

DEPARTMENT OF COMPUTER APPLICATIONS

CADX116 - MACHINE LEARNING ALGORITHMS LABORATORY



LAB RECORD

NAME	:	
RRN	:	
LAB	:	CADX116 - MACHINE LEARNING ALGORITHMS LABORATORY



DEPARTMENT OF COMPUTER APPLICATIONS

ACADEMIC YEAR (JULY 2024 - DECEMBER 2024)

COURSE CODE : CADX116

COURSE NAME : MACHINE LEARNING ALGORITHMS

LABORATORY

PROGRAMME : BCA (DATA SCIENCE)

SEMESTER : V



BONAFIDE CERTIFICATE

This is a Certified Record Book of	
RRN:	submitted for the Semester End
Practical Examination held on	, for the <u>CADX116</u> -
MACHINE LEARNING ALGORITHMS	<u>LABORATORY</u> during <u>2024 - 2025</u> .
	Signature of Faculty

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11.		Implement Logistic Regression		
12.		Implement Neural Network Model		

K.NO : ATE : IMPLEMENT NAÏVE BAYES CLASSIFIER								
URCE C	ODE :							
Step 1:								
	[] import pandas as pd from sklearn import tree from sklearn.preprocessing import LabelEncoder from sklearn.naive_bayes import GaussianNB							
Step 2:			\wedge	\forall		1		
[] data=pd.read_csv("Naive_Bayes_PlayTennis.csv") print("the First 5 values of data is :\n") data.head()								
Output: the First 5 values of data is:								
		Outlook	Temperature	Humidity	Windy	Play Tennis		
O Sunny Hot High False No								
1 Sunny Hot High True No								
2 Overcast Hot High False Yes								
3 Rainy Mild High False Yes								
4 Rainy Cool Normal False Yes								

 $print("the\ First\ 5\ values\ of\ data\ is\ :\n")$

X.head()

Output:

the First 5 values of data is:

	Outlook	Temperature	Humidity	Windy
0	Sunny	Hot	High	False
1	Sunny	Hot	High	True
2	Overcast	Hot	High	False
3	Rainy	Mild	High	False
4	Rainy	Cool	Normal	False

Step 4:

```
[] y=data.iloc[:,-1]

print("the First 5 values of data is :\n")

y.head()
```

Output:

the First 5 values of data is:

- 0 No
- 1 No
- 2 Yes
- 3 Yes
- 4 Yes

Name: PlayTennis, dtype: object

Step 5:

```
[] for column in X.columns:

X[column]=LabelEncoder().fit_transform(X[column])

X.head()
```

	Outlook	Temperature	Humidity	Windy
0	2	1	0	0
1	2	1	0	1
2	0	1	0	0
3	1	2	0	0
4	1	0	1	0

	y=LabelEncoder().fit_transform(y)
	print(y)
Output:	
	[0 0 1 1 1 0 1 0 1 1 1 1 1 0]
Step 7:	
	from sklearn.model_selection import train_test_split
Step 8:	
	X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.20)
Step 9:	
	from sklearn.naive_bayes import GaussianNB
	classifier = GaussianNB()
	classifier.fit(X_train,y_train)
Output:	
	▼ GaussianNB
	GaussianNB()
Step 10:	
	from sklearn.metrics import accuracy_score
Step 11:	
	<pre>print("Accuracy is:",accuracy_score(classifier.predict(X_test),(y_test)))</pre>
Output:	
	Accuracy is: 0.666666666666666666666666666666666666

EX.NO : DATE :	
	IMPLEMENT LINEAR REGRESSION
SOURCE	CODE:
Step 1	:
	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.linear_model import LinearRegression from sklearn.metrics import mean_squared_error, r2_score
Step 2	
	data=pd.read_csv("Linear_Regression_Advertising.csv") print("the First 5 values of data is :\n") data.head()

the First 5 values of data is ?

	TV	Radio	Newspaper	Sales
0	230.1	37.8	69.2	22.1
1	44.5	39.3	45.1	10.4
2	17.2	45.9	69.3	9.3
3	151.5	41.3	58.5	18.5
4	180.8	10.8	58.4	12.9

Step 3:

```
[] X = data[['TV', 'Radio', 'Newspaper']]

y = data['Sales']

X.head()
```

Output:

	TV	Radio	Newspaper
0	230.1	37.8	69.2
1	44.5	39.3	45.1
2	17.2	45.9	69.3
3	151.5	41.3	58.5
4	180.8	10.8	58.4

Step 4:

[] y.head()

Output:

0 22.1

1 10.4

2 9.3

3 18.5

4 12.9

Name: Sales, dtype: float64

Step 5:

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

model = LinearRegression()

model.fit(X_train, y_train)

Output:

LinearRegression
LinearRegression()

Step 6:

 $[] y_pred = model.predict(X_test)$

Step 7:

```
[] mse = mean_squared_error(y_test, y_pred)

r2 = r2_score(y_test, y_pred)

print(f'Mean Squared Error: {mse}')

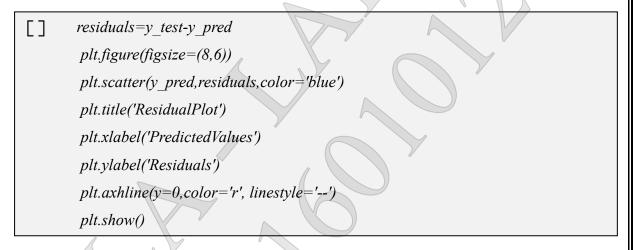
print(f'R^2 Score: {r2}')
```

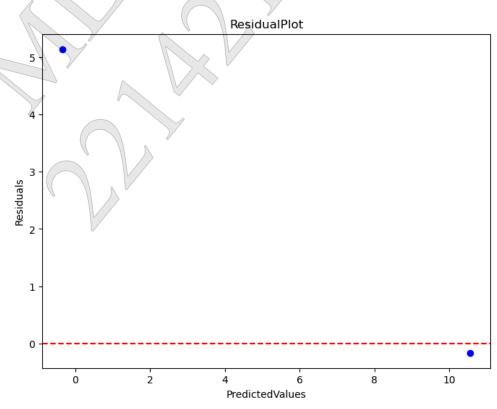
Output:

