COMPARE THE VARIOUS UNSUPERVISED LEARNING ALGORITHM BY USING THE APPROPRIATE DATASETS.(K MEANS CLUSTERING, K MODE).

## **PROCEDURE**:

#### 1.IMPORT REQUIRED LIBRARIES:

• Load essential libraries such as NumPy, Pandas, Matplotlib, Seaborn, and Scikit-learn.

#### 2.SELECT AN APPROPRIATE DATASET:

- Choose a dataset suitable for clustering tasks, such as:
  - a) K-Means Clustering: Works well with numerical datasets (e.g., Iris Dataset).
  - b) K-Modes Clustering: Used for categorical data (e.g., Mushroom Dataset).

#### 3.PREPROCESS THE DATA:

• Handle missing values, normalize numerical data, and encode categorical variables if necessary.

#### **4.APPLY K-MEANS CLUSTERING:**

- Determine the optimal number of clusters using the Elbow Method or Silhouette Score.
- Train the K-Means model and assign cluster labels to the dataset.

#### **5.APPLY K-MODES CLUSTERING:**

- Convert categorical data into an appropriate format.
- Train the K-Modes model and assign cluster labels.

#### 6.EVALUATE THE CLUSTERING PERFORMANCE:

• Use metrics such as Inertia, Silhouette Score, or visualization techniques (e.g., scatter plots, cluster heatmaps) to analyze results.

#### 7. COMPARE THE ALGORITHMS:

• Compare the clustering results in terms of accuracy, interpretability, and efficiency.

#### 8.DRAW CONCLUSIONS:

• Identify which algorithm performs better based on dataset characteristics and clustering effectiveness.



## **PROGRAM**:

# Install the necessary libraries
!pip install scikit-learn matplotlib kmodes
# Import necessary libraries
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.datasets import load\_iris
from sklearn.cluster import KMeans
from kmodes.kmodes import KModes

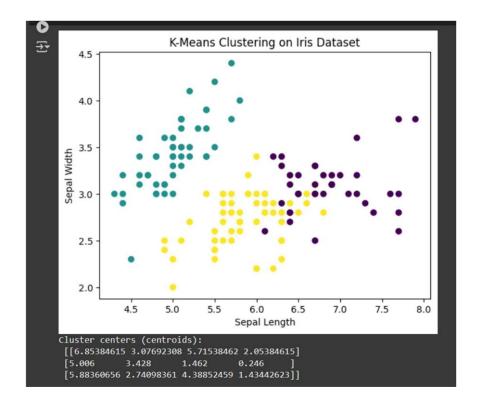
```
Requirement already satisfied: scikit-learn in /usr/local/lib/python3.11/dist-packages (1.6.1)
Requirement already satisfied: matplotlib in /usr/local/lib/python3.11/dist-packages (3.10.0)
Collecting kmodes

Downloading kmodes-0.12.2-py2.py3-none-any.whl.metadata (8.1 kB)
Requirement already satisfied: numpy>=1.19.5 in /usr/local/lib/python3.11/dist-packages (from scikit-learn) (1.26.4)
Requirement already satisfied: scipy>=1.6.0 in /usr/local/lib/python3.11/dist-packages (from scikit-learn) (1.13.1)
Requirement already satisfied: scipy>=1.6.1 in /usr/local/lib/python3.11/dist-packages (from scikit-learn) (1.4.2)
Requirement already satisfied: threadpoolctl>=3.1.0 in /usr/local/lib/python3.11/dist-packages (from scikit-learn) (3.5.0)
Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (1.3.1)
Requirement already satisfied: fonttools>=4.22.0 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (4.56.0)
Requirement already satisfied: packaging>=20.0 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.2.2)
Requirement already satisfied: packaging>=20.0 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.2.2)
Requirement already satisfied: packaging>=20.0 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (1.1.1.0)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (3.2.1)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (3.2.1)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.8.2)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.8.2)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.8.2)
Requirement already satisfied: pyparsing>=23.1 in /usr/local/lib/python3.11/dist-packages (from matplotlib) (2.8.2)
Requirement already sati
```

```
# Load the Iris dataset iris =
load iris()
X = iris.data
# Features: Sepal length, Sepal width, Petal length, Petal width # Apply K-
Means clustering with 3 clusters
kmeans = KMeans(n_clusters=3, random_state=42)
kmeans.fit(X)
# Get the cluster labels
labels = kmeans.labels
# Visualize the K-Means clustering results
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis')
plt.title('K-Means Clustering on Iris Dataset')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
#Print the cluster centers (centroids)
print("Cluster centers (centroids):\n", kmeans.cluster_centers_)
```

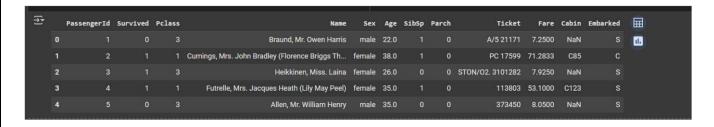


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# Load Titanic dataset from a URL url = "https://raw.githubusercontent.com/datasciencedojo/datasets/master/titanic.csv" data = pd.read\_csv(url)

