#### **UCS2612 Machine Learning Laboratory**

#### A9. Applications of dimensionality reductiontechniques

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## Aim

Develop a python program to perform dimensionality reduction using PCA and LDA. Visualize the features from the dataset and interpret the results obtained by the model using Matplotlib library.

Dataset:- http://www3.dsi.uminho.pt/pcortez/wine/winequality.zip

## Import Libraries

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.metrics import roc curve, roc auc score
from sklearn.model selection import train test split
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.preprocessing import LabelEncoder
from sklearn.preprocessing import StandardScaler, MinMaxScaler
from sklearn.preprocessing import StandardScaler
from sklearn.ensemble import RandomForestClassifier,
GradientBoostingClassifier
from sklearn.metrics import accuracy score, classification report,
confusion matrix
import numpy as np
import matplotlib.pyplot as plt
```

#### Read dataset

```
# importing or loading the dataset
data = pd.read csv(""C:\Users\SSN\Desktop\ML
Lab\A9\winequality.zip"", header=0, sep=";")
data.head()
   fixed acidity volatile acidity citric acid residual sugar
chlorides \
0
             7.4
                              0.70
                                            0.00
                                                             1.9
0.076
             7.8
                              0.88
                                            0.00
                                                             2.6
0.098
                              0.76
             7.8
                                           0.04
                                                             2.3
0.092
            11.2
                              0.28
                                            0.56
                                                             1.9
0.075
             7.4
                              0.70
                                            0.00
                                                             1.9
0.076
   free sulfur dioxide total sulfur dioxide density pH sulphates
0
                  11.0
                                         34.0 0.9978 3.51
                                                                    0.56
                  25.0
                                         67.0 0.9968 3.20
1
                                                                    0.68
```

2	15	.0	5	54.0	0.9970	3.26	0.65			
3	17	.0	6	50.0	0.9980	3.16	0.58			
4	11	.0	3	34.0	0.9978	3.51	0.56			
alcoho 0 9. 1 9. 2 9. 3 9. 4 9.	4 5 8 5 8 5 8 6									
data.describe()										
	xed acidity 1599.000000 8.319637 1.741096 4.600000 7.100000 7.900000 9.200000 15.900000	0.9	1	0.27 0.19 0.00 0.09 0.26 0.42		esidual su 1599.00 2.53 1.40 0.90 1.90 2.20 2.60 15.50	0000 8806 9928 0000 0000 0000			
		free sulfur	dioxide	total	sulfur o	dioxide				
density	99.000000	1599	.000000		1599	.000000				
1599.0000 mean	0.087467	15	.874922		46	.467792				
0.996747 std	0.047065	10	.460157		32	.895324				
0.001887 min 0.990070	0.012000	1	.000000		6	.000000				
25%	0.070000	7	.000000		22	.000000				
0.995600 50%	0.079000	14	.000000		38	.000000				
0.996750 75%	0.090000	21	.000000		62	.000000				
0.997835 max 1.003690	0.611000	72	.000000		289	.000000				
count 15 mean std	pH 599.000000 3.311113 0.154386	sulphates 1599.000000 0.658149 0.169507	alcc 1599.000 10.422 1.065	) ) ) ) ) ) ) )	qual 1599.000 5.636 0.807	000 023				

$egin{array}{cccccccccccccccccccccccccccccccccccc$
--

# **Data Pre-processing**

data.corr()  fixed acidity volatile acidity citric ac		
1 000000	id \	
fixed acidity 1.000000 -0.256131 0.6717	)3	
volatile acidity -0.256131 1.000000 -0.5524	96	
citric acid 0.671703 -0.552496 1.0000	00	
residual sugar 0.114777 0.001918 0.1435	77	
chlorides 0.093705 0.061298 0.2038	0.203823	
free sulfur dioxide -0.153794 -0.010504 -0.0609	-0.060978	
total sulfur dioxide -0.113181 0.076470 0.0355	0.035533	
density 0.668047 0.022026 0.3649	0.364947	
pH -0.682978 0.234937 -0.5419	-0.541904	
sulphates 0.183006 -0.260987 0.3127	0.312770	
alcohol -0.061668 -0.202288 0.1099	0.109903	
quality 0.124052 -0.390558 0.2263	73	
residual sugar chlorides free sulfur		
dioxide \		
fixed acidity 0.114777 0.093705 -0.15	3794	
volatile acidity 0.001918 0.061298 -0.01	2504	
volatile actualty 0.001910 0.001290 -0.01	1504	
citric acid 0.143577 0.203823 -0.06	0978	
residual sugar 1.000000 0.055610 0.18	7049	
chlorides 0.055610 1.000000 0.00	5562	
free sulfur dioxide 0.187049 0.005562 1.00	0000	
0.000000 0.045400	7.6.6.6	
total sulfur dioxide 0.203028 0.047400 0.66	/666	
density 0.355283 0.200632 -0.02	1016	
density 0.355283 0.200632 -0.02	1940	
рH -0.085652 -0.265026 0.07	0377	
0.000000		
sulphates 0.005527 0.371260 0.05	1658	
alcohol 0.042075 -0.221141 -0.06	9408	
quality 0.013732 -0.128907 -0.05	0656	

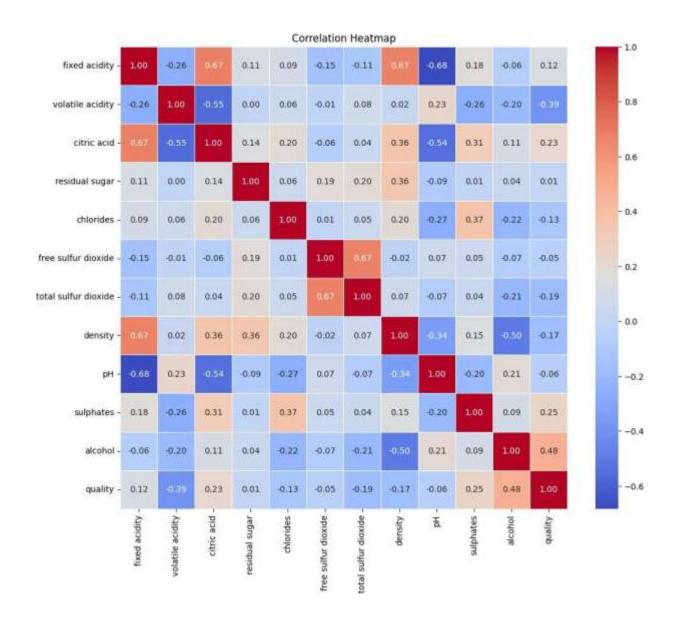
```
total sulfur dioxide density pH
sulphates \
                                -0.113181 0.668047 -0.682978
fixed acidity
0.183006
volatile acidity
                                 0.076470 0.022026 0.234937 -
0.260987
citric acid
                                 0.035533 0.364947 -0.541904
0.312770
residual sugar
                                  0.203028 0.355283 -0.085652
0.005527
chlorides
                                  0.047400 0.200632 -0.265026
0.371260
free sulfur dioxide
                                 0.667666 -0.021946 0.070377
0.051658
total sulfur dioxide
                                 1.000000 0.071269 -0.066495
0.042947
                                  0.071269 1.000000 -0.341699
density
0.148506
                                 -0.066495 -0.341699 1.000000 -
Нф
0.196648
sulphates
                                  0.042947 0.148506 -0.196648
1.000000
                                 -0.205654 -0.496180 0.205633
alcohol
0.093595
                                 -0.185100 -0.174919 -0.057731
quality
0.251397
                       alcohol quality
                     -0.061668 0.124052
fixed acidity
                   -0.202288 -0.390558
volatile acidity
citric acid
                    0.109903 0.226373
residual sugar
                     0.042075 0.013732
chlorides
                    -0.221141 -0.128907
free sulfur dioxide -0.069408 -0.050656
total sulfur dioxide -0.205654 -0.185100
density
                    -0.496180 -0.174919
                     0.205633 -0.057731
рН
                     0.093595 0.251397
sulphates
alcohol
                      1.000000 0.476166
                     0.476166 1.000000
quality
data.dropna(inplace=True)
scaler standard = StandardScaler()
data standardized = scaler standard.fit transform(data)
scaler normal = MinMaxScaler()
data normalized = scaler normal.fit transform(data)
```