Mean Squared Error: 13.201451790963432

R^2 Score: -0.6838586468065599

Step 8:





Step 9:

```
[] plt.figure(figsize=(8,6))

plt.scatter(y_test,y_pred,color='green')

plt.title('PredictedvsActual')

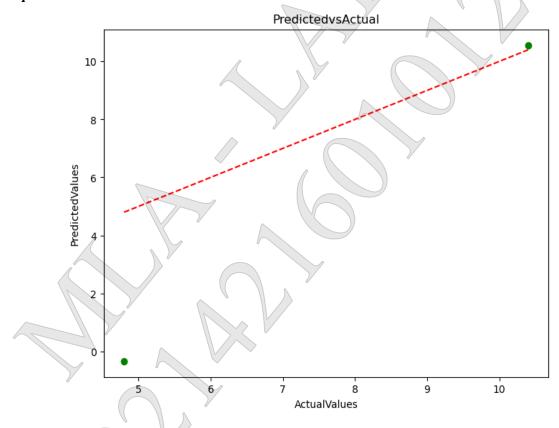
plt.xlabel('ActualValues')

plt.ylabel('PredictedValues')

plt.plot([min(y_test),max(y_test)], [min(y_test),max(y_test)],color='red',

linestyle='--')

plt.show()
```



DATE : IMPLEMENT SUPPORT VECTOR MACHINE (SVM) ALGORITHM		
SOURCE CODE: Step 1: [] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC	EX.NO:	
SOURCE CODE: Step 1: [] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC	DATE :	
SOURCE CODE: Step 1: [] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		IMPLEMENT SUPPORT VECTOR MACHINE
Source code: Step 1: [] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		
[] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		(SVM) ALGORITHM
[] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		
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[] import pandas as pd import numpy as np from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC	SOURCE COL	DE:
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from sklearn.model_selection import train_test_split,cross_val_score from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC	[]	import pandas as pd
from sklearn.preprocessing import StandardScaler, LabelEncoder from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		import numpy as np
from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, fl_score from sklearn.svm import SVC		from sklearn.model_selection import train_test_split,cross_val_score
recall_score, f1_score from sklearn.svm import SVC		from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.svm import SVC		from sklearn.metrics import confusion_matrix, accuracy_score, precision_score,
		recall_score, f1_score
		from sklearn.svm import SVC
Step 2:		
Step 2:		
	Step 2:	

[] data=pd.read_csv("SVM_PurchasePrediction.csv")

print("the First 5 values of data is :\n")

data.head()

Output:

the First 5 values of data is:

	Age	Income	Gender	Purchase
0	25	50000	Male	0
1	45	64000	Female	1
2	35	58000	Female	0
3	50	72000	Male	1
4	23	48000	Male	0

```
Step 3:
     le=LabelEncoder()
             data['Gender']=le.fit transform(data['Gender'])
Step 4:
             data.fillna(method='ffill',inplace=True)
Step 5:
             X = data.drop('Purchase',axis=1)
     y = data['Purchase']
Step 6:
             sc=StandardScaler()
      X \ scaled = sc.fit \ transform(X)
Step 7:
     classifiers={
               'Linear SVM': SVC(kernel='linear',random state=0),
               'Polynomial SVM': SVC(kernel='poly',degree=3,random state=0),
               'RBF SVM': SVC(kernel='rbf',random state=0)
Step 8:
           for clf name, clf in classifiers.items():
               scores=cross val score(clf,X scaled,y,cv=5,scoring='accuracy')
               print(f"{clf name} Cross-validation Accuracy: {scores.mean():.2f}(+/-
             {scores.std()*2:.2f})")
Output:
         Linear SVM Cross-validation Accuracy : 0.80(+/-0.33)
         Polynomial SVM Cross-validation Accuracy: 0.80(+/-0.53)
         RBF SVM Cross-validation Accuracy: 0.80(+/-0.53)
Step 9:
              X_{train}, X_{test}, y_{train}, y_{test} = train_{test_{split}}(X, y, test_{size}=0.25, random_{state}=0)
     Г٦
              classifier=SVC(kernel='linear',random state=0)
              classifier.fit(X train, y train)
```

```
Output:
                               SVC
            SVC(kernel='linear', random_state=0)
Step 10:
     y_pred=classifier.predict(X_test)
Step 11:
            cm=confusion_matrix(y_test,y_pred)
      accuracy=accuracy_score(y_test,y_pred)
            precision=precision_score(y_test,y_pred)
            recall=recall_score(y_test,y_pred)
            fl=fl_score(y_test, y_pred)
Step 12:
     print(f"Confusion Matrix :\n",cm)
            print(f"Accuracy:{accuracy:.2f}")
            print(f"Precision : {precision:.2f}")
            print(f"Recall :{recall:.2f}")
            print(f"F1 Performance score :{f1:.2f}")
Output:
              Confusion Matrix.
               [[4]]
              Accuracy:1.00
              Precision:1.00
              Recall:1.00
              F1 Performance score :1.00
```

EX.NO:	
DATE:	
	IMPLEMENT DECISION TREE CLASSIFIER
SOURCE COL	DE:
Step 1:	
	import pandas as pd import numpy as np from sklearn.model_selection import train_test_split from sklearn.tree import DecisionTreeClassifier from sklearn.metrics import confusion_matrix,accuracy_score,precision_score,recall_score,fl_score
Step 2:	
Output:	data=pd.read_csv("DecisionTree_TargetIncome.csv") print("the First 5 values of data is :\n") data.head() First 5 values of data is :
	Age Income Gender TargetVariable