# Display the first few rows of the dataset to understand its structure data.head()



- # Select categorical columns for clustering categorical\_data = data[['Sex', 'Embarked', 'Pclass']]
- # Encode categorical data as numeric codes categorical\_data = categorical\_data.apply(lambda col: col.astype('category').cat.codes)
- # Apply K-Modes clustering with 3 clusters kmodes = KModes(n\_clusters=3, init='Huang', n\_init=5, verbose=1) clusters = kmodes.fit\_predict(categorical\_data)
- # Add the cluster labels to the original dataset



data['Cluster'] = clusters
# Display the data with the assigned clusters
print(data[['Sex', 'Embarked', 'Pclass', 'Cluster']].head())

# Print the cluster centers (modes)
print("Cluster centers (modes):\n", kmodes.cluster\_centroids\_)

```
Init: initializing centroids

→ Init: initializing clusters

    Starting iterations...
    Run 1, iteration: 1/100, moves: 96, cost: 509.0
    Init: initializing centroids
    Init: initializing clusters
    Starting iterations...
    Run 2, iteration: 1/100, moves: 179, cost: 603.0
    Init: initializing centroids
    Init: initializing clusters
    Starting iterations...
    Run 3, iteration: 1/100, moves: 0, cost: 575.0
    Init: initializing centroids
    Init: initializing clusters
    Starting iterations...
    Run 4, iteration: 1/100, moves: 84, cost: 644.0
    Init: initializing centroids
    Init: initializing clusters
    Starting iterations...
    Run 5, iteration: 1/100, moves: 0, cost: 577.0
    Best run was number 1
         Sex Embarked Pclass Cluster
        male S
    1 female C
2 female S
3 female S
4 male S
    Cluster centers (modes):
     [[1 2 2]
     [0 0 0]
     [0 2 2]]
```



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## PRACTICAL - 12

❖ <u>AIM</u>: BUILD AN ARTIFICIAL NEURAL NETWORK BY IMPLEMENTING THE BACKPROPAGATION ALGORITHM AND TEST THE SAME USING APPROPRIATE DATA SETS.

## **PROCEDURE**:

#### 1. UNDERSTAND THE BACKPROPAGATION ALGORITHM

- Forward propagation: Compute the output using initial weights.
- Compute the loss: Measure the difference between predicted and actual output.
- Backpropagation: Compute gradients using the chain rule.
- Weight update: Adjust weights using gradient descent.

#### 2. IMPORT REQUIRED LIBRARIES

- Use **NumPy** for numerical computations.
- Use **pandas** and **scikit-learn** for dataset handling.

#### 3. DEFINE THE NEURAL NETWORK STRUCTURE

- Input layer: Number of features in the dataset.
- Hidden layer: Choose an appropriate number of neurons.
- Output layer: Number of target classes.
- Activation function: Use sigmoid or ReLU for hidden layers and softmax for multi-class classification.

#### 4. INITIALIZE WEIGHTS AND BIASES

• Use random values for weights and set biases to zero.

#### 5. IMPLEMENT FORWARD PROPAGATION

• Compute weighted sums and apply activation functions layer by layer.

#### 6. COMPUTE THE LOSS FUNCTION

• Use Mean Squared Error (MSE) for regression and Cross-Entropy loss for classification.

#### 7. IMPLEMENT BACKPROPAGATION

- Compute gradients of the loss function with respect to weights and biases.
- Adjust weights using Gradient Descent or Adam Optimizer.

#### 8. TRAIN THE NETWORK

- Feed training data into the model.
- Update weights iteratively over multiple epochs.



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• Monitor loss reduction during training.

#### 9. EVALUATE THE MODEL

- Use a test dataset to check the model's accuracy.
- Compare predicted outputs with actual values.

### 10. OPTIMIZE THE MODEL

• Adjust learning rate, hidden layers, and activation functions for better performance.

#### 11. TEST THE MODEL WITH A REAL DATASET

• Use datasets like Iris, MNIST, or any tabular dataset for classification.

