```
PR1 DW1 Perform the following operations using Python on any opensource dataset
import pandas as pd
import numpy as np
~data={'NAME':['jay','prince','nisha','neha','raj','riya'],
   'Age':[17,20,21,18,19,25],
   'Gender':['M','M','F','F','M','M'],
   'Marks':[90,75,80,84,71,86] }
~df = pd.DataFrame(data)
~print("Original Data:")
 print(df)
~data1 = {
  'NAME': ['jay', 'prince', 'nisha', 'neha', 'raj', 'riya'],
  'Age': [17, 20, 21, 18, 19, 25],
  'Gender': ['M', 'M', 'F', 'F', 'M', 'M'],
  'Marks': [90, 75, np.nan, 84, np.nan, 86] # Proper missing values }
~df2 = pd.DataFrame(data1)
 print("\nData with Missing Marks:")
 print(df2)
~df['Gender'] = df['Gender'].map({'M': 0, 'F': 1}).astype(float)
 df = df[df['Marks'] > 75]
 df = df.drop(['Age'], axis=1)
 print("\nCleaned Data (Marks > 75, Gender M/F mapped to 0/1):")
 print(df)
or
df = pd.read csv("iris.csv")
print(df.head())
print("\nMissing Values in Each Column:")
print(df.isnull().sum())
print("\nData Description:")
print(df.describe())
df['species'] = df['species'].map({'setosa': 0, 'versicolor': 1, 'virginica': 2})
df = df[df['petal length'] > 1.5]
df = df.drop(['sepal_width'], axis=1)
print("\nCleaned Dataset:")
print(df.head())
```

```
PR2 DW2 Create an "Academic performance" dataset of students
~import pandas as pd
 import numpy as np
~data_frame=pd.read_csv("Academic_performace.csv")
 print(data_frame)
~print(data_frame.head())
~print(data_frame.tail())
~print(data_frame.describe())
~print(data_frame.info())
~print(data_frame.shape)
~print(data_frame.isnull().any().any())
~print(data_frame.isnull().sum())
~import seaborn as sns
import matplotlib.pyplot as plt
from scipy import stats
x=data_frame['Sno']
y=data_frame['AnnouncementsView']
sns.regplot(x='Sno', y='AnnouncementsView',data=data_frame)
sns.boxplot(x=data_frame['AnnouncementsView'])
z=np.abs(stats.zscore(data_frame['AnnouncementsView']))
print(z)
threshold=3
print(np.where(z>3))
~df=pd.DataFrame({
  'Income':[15000,1000,120000,10000],
  'Age':[25,18,42,51],
  'Department':['HR','Legal','Marketing','Management']})
print(df)
df scaled=df.copy()
```

col names=['Income','Age']