	Age	Income	Gender	TargetVariable
0	25	50000	Male	0
1	45	64000	Female	1
2	35	58000	Female	0
3	50	72000	Male	1
4	23	48000	Male	0

Step 3:

data=pd.get_dummies(data, columns=['Gender'],drop_first=True)

```
Step 4:
      print(data.columns)
Step 5:
      X = data.drop('TargetVariable',axis=1)
             y = data['TargetVariable']
Step 6:
              X_{train}, X_{test}, y_{train}, y_{test} = train_{test} split(X, y, test_{size} = 0.25, random_{state} = 0)
     classifier=DecisionTreeClassifier(random state=0)
              classifier.fit(X train, y train)
Output:
                 DecisionTreeClassifier
        DecisionTreeClassifier(random_state=0)
Step 7:
      y_pred=classifier.predict(X_test)
Step 8:
             cm=confusion_matrix(y_test,y_pred)
      accuracy=accuracy_score(y_test,y_pred)
             precision=precision_score(y_test,y_pred)
             recall=recall score(y test,y pred)
             fl=fl_score(y_test,y_pred)
Step 9:
      print(f"Confusion Matrix :\n",cm)
             print(f"Accuracy : {accuracy:.2f}")
             print(f"Precision : {precision:.2f}")
             print(f"Recall :{recall:.2f}")
             print(f"F1 score :{f1:.2f}")
```

Output: Confusion Matrix: [[0 0] [1 3]] Accuracy:0.75 Precision:1.00 Recall:0.75 F1 score:0.86

EX.NO : DATE :	
DAIE .	IMPLEMENT RANDOM FOREST ALGORITHM
SOURCE COL	DE:
Step 1:	
	import pandas as pd from sklearn.model_selection import train_test_split from sklearn.ensemble import RandomForestClassifier from sklearn.metrics import accuracy_score, classification_report import warnings warnings.filterwarnings('ignore')
Step 2:	
	titanic_data=pd.read_csv('RandomForest_TitanicSurvival.csv') titanic_data.head()
Output:	

	Passenger Id	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin	Embarked
0	1	<i>y</i> 0	3	Braund, Mr. Owen Harris	male	22	1	0	A/5 21171	7.25	-	S
1	2		1	Cumings, Mrs. John Bradley (Florence Briggs Thayer)	female	38	1	0	PC 17599	71.2833	C85	С
2	3	1	3	Heikkinen, Miss. Laina	female	26	0	0	STON/O2. 3101282	7.925	-	S
3	4	1	1	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35	1	0	113803	53.1	C123	S
4	5	0	3	Allen, Mr. William Henry	male	35	0	0	373450	8.05	-	S

	titanic_data=titanic_data.dropna(subset=['Survived'])
Step 4:	
[]	X=titanic_data[['Pclass','Sex','Age','SibSp','Parch','Fare']]
	y=titanic_data['Survived']
Step 5:	
	$X.loc[:,'Sex'] = X['Sex'].map(\{'female':0,'male':1\})$ X.loc[:,'Age'] = X['Age'].fillna(X['Age'].median())
Step 6:	
	X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=
	$rf_classifier = RandomForestClassifier (n_estimators = 100, random_state = 42)$
	rf_classifier.fit(X_train, y_train)
Output:	
Output:	RandomForestClassifier
•	RandomForestClassifier ndomForestClassifier(random_state=42)
•	
•	
Rail Step 7:	
Rai	ndomForestClassifier(random_state=42)
Rail Step 7:	ndomForestClassifier(random_state=42)
Ral Step 7:	ndomForestClassifier(random_state=42)
Rail Step 7: [] Step 8:	y_pred=rf_classifier.predict(X_test)
Rail Step 7: [] Step 8:	y_pred=rf_classifier.predict(X_test) accuracy=accuracy_score(y_test,y_pred)
Rail Step 7: [] Step 8:	y_pred=rf_classifier.predict(X_test) accuracy=accuracy_score(y_test,y_pred)
Ral Step 7: [] Step 8:	y_pred=rf_classifier.predict(X_test) accuracy=accuracy_score(y_test,y_pred)

Output	
Ouipui	٠

Accurancy:0.80

Classification Report :

J	precision	recall	f1-score	support
0	0.82	0.85	0.83	105
1	0.77	0.73	0.75	74
accuracy			0.80	179
macro avg	0.79	0.79	0.79	179
weighted avg	0.80	0.80	0.80	179

EX.NO: DATE:	
	IMPLEMENT K-MEANS CLUSTERING ALGORITHM
SOURCE CO.	DE:
Step 1:	
	import pandas as pd import numpy as np from sklearn.cluster import KMeans import matplotlib.pyplot as plt
Step 2:	
	data=pd.read_csv("KMeans_FeatureClusters.csv") print("the First 5 values of data is :\n") data.head()
Output:	

the First 5 values of data is:

	Feature1	Feature2
0	2.3	3.4
1	1.5	1.8
2	7.6	6.5
3	2.1	4.2
4	8.0	7.0

Step 3:

[] kmeans=KMeans(n_clusters=2,random_state=0)
kmeans.fit(df)

Output:

C:\ProgramData\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:870:

FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

warnings.warn(

C:\ProgramData\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:1382:

UserWarning: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks than available threads. You can avoid it by setting the environment variable OMP NUM THREADS=1.

warnings.warn(

KMeans

KMeans(n_clusters=2, random_state=0)

Step 4:

[] df['cluster']=kmeans.labels_

Step 5:

[] print(df.head())

	Feature1	Feature2	Cluster
0	2.3	3.4	1
1	1.5	1.8	1
2	7.6	6.5	0
3	2.1	4.2	1
4	8.0	7.0	0