## **PROGRAM**:

```
# Step 1 Import necessary libraries
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make classification
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
# Step 2 Part 1
# Define the Neural Network class with backpropagation
class NeuralNetwork:
  def init (self, input size, hidden size, output size, learning rate=0.01):
     # Initialize network parameters
    self.input size = input size
    self.hidden size = hidden size
    self.output size = output size
    self.learning rate = learning rate
# Step 2 Part 2: Initialize weights and biases with random values
     self.weights input hidden = np.random.randn(self.input size, self.hidden size)
     self.bias hidden = np.random.randn(1, self.hidden size)
    self.weights hidden output = np.random.randn(self.hidden size, self.output size)
    self.bias output = np.random.randn(1, self.output size)
```



```
def sigmoid(self, x):
    return 1/(1 + np.exp(-x))
  def sigmoid derivative(self, x):
    return x * (1 - x)
# Step 2 Part 3
  def forward(self, X):
     # Forward pass: calculate hidden layer output and final output
     self.hidden input = np.dot(X, self.weights input hidden) + self.bias hidden
     self.hidden output = self.sigmoid(self.hidden input)
     self.final input = np.dot(self.hidden output, self.weights hidden output) + self.bias output
     self.final output = self.sigmoid(self.final input)
     return self.final output
# Step 2 Part 4
  def backward(self, X, y):
     # Backpropagation: calculate gradients and update weights
     output error = y - self.final output
     output delta = output error * self.sigmoid derivative(self.final output)
     hidden error = output delta.dot(self.weights hidden output.T)
     hidden delta = hidden error * self.sigmoid derivative(self.hidden output)
# Update weights and biases using the gradients
     self.weights input hidden += X.T.dot(hidden_delta) * self.learning_rate
     self.bias hidden += np.sum(hidden delta, axis=0, keepdims=True) * self.learning rate
     self.weights hidden output += self.hidden output.T.dot(output delta) * self.learning rate
     self.bias output += np.sum(output delta, axis=0, keepdims=True) * self.learning rate
# Step 2 Part 5
  def train(self, X, y, epochs=1000):
# Training the network: run forward pass and backward pass for multiple epochs
     for epoch in range(epochs):
       self.forward(X)
       self.backward(X, y)
```



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```
if epoch \% 100 == 0:
          loss = np.mean(np.square(y - self.final output))
          print(f'Epoch {epoch}, Loss: {loss}')
# Step 2 Part 6
  def predict(self, X):
     # Make predictions with the trained network
     return self.forward(X)
# Step 3
# Generate a synthetic dataset for classification
X, y = make classification(n samples=1000, n features=20, n informative=15, n classes=2)
# Scale the data for better performance in training
scaler = StandardScaler()
X = scaler.fit transform(X)
# Split the data into training and testing sets (80% training, 20% testing)
X train, X test, y train, y test = train test split(X, y.reshape(-1, 1), test size=0.2, random state=42)
# Step 4
# Initialize the neural network with input size, hidden size, and output size
nn = NeuralNetwork(input size=20, hidden size=10, output size=1, learning rate=0.01)
# Train the network with the training data (1000 epochs)
nn.train(X train, y train, epochs=1000)
```

```
Epoch 0, Loss: 0.2502172022855033
Epoch 100, Loss: 0.08991957284799572
Epoch 200, Loss: 0.06760875450455225
Epoch 300, Loss: 0.05107551543490075
Epoch 400, Loss: 0.042791161933662704
Epoch 500, Loss: 0.036625564655806596
Epoch 600, Loss: 0.03232724083004919
Epoch 700, Loss: 0.028755841917128835
Epoch 800, Loss: 0.02591051392948267
Epoch 900, Loss: 0.02342603045676408
```



#step5

```
# Test the trained model on the test set

predictions = nn.predict(X_test)

# Convert predictions to binary (0 or 1)

predictions = (predictions > 0.5).astype(int)

# Calculate accuracy: compare predicted values with actual values accuracy = np.mean(predictions == y_test)

print(f'Accuracy: {accuracy * 100:.2f}%')
```

## → Accuracy: 87.00%

```
#step6
# Neural network class with loss tracking for visualization
class NeuralNetworkWithLoss(NeuralNetwork):
  def init (self, input size, hidden size, output size, learning rate=0.01):
     super(). init (input size, hidden size, output size, learning rate)
    self.loss history = []
#step6 part2
def train(self, X, y, epochs=1000):
     for epoch in range(epochs):
       self.forward(X)
       self.backward(X, y)
       if epoch \% 100 == 0:
          loss = np.mean(np.square(y - self.final output))
         self.loss history.append(loss)
         print(f'Epoch {epoch}, Loss: {loss}')
#step6 prac3
# Re-train using the modified class to store the loss
nn with loss = NeuralNetworkWithLoss(input size=20, hidden size=10, output size=1,
learning rate=0.01)
nn with loss.train(X train, y train, epochs=1000)
```



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```
Epoch 0, Loss: 0.46936617477890524
Epoch 100, Loss: 0.12411160196289518
Epoch 200, Loss: 0.09914075921479235
Epoch 300, Loss: 0.07783312799646006
Epoch 400, Loss: 0.0595244840988669
Epoch 500, Loss: 0.046070768146957236
Epoch 600, Loss: 0.036989619528565096
Epoch 700, Loss: 0.03136187735015099
Epoch 800, Loss: 0.027356785929926294
Epoch 900, Loss: 0.024565783779983114
```

```
# Plot loss over epochs

plt.plot(nn_with_loss.loss_history)

plt.title('Loss over Epochs')

plt.xlabel('Epochs')

plt.ylabel('Loss')

plt.show()
```