features=df scaled[col names]

```
PR3 Descriptive Statistics - Measures of Central Tendency and variability
~import pandas as pd
import numpy as np
import statistics as st
~data_frame = pd.read_csv("/home/ubuntu/DSBDA/test_AV3.csv")
~print(data_frame)
~print(data_frame.info())
~print(data_frame.mean)
~print(data_frame['ApplicantIncome'].mean())
~print(data_frame['ApplicantIncome'].median())
~print(data_frame.mode())
~print(data_frame['ApplicantIncome'].mode())
~print(data_frame.median)
~print(data_frame['ApplicantIncome'].std())
~print(data_frame['LoanAmount'].std())
~print(data_frame['LoanAmount'].var())
~data_frame = pd.read_csv("iris.csv")
~print(data_frame)
~print(data_frame.info())
~setosa = data_frame['species'] == 'setosa'
print(data_frame[setosa].describe())
~virginica = data_frame['species'] == 'virginica'
 print(data_frame[virginica].describe())
~versicolor = data_frame['species'] == 'versicolor'
 print(data frame[versicolor].describe())
~print('\nIris-versicolor')
 ver = data frame['species'] == 'versicolor'
 print(data frame[ver].describe())
~print('\nIris-virginica')
 virg = data frame['species'] == 'virginica'
 print(data_frame[virg].describe())
grouped = data_frame.groupby('species')
for name, group in grouped:
  print(group.select dtypes(include=np.number).values.tolist())
```

```
PR4 Linear RegressionModel
import pandas as pd
import numpy as np
from sklearn.linear_model import LinearRegression
from sklearn.model_selection import train_test_split
from sklearn.metrics import mean_squared_error
df = pd.read_csv("Boston.csv")
print(df.head())
print(df.describe())
X = df.drop("MEDV", axis=1)
y = df["MEDV"]
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.33,
random state=42)
model = LinearRegression()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print("Model Coefficients:", model.coef_)
print("\nFirst 10 Predicted vs Actual Prices:")
for i in range(10):
  print(f"Predicted: {y pred[i]}, Actual: {y test.iloc[i]}")
mse = mean squared error(y test, y pred)
rmse = np.sqrt(mse)
print(f"\nMean Squared Error (MSE): {mse}")
print(f"Root Mean Squared Error (RMSE): {rmse}")
```

```
PR5 logistic regression
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import (
  confusion_matrix,
  accuracy_score,
  precision_score,
  recall_score,
  f1_score
dataset = pd.read_csv('Social_Network_Ads.csv')
print(dataset.head())
print(dataset.tail())
X = dataset[['Age', 'EstimatedSalary']]
Y = dataset['Purchased']
X train, X test, Y train, Y test = train test split(
  X, Y, test size=0.25, random state=0
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X test = scaler.transform(X test)
classifier = LogisticRegression(random state=0)
classifier.fit(X train, Y train)
Y pred = classifier.predict(X test)
cm = confusion matrix(Y test, Y pred)
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tp = cm[1, 1]
fp = cm[0, 1]
tn = cm[0, 0]
fn = cm[1, 0]
accuracy = accuracy score(Y test, Y pred)
error rate = 1 - accuracy
precision = precision score(Y test, Y pred)
recall = recall_score(Y_test, Y_pred)
f1 = f1_score(Y_test, Y_pred)
print("Confusion Matrix:")
print(cm)
print(f"True Positive: {tp}")
print(f"False Positive: {fp}")
print(f"True Negative: {tn}")
print(f"False Negative: {fn}")
print(f"Accuracy: {accuracy:.4f}")
print(f"Error Rate: {error_rate:.4f}")
print(f"Precision: {precision:.4f}")
print(f"Recall: {recall:.4f}")
print(f"F1 Score: {f1:.4f}")
plt.figure(figsize=(6, 6))
sns.heatmap(
  cm,
  annot=True,
  fmt="d",
  cmap="Blues",
  xticklabels=["Predict No", "Predict Yes"],
  yticklabels=["Actual No", "Actual Yes"]
plt.title("Confusion Matrix")
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.show()
```

```
Naïve Bayes classificationalgorithm
PR6
import pandas as pd
import numpy as np
from sklearn.model selection import train test split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import confusion_matrix, accuracy_score, precision_score,
recall_score,f1_score
data = pd.read csv('iris.csv')
X = data.drop('species', axis=1)
y = data['species']
print("Value of X=",X)
print( "Value of Y=",y)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
print("X train shape:", X train.shape)
print("y train shape:", y train.shape)
model= GaussianNB()
model.fit(X train, y train)
y pred = model.predict(X test)
cm = confusion_matrix(y_test, y_pred)
print("Confusion Matrix:\n", cm)
TP = cm[0, 0] # True positives (class 0)
FP = cm[0, 1] # False positives (class 1)
TN = cm[1, 1] # True negatives (class 1)
FN = cm[1, 0]
accuracy = accuracy score(y test, y pred)
error rate = 1 - accuracy
precision = precision_score(y_test, y_pred, average='macro')
recall = recall_score(y_test, y_pred, average='macro')
f1 = f1 score(y test, y pred, average='macro')
print(f"Accuracy: {accuracy:.4f}")
print(f"Error Rate: {error rate:.4f}")
print(f"Precision: {precision:.4f}")
print(f"Recall: {recall:.4f}")
print(f"F1 Score: {f1:.4f}")
```

```
PR7 Tokenization ,POSTagging, stopwords removal
import nltk
from nltk.tokenize import word tokenize
from nltk.corpus import stopwords
from nltk.stem import PorterStemmer, WordNetLemmatizer
from nltk import pos_tag
from sklearn.feature extraction.text import TfidfVectorizer
import string
~nltk.download('punkt tab')
 nltk.download('stopwords')
 nltk.download('wordnet')
nltk.download('averaged_perceptron_tagger_eng')
document =""" Natural Langauage processing is a field of AI It aims to enable to
understand, interpret and generate human langaueg """
tokens= word tokenize(document)
pos tags = pos tag(tokens)
stop words = set(stopwords.words('english'))
filtered tokens=[word for word in tokens if word.lower() not in stop words and word
not in
string.punctuation]
stemmer =PorterStemmer()
stemmed_tokens=[stemmer.stem(word) for word in filtered_tokens]
lemmatizer = WordNetLemmatizer()
lemmatized_tokens =[lemmatizer.lemmatize(word)for word in filtered_tokens]
print("\n original doc\n",document)
print("\n tokens\n", tokens)
print("\n pos tag\n",pos_tag)
print("\n filtered token after removed stop word removal\n", filtered tokens)
print("\n stemmed tokens\n",stemmed tokens)
print("\n Lemmetaized Tokens\n",lemmatized tokens)
~corpus = [document]
 vectorizer = TfidfVectorizer()
 x = vectorizer.fit transform(corpus)
 tfidf matrix = x.toarray()
 terms = vectorizer.get_feature_names_out()
~print("\nTerm Frequency and Inverse Document Frequency (TF-IDF):")
for i, term in enumerate(terms):
    print(f"Term: {term}, TF-IDF: {tfidf matrix[0][i]:.4f}")
```

```
PR8 DV1 titanic 891rows
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
titanic = sns.load_dataset('titanic')
print(titanic.head())
plt.title('Distribution of Fare Prices of Titanic Passengers')
plt.xlabel('Fare')
plt.ylabel('Density')
plt.show()
plt.figure(figsize=(10, 6))
sns.histplot(titanic['fare'], bins=30, kde=True, color='blue', stat='density')
plt.title('Distribution of Fare Prices of Titanic Passengers')
plt.xlabel('Fare')
plt.ylabel('Density')
plt.grid(True)
plt.show()
PR9 DV2 titanic Plotaboxplot
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
titanic = sns.load dataset('titanic')
titanic.head(10)
titanic.info()
titanic.describe()
titanic.loc[:,["survived","alive"]]
sns.boxplot(x='sex', y='age', data=titanic)
plt.show()
sns.boxplot(x='sex', y='age', hue='survived', data=titanic, palette="Set2")
```