Step 6:

```
[] plt.figure(figsize=(10,8))

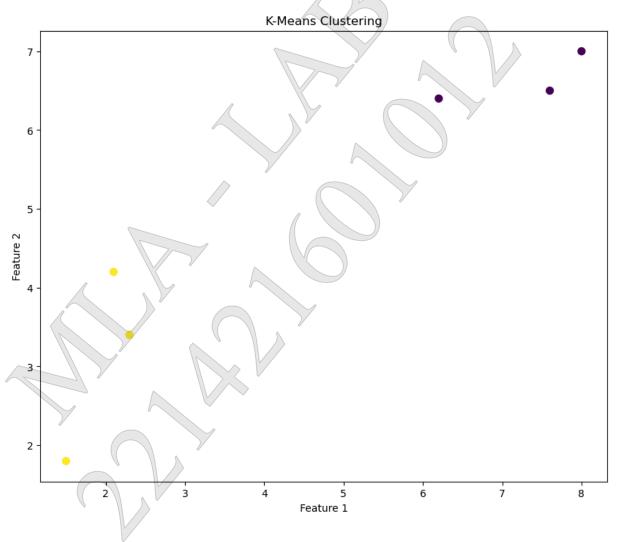
plt.scatter(df['Feature1'],df['Feature2'],c=df['cluster'],cmap='viridis',s=50)

plt.title('K-Means Clustering')

plt.xlabel('Feature 1')

plt.ylabel('Feature 2')

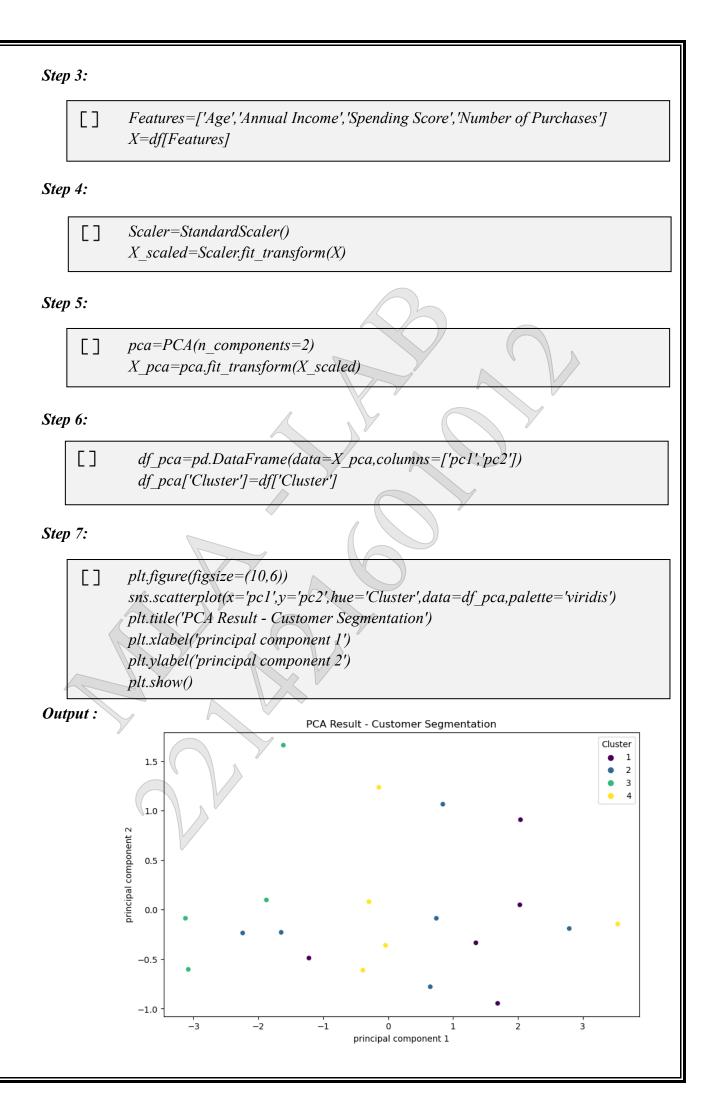
plt.show()
```



EX.NO: DATE:		
	IMP	PLEMENT PRINCIPAL COMPONENT ANALYSIS (PCA)
SOURCE	COL	DE:
Step	1:	
		import pandas as pd import numpy as np from sklearn.decomposition import PCA
		from sklearn.preprocessing import StandardScaler import matplotlib.pyplot as plt import seaborn as sns
Step	2:	
		data=pd.read_csv("PCA_CustomerSegmentation.csv") print("the First 5 values of data is :\n") data.head()
Out <u>f</u>	out:	

the First 5 values of data is:

	Age	Annual Income	Spending Score	Number of Purchases	Cluster
0	22	35000	45	15	3
1	28	45000	50	18	2
2	33	60000	55	22	1
3	38	70000	60	25	4
4	45	80000	65	30	2



Step 8:

[] explained_variance=pca.explained_variance_ratio_
print(f'Explained Variance by each Component: {explained_variance}')
print(f'Total Explained Variance : {np.sum(explained_variance)}')

Output:

Explained Variance by each Component: [0.87276806 0.1144451]

Total Explained Variance: 0.9872131621734076



EX.NO:										
DATE:										
IMPLEMENT K-NEAREST NEIGHBORS										
(KNN) ALGORITHM										
SOUDCE COL	$n_{m E}$.									
SOUNCE COL	SOURCE CODE:									
Step 1:										
	import pandas as pd									
	import plotly.graph_objects as go									
	import plotly.offline as pyoff									
	from sklearn.model_selection import train_test_split									
	from sklearn.preprocessing import StandardScaler									
	from sklearn.neighbors import KNeighborsClassifier									
	from sklearn.metrics import classification_report,confusion_matrix,accuracy_score									
Step 2:										
	data=pd.read_csv("KNN_CellClassification.csv")									
	data.info()									
Output:										
	<class 'pandas.core.frame.dataframe'=""></class>									
	RangeIndex: 698 entries, 0 to 697 Data columns (total 11 columns):									
	# Column Non-Null Count Dtype									
	0 1000025 698 non-null int64									
	0									
	2 1 698 non-null int64									
	3									
	5 2 698 non-null int64									
	6 1.3 698 non-null object 7 3 698 non-null int64									
	8 1.4 698 non-null int64									
	9 1.5 698 non-null int64									
	10 2.1 698 non-null int64 dtypes: int64(10), object(1)									
	memory usage: 60.1+ KB									