```
data standardized = pd.DataFrame(data standardized,
columns=data.columns)
data normalized = pd.DataFrame(data normalized, columns=data.columns)
data standardized.head()
  fixed acidity volatile acidity citric acid residual sugar
chlorides \
      -0.528360
                         0.961877 -1.391472
                                                    -0.453218 -
0.243707
                         1.967442 -1.391472
      -0.298547
                                                     0.043416
0.223875
2 -0.298547
                                                    -0.169427
                         1.297065 -1.186070
0.096353
3 1.654856
                        -1.384443 1.484154
                                                    -0.453218 -
0.264960
4 -0.528360
                         0.961877 - 1.391472
                                                    -0.453218 -
0.243707
  free sulfur dioxide total sulfur dioxide density pH
sulphates \
            -0.466193
                                  -0.379133 0.558274 1.288643 -
0.579207
                                   0.624363 0.028261 -0.719933
             0.872638
0.128950
                                   0.229047 0.134264 -0.331177 -
            -0.083669
0.048089
                                   0.411500 0.664277 -0.979104 -
             0.107592
0.461180
            -0.466193
                                  -0.379133 0.558274 1.288643 -
0.579207
  alcohol quality
0 -0.960246 -0.787823
1 -0.584777 -0.787823
2 -0.584777 -0.787823
3 -0.584777 0.450848
4 -0.960246 -0.787823
data normalized.head()
  fixed acidity volatile acidity citric acid residual sugar
chlorides \
       0.247788
                         0.397260
                                         0.00
                                                     0.068493
0.106845
      0.283186
                         0.520548
                                         0.00
                                                     0.116438
0.143573
       0.283186
                         0.438356
                                         0.04
                                                     0.095890
0.133556
      0.584071
                                         0.56
                         0.109589
                                                     0.068493
```