## PR10 Data Visualization III:- Download the Iris flower dataset import pandas as pd import seaborn as sns import matplotlib.pyplot as plt df = pd.read\_csv('iris.csv') print("Features And Types in the iris dataset") print (df.dtypes) df.drop(columns='Species').hist(bins=20, figsize=(10,8), color='skyblue',edgecolor='black') plt.suptitle('Histogram of Each Feature',fontsize=16) plt.tight\_layout() plt.show() plt.figure(figsize=(10,8)) sns.boxplot(x='Species',y='Spetal\_length',data=df) sns.boxplot(x='Species',y='Spetal\_length',data=df) sns.boxplot(x='Species',y='Spetal\_length',data=df) sns.boxplot(x='Species',y='Spetal\_length',data=df) plt.tight\_layout() plt.show() sns.pairplot(df,hue='Species')

plt.suptitle('pairplot of iris dataset',fontsize=16)

plt.tight layout()

plt.show()

```
PR13 Scala
//checks if a number is positive, negative, or zero
object Number{
def main (args: Array[String]) {
var number = (-100);
if(number==0) {
println("number is zero");
                                      }
else if(number>0) {
println("number is positive");
else { println("number is negative");
                                                 }}}
//Print name
object Name {
def main (args: Array[String]) {
println("my name is name");
} }
// concanate string
object String {
def main (args: Array[String]) {
var text: String = "Scala in programming langauge";
println("string is "+text);
}}
// compares two numbers
object Find Number{
def main (args: Array[String]) {
var number1 = 20;
var number 2 = 40;
if(number1>number2) {
println("large number is "+number1); }
else
{ println("large number is "+number2);
                                           } } }
// addition of no
object Addition {
 def main(args: Array[String]) {
  val num1 = 10
  val num2 = 20
  val sum = num1 + num2
  println("The sum is: " + sum) }
```