Step 3: [] data.head()

Output:

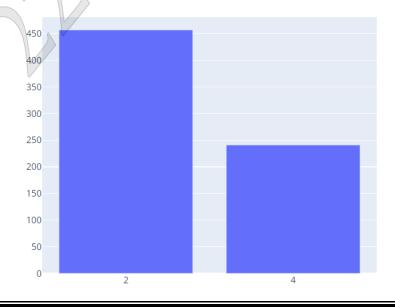
	1000025	5	1	1.1	1.2	2	1.3	3	1.4	1.5	2.1
0	1002945	5	4	4	5	7	10	3	2	1	2
1	1015425	3	1	1	1	2	2	3	1	1	2
2	1016277	6	8	8	1	3	4	3	7	1	2
3	1017023	4	1	1	3	2	1	3) 1	1	2
4	1017122	8	10	10	8	7	10	9	7	1	4

Step 4:

[] data.columns=['Id','Clump Thickness','uniformity of cell size','uniformity of cell shape','Marginal Adhesion','Single Epithelial cell size','Bare Nuclei','Bland Chromatin','Normal Nucleoli','Mitoses','Class']

Step 5:

```
[] target_balance=data['Class'].value_counts().reset_index()
    target_balance.columns=['Class','Count']
    target_class=go.Bar(
        name="Target Balance",
        x=target_balance['Class'].astype(str),
        y=target_balance['Count']
    )
    fig=go.Figure(target_class)
    pyoff.iplot(fig)
```



Step 6:

```
[] beg_class_pat=data.loc[data['Class']==2]

mal_class_pat=data.loc[data['Class']==4]

Mith_10_beg=beg_class_pat['Mitoses'].value_counts().reset_index()

Mith_10_beg.columns=['Mitoses','Count']

Mith_10_mal=mal_class_pat['Mitoses'].value_counts().reset_index()

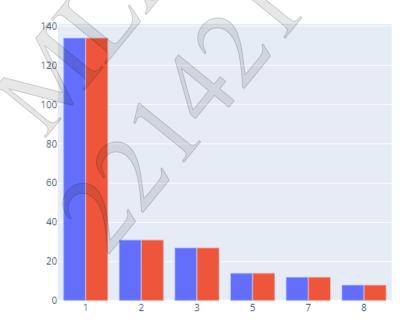
Mith_10_mal.columns=['Mitoses','Count']
```

Step 7:

```
[] fig=go.Figure(data=[
go.Bar(name='Levels of Mitoses in Benign
Group',x=Mith_10_beg['Mitoses'].astype(str),y=Mith_10_mal['Count']),
go.Bar(name='Levels of Mitoses in Malignant
Group',x=Mith_10_beg['Mitoses'].astype(str),y=Mith_10_mal['Count'])])
fig.update_layout(barmode='group')
fig.show()
```

Levels of Mitoses in Benign Group Levels of Mitoses in Malignant Group

Output:



Step 8:

```
[] x=data.drop(columns=['Id','Class'])
y=data['Class']
```

```
Step 9:
     print("Unique Values in 'Bara Nuclei':",x['Bare Nuclei'].unique())
Output:
              Unique Values in 'Bara Nuclei': ['10' '2' '4' '1' '3' '9' '7' '?' '5' '8' '6']
Step 10:
     x['Bare Nuclei']=pd.to_numeric(x['Bare Nuclei'],errors='coerce')
Step 11:
     x=x.fillna(x.median())
Step 12:
     x train,x test,y train,y test=train test split(x,y,test size=0.3,random state=42)
Step 13:
            scaler=StandardScaler()
      x train scaled=scaler.fit transform(x train)
            x test scaled=scaler.transform(x test)
Step 14:
     knn=KNeighborsClassifier(n neighbors=5)
            knn.fit(x_train_scaled,y_train)
Output:
            ▼ KNeighborsClassifier
            KNeighborsClassifier()
Step 15:
            y pred=knn.predict(x test scaled)
```

Step 16:

[] print("Accuracy: ",accuracy_score(y_test,y_pred))
print()
print("Confusion Matrix: \n",confusion_matrix(y_test,y_pred))
print("Classification Report: \n",classification_report(y_test,y_pred))

Output:

Accuracy: 0.9714285714285714

Confusion Matrix:

[[131 4]

[2 73]]

Classification Report:

	precision	recall	fl-score	suppor
			\	
2	0.98	0.97	0.98	135
4	0.95	0.97	0.96	75
		(7
accuracy			0.97	210
macro avg	0.97	0.97	0.97	210
weighted avg	0.97	0.97	0.97	210

EX.NO:	
DATE:	
	DEMONSTRATE WEKA TOOL
SOURCE COL	DE:
Step 1:	
	import pandas as pd from sklearn.model_selection import train_test_split from sklearn.preprocessing import StandardScaler from sklearn.ensemble import RandomForestClassifier from sklearn.metrics import classification_report, accuracy_score, confusion_matrix
Step 2:	data=pd.read_csv("Weka_CustomerChurn.csv") print("the First 5 values of data is :\n")
	data.head()

Output:

the First 5 values of data is :

	Customer ID	Gender	Senior Citizen	Partner	Dependents	Tenure	Phone Service	Monthly Charges	Total Charges	Churn
0	1	Male	0	Yes	No	1	No	29.85	29.85	No
1	2	Female	1	No	No	34	Yes	56.95	1889.50	No
2	3	Female	0	Yes	Yes	2	Yes	53.85	108.15	Yes
3	4	Male	0	No	Yes	45	Yes	42.30	1840.75	No
4	5	Female	1	No	No	2	No	70.70	151.65	Yes

```
Step 3:
             X = data.drop(columns = ['Churn'])
      y = data['Churn'].apply(lambda x: 1 if x == 'Yes' else 0)
Step 4:
             X = pd.get \ dummies(X, drop \ first=True)
Step 5:
             X_{train}, X_{test}, y_{train}, y_{test} = train_{test} split(X, y, test_{size} = 0.2, random_{state} = 42)
     Step 6:
     scaler = StandardScaler()
              X train = scaler.fit transform(X train)
              X test = scaler.transform(X test)
Step 7:
     model = RandomForestClassifier(n estimators=100, random state=42)
Step 8:
             model.fit(X train, y train)
Output:
                  RandomForestClassifier
       RandomForestClassifier(random_state=42)
Step 9:
      y pred = model.predict(X test)
Step 10:
      accuracy = accuracy score(y test, y pred)
             conf_matrix = confusion_matrix(y_test, y_pred)
             report = classification report(y test, y pred)
```

Output:

C:\ProgramData\anaconda3\Lib\site-packages\sklearn\metrics_classification.py:1344: UndefinedMetricWarning: Precision and F-score are ill-defined and being set to 0.0 in labels with no predicted samples. Use `zero_division` parameter to control this behavior. _warn_prf(average, modifier, msg_start, len(result))

C:\ProgramData\anaconda3\Lib\site-packages\sklearn\metrics_classification.py:1344: UndefinedMetricWarning: Precision and F-score are ill-defined and being set to 0.0 in labels with no predicted samples. Use `zero_division` parameter to control this behavior. _warn_prf(average, modifier, msg_start, len(result))

C:\ProgramData\anaconda3\Lib\site-packages\sklearn\metrics_classification.py:1344: UndefinedMetricWarning: Precision and F-score are ill-defined and being set to 0.0 in labels with no predicted samples. Use `zero_division` parameter to control this behavior. warn prf(average, modifier, msg_start, len(result))

Step 11:

```
[] print(f'Accuracy: {accuracy}')
print('Confusion Matrix:')
print(conf_matrix)
print('Classification Report:')
print(report)
```

Output:

Accuracy: 0.5 Confusion Matrix:

[[0 1] [0 1]]

Classification Report:

	precision	recall	f1-score	suppor	į
0	0.00	0.00	0.00	1	
	0.50	1.00	0.67	1	
accuracy			0.50	2	
macro avg	0.25	0.50	0.33	2	
weighted avg	0.25	0.50	0.33	2	

EX.NO:	
DATE:	
	DIJII D AN UNGUDEDIJICED MODEL
	BUILD AN UNSUPERVISED MODEL
	USING [Scikit-Learn and PyTorch]
SOURCE COL	N.F
SOURCE COL	DE:
Step 1:	
Siep 1.	
	import pandas as pd
	from sklearn.datasets import load_iris
	from sklearn.preprocessing import StandardScaler
Step 2:	
	iris=load iris()
	X=pd.DataFrame(iris.data,columns=iris.feature_names)
Step 3:	
[]	Scaler=StandardScaler() V. Scaled=Scaler fit transform(V)
	X_Scaled=Scaler.fit_transform(X)
Step 4:	
	from sklearn.cluster import KMeans
	from sklearn.metrics import silhouette_score
	kmeans=KMeans(n_clusters=3,random_state=42)
	kmeans_labels=kmeans.fit_predict(X_Scaled)
Output :	
-	
Futu	rogramData\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:870: reWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the e of `n_init` explicitly to suppress the warning
	nings.warn(rogramData\anaconda3\Lib\site-packages\sklearn\cluster\ kmeans.py:1382:
	Warning: KMeans is known to have a memory leak on Windows with MKL, when there
are le	ess chunks than available threads. You can avoid it by setting the environment variable
	P_NUM_THREADS=1. rnings.warn(

```
Step 5:
     kmeans silhouette=silhouette score(X Scaled, kmeans labels)
            print(f'K-Means Silhouette Score: {kmeans silhouette}')
Output:
           K-Means Silhouette Score: 0.45994823920518635
Step 6:
     import torch
              import torch.nn as nn
              import torch.optim as optim
             from torch.utils.data import DataLoader, TensorDataset
             X tensor=torch.tensor(X Scaled, dtype=torch.float32)
Step 7:
            data\ loader=DataLoader(TensorDataset(X\ tensor),batch\ size=16,shuffle=True)
     Step 8:
     class Autoencoder(nn.Module):
               def __init__(self):
                 super(Autoencoder, self).__init__()
                 self.encoder=nn.Sequential(
                   nn.Linear(4,2),
                   nn.ReLU()
                 self.decoder=nn.Sequential(
                   nn.Linear(2,4),
                   nn.ReLU()
               def forward(self,x):
                 encoded=self.encoder(x)
                 decoded=self.decoder(encoded)
                 return decoded
```

```
Step 9:
     Г٦
            autoencoder=Autoencoder()
            criterion=nn.MSELoss()
            optimizer = optim.Adam(autoencoder.parameters(), lr = 0.01)
Step 10:
     num epochs=100
            for epoch in range(num epochs):
              for data in data loader:
                 inputs, =data
                 optimizer.zero grad()
                 outputs=autoencoder(inputs)
                 loss=criterion(outputs, inputs)
                 loss.backward()
                 optimizer.step()
Step 11:
            with torch.no grad():
     encoded data=autoencoder.encoder(X tensor).numpy()
Step 12:
            encoded kmeans=KMeans(n clusters=3,random state=42)
            encoded kmeans labels=encoded kmeans.fit predict(encoded data)
Output:
         C:\ProgramData\anaconda3\Lib\site-packages\sklearn\cluster\ kmeans.py:870:
         FutureWarning: The default value of `n init` will change from 10 to 'auto' in 1.4. Set
         the value of `n init` explicitly to suppress the warning
          warnings.warn(
         C:\ProgramData\anaconda3\Lib\site-packages\sklearn\cluster\ kmeans.py:1382:
         UserWarning: KMeans is known to have a memory leak on Windows with MKL,
         when there are less chunks than available threads. You can avoid it by setting the
         environment variable OMP NUM THREADS=1.
          warnings.warn(
```

Step 13: encoded_silhouette = silhouette_score(X_Scaled, encoded_kmeans_labels) []print(f'Autoencoder + kmeans silhouette score : {encoded_silhouette}') Output: Autoencoder + kmeans silhouette score : 0.302958422921628

	IMPLEMENT LOGISTIC REGRESSION
E COL	DE:
<i>1</i> :	
[]	import pandas as pd from sklearn.model_selection import train_test_split from sklearn.preprocessing import StandardScaler, LabelEncoder
	from sklearn.linear_model import LogisticRegression
	from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
p 2:	
[]	data=pd.read_csv("LogisticRegression_HeartDisease.csv") print("the First 5 values of data is : \n") data.head()
	0 1:

Output:

the First 5 values of data is:

	Age	CholesterolLevel	BloodPressure	SmokingStatus	HeartDisease
0	68	213.510222	119.336301	No	No
1	58	209.232228	109.821694	Yes	No
2	44	194.023226	154.003458	No	No
3	72	152.111973	162.855094	Yes	No
4	37	179.826361	121.379939	No	Yes

Step 3:

```
[] label_encoder = LabelEncoder()

data['SmokingStatus'] = label_encoder.fit_transform(data['SmokingStatus'])

data['HeartDisease'] = label_encoder.fit_transform(data['HeartDisease'])
```

Step 4:

X = data[['Age', 'CholesterolLevel', 'BloodPressure', 'SmokingStatus']] X.head()

Output:

	Age	CholesterolLevel	BloodPressure	SmokingStatus
0	68	213.510222	119.336301	0
1	58	209.232228	109.821694	1
2	44	194.023226	154.003458	0
3	72	152.111973	162.855094	1
4	37	179.826361	121,379939	0

Step 5:

[] y = data[['HeartDisease']] y.head()

Output:

		HeartDisease
	0	0
	1) 0
	2	0
1	3	0
/	4	1

Step 6:

 $[\]$ $X_{train}, X_{test}, y_{train}, y_{test} = train_{test_split}(X, y, test_{size}=0.2, random_{state}=42)$

Step 7:

[] scaler = StandardScaler()

X_train = scaler.fit_transform(X_train)

X_test = scaler.transform(X_test)

Step 8: model = LogisticRegression()model.fit(X_train, y_train) Output: C:\ProgramData\anaconda3\Lib\site-packages\sklearn\utils\validation.py:1143: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n_samples,), for example using ravel(). $y = column \ or \ ld(y, warn=True)$ ▼ LogisticRegression LogisticRegression() Step 9: y pred = model.predict(X test)**Step 10:** accuracy = accuracy score(y test, y pred) conf matrix = confusion matrix(y test, y pred)class report = classification report(y test, y pred) **Step 11:** print(f'Accuracy: {accuracy:.2f}') print('Confusion Matrix:') print(conf matrix) print('Classification Report:') print(class report) Output: Accuracy: 0.60 Confusion Matrix: [[7 3] [5 5]] Classification Report: precision recall f1-score support 10 0.58 0.70 0.64 0.62 0.56 10 0.50 0.60 20 accuracy 20 0.60 0.60 0.60 macro avg weighted avg 0.60 0.60 0.60 20

	NO: TE:										
			<i>IMPLE</i>	EMENT	NEUR	AL NE	TWORK	MODE	EL.		
so	URCE (CODE:									
	Step 1:										
		imp fro fro fro fro imp	port numpy port panda m sklearn., m tensorflo m tensorflo m sklearn., port matplo	s as pd model_sel preproces. ow.keras.h ow.keras.l metrics im otlib.pyplo	sing impo nodels imp ayers impo port class	rt Standar oort Seque ort Dense	dScaler ential	nfusion_m	atrix		
	Step 2:										
	Outpu	prodf.	=pd.read int("the F head() t 5 values	irst 5 valı	ues of dat		ctiveMair	ntenance.	csv")		
	sensor_1	sensor_2	sensor_3	sensor_4	sensor_5	sensor_6	sensor_7	sensor_8	sensor_9	sensor_10	failure
0	0.374540	0.950714	0.731994	0.598658	0.156019	0.155995	0.058084	0.866176	0.601115	0.708073	0

	sensor_1	sensor_2	sensor_3	sensor_4	sensor_5	sensor_6	sensor_7	sensor_8	sensor_9	sensor_10	failure
0	0.374540	0.950714	0.731994	0.598658	0.156019	0.155995	0.058084	0.866176	0.601115	0.708073	0
1	0.020584	0.969910	0.832443	0.212339	0.181825	0.183405	0.304242	0.524756	0.431945	0.291229	1
2	0.611853	0.139494	0.292145	0.366362	0.456070	0.785176	0.199674	0.514234	0.592415	0.046450	1
3	0.607545	0.170524	0.065052	0.948886	0.965632	0.808397	0.304614	0.097672	0.684233	0.440152	1
4	0.122038	0.495177	0.034389	0.909320	0.258780	0.662522	0.311711	0.520068	0.546710	0.184854	0

Step 3:

X = df.drop('failure', axis=1).values # Features (sensor data)y = df['failure'].values # Target (failure labels)

```
Step 4:
      scaler = StandardScaler()
             X \ scaled = scaler.fit \ transform(X)
Step 5:
     X train, X test, y train, y test = train test split(X scaled, y, test size=0.15,
             random\ state=42)
             X train, X val, y train, y val = train test split(X train, y train, test size=0.15,
             random state=42)
Step 6:
      Г٦
             model = Sequential()
             model.add(Dense(64, input dim=X.shape[1], activation='relu'))
             model.add(Dense(32, activation='relu'))
             model.add(Dense(1, activation='sigmoid'))
Output:
     C:\ProgramData \anaconda3\Lib\site-packages\keras\src\layers\core\dense.py:87: UserWarning:
    Do not pass an 'input_shape'/'input_dim' argument to a layer. When using Sequential models,
    prefer using an 'Input(shape)' object as the first layer in the model instead.
     super(). init (activity regularizer=activity regularizer, **kwargs)
Step 7:
     []
             model.compile(optimizer='adam', loss='binary_crossentropy', metrics=['accuracy'])
Step 8:
             # Train the model with fewer epochs
      history = model.fit(X train, y train, validation data=(X val, y val), epochs=10,
             batch size=32)
             # Evaluate the model on the test set
             test loss, test accuracy = model.evaluate(X test, y test)
             print(f"Test Loss: {test loss}")
             print(f"Test Accuracy: {test accuracy}")
             # Predict and generate classification report
             y pred = (model.predict(X test) > 0.5).astype("int32")
             print(classification report(y test, y pred))
```

Output:

```
Epoch 1/10
23/23
                                                      0s 4ms/step - accuracy: 0.7976 - loss: 0.5117 -
val accuracy: 0.5156 - val loss: 0.7832
Epoch 2/10
23/23
                                                      0s 2ms/step - accuracy: 0.7784 - loss: 0.5235 -
val accuracy: 0.5156 - val loss: 0.7902
Epoch 3/10
23/23 -
                                                      Os 3ms/step - accuracy: 0.7891 - loss: 0.5068 -
val accuracy: 0.5000 - val loss: 0.7948
Epoch 4/10
23/23
                                                      0s 3ms/step - accuracy: 0.7687 - loss: 0.5194 -
val accuracy: 0.4922 - val loss: 0.8020
Epoch 5/10
23/23
                                                       0s 2ms/step - accuracy: 0.7619 - loss: 0.5247 -
val accuracy: 0.5312 - val loss: 0.8121
Epoch 6/10
23/23
                                                       0s 2ms/step - accuracy: 0.7967 - loss: 0.5169 -
val accuracy: 0.5234 - val loss: 0.8118
Epoch 7/10
23/23
                                                      0s 2ms/step - accuracy: 0.7831 - loss: 0.5106 -
val accuracy: 0.5000 - val loss: 0.8141
Epoch 8/10
23/23
                                                       0s 3ms/step - accuracy: 0.7767 - loss: 0.5078 -
val accuracy: 0.5391 - val loss: 0.8291
Epoch 9/10
23/23
                                                       0s 2ms/step - accuracy: 0.7946 - loss: 0.4805 -
val_accuracy: 0.4922 - val_loss: 0.8261
Epoch 10/10
23/23
                                                      0s 2ms/step - accuracy: 0.8199 - loss: 0.4780 -
val accuracy: 0.5156 - val loss: 0.8287
                                                    0s 2ms/step - accuracy: 0.4935 - loss: 0.7497
Test Loss: 0.7296823263168335
Test Accuracy: 0.5
                                                    Os 2ms/step
5/5/
                            precision
                                         recall f1-score
                                                             support
                              0.48
                                         0.44
                       0
                                                   0.46
                                                               73
                              0.51
                                        0.56
                                                  0.53
                                                               77
             accuracy
                                                  0.50
                                                            150
           macro avg
                            0.50
                                       0.50
                                                 0.50
                                                            150
        weighted avg
                            0.50
                                      0.50
                                                 0.50
                                                            150
```

Step 9:

```
[] # Confusion Matrix

cm = confusion_matrix(y_test, y_pred)

plt.figure(figsize=(6, 4))

sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['No Failure',

'Failure'], yticklabels=['No Failure', 'Failure'])

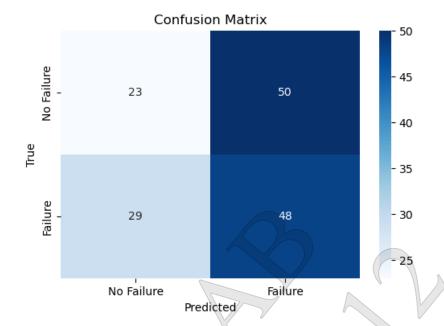
plt.title('Confusion Matrix')

plt.xlabel('Predicted')

plt.ylabel('True')

plt.show()
```





Step 10:

```
[] # Plot learning curves for loss

plt.figure(figsize=(10, 5))

plt.plot(history.history['loss'], label='Training Loss')

plt.plot(history.history['val_loss'], label='Validation Loss')

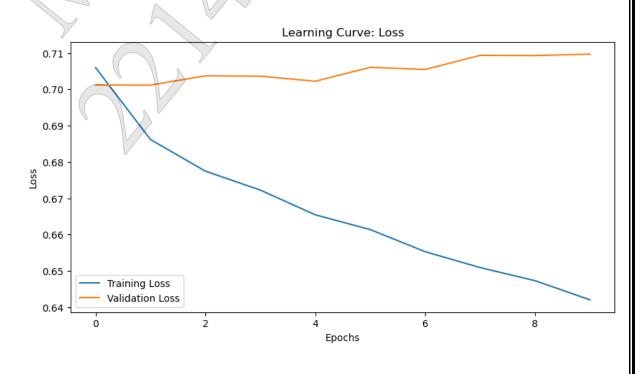
plt.title('Learning Curve: Loss')

plt.xlabel('Epochs')

plt.ylabel('Loss')

plt.legend()

plt.show()
```



Step 11:

```
# Plot learning curves for accuracy

plt.figure(figsize=(10, 5))

plt.plot(history.history['accuracy'], label='Training Accuracy')

plt.plot(history.history['val_accuracy'], label='Validation Accuracy')

plt.title('Learning Curve: Accuracy')

plt.xlabel('Epochs')

plt.ylabel('Accuracy')

plt.legend()

plt.show()
```