```
0.105175
4 0.247788
                        0.397260 0.00 0.068493
0.106845
  free sulfur dioxide total sulfur dioxide density pH
sulphates \
                                 0.098940 0.567548 0.606299
             0.140845
0
0.137725
                                 0.215548 0.494126 0.362205
             0.338028
0.209581
                                 0.169611 0.508811 0.409449
             0.197183
0.191617
                                 0.190813 0.582232 0.330709
             0.225352
0.149701
                                 0.098940 0.567548 0.606299
             0.140845
0.137725
  alcohol quality
0 0.153846
                0.4
1 0.215385
               0.4
2 0.215385
               0.4
3 0.215385
               0.6
4 0.153846
               0.4
```

### **EDA**

```
plt.figure(figsize=(12, 10))
sns.heatmap(data.corr(), annot=True, cmap='coolwarm', fmt='.2f',
linewidths=0.5)
plt.title('Correlation Heatmap')
plt.show()
```



# Splitting the data into testing and training

```
# distributing the dataset into two components X and Y
X_red = data.iloc[:, 0:11].values
y_red = data.iloc[:, 11].values

X_train_red, X_test_red, y_train_red, y_test_red =
train_test_split(X_red, y_red, test_size=0.2, random_state=0)
```

## Feature engineering

```
# performing preprocessing part
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()

X_train_red = sc.fit_transform(X_train_red)
X_test_red = sc.transform(X_test_red)
```

## **Building PCA model**

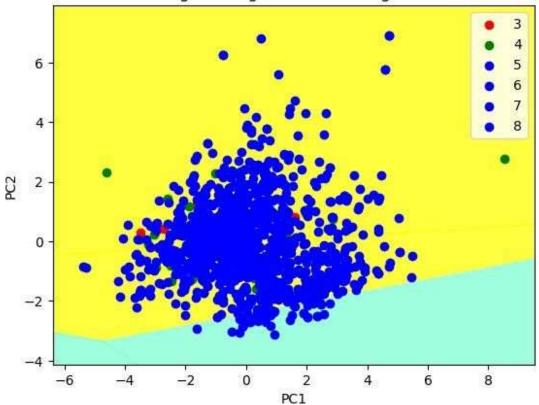
```
from sklearn.decomposition import PCA
PCa = PCA (n components = 2)
X train red = PCa.fit transform(X train red)
X test red = PCa.transform(X test red)
explained variance = PCa.explained variance ratio
from sklearn.linear model import LogisticRegression
classifier = LogisticRegression(random state = 0)
classifier.fit(X train red, y train red)
LogisticRegression(random state=0)
y pred red = classifier.predict(X test red)
from sklearn.metrics import confusion matrix
cm = confusion matrix(y test red, y pred red)
print(cm)
accuracy score (y test red, y pred red)
[[0 0 0 2 0 0]
 [0 0 4 7 0 0]
 [ 0 0 89 45 1 0]
 [ 0 0 55 81 6 0]
 [ 0 0 4 21 2 0]
[0 0 0 2 1 0]
0.5375
```

### Visualisation of PCA model

```
# result through scatter plot
from matplotlib.colors import ListedColormap
```

```
X set, y set = X train red, y train red
X1, X2 = np.meshgrid(np.arange(start = X set[:, 0].min() - 1,
                     stop = X set[:, 0].max() + 1, step = 0.01),
                     np.arange(start = X \text{ set}[:, 1].min() - 1,
                     stop = X set[:, 1].max() + 1, step = 0.01))
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(),
             X2.ravel()).T).reshape(X1.shape), alpha = 0.75,
             cmap = ListedColormap(('yellow', 'white', 'aquamarine')))
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y set)):
   plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
                c = ListedColormap(('red', 'green', 'blue'))(i), label
= j)
plt.title('Logistic Regression (Training set)')
plt.xlabel('PC1') # for Xlabel plt.ylabel('PC2')
# for Ylabel
plt.legend() # to show legend
# show scatter plot
plt.show()
C:\Users\nithi\AppData\Local\Temp\ipykernel 4776\4008791166.py:18:
UserWarning: *c* argument looks like a single numeric RGB or RGBA
sequence, which should be avoided as value-mapping will have
precedence in case its length matches with *x* & *y*. Please use the
*color* keyword-argument or provide a 2D array with a single row if
you intend to specify the same RGB or RGBA value for all points.
 plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
```

#### Logistic Regression (Training set)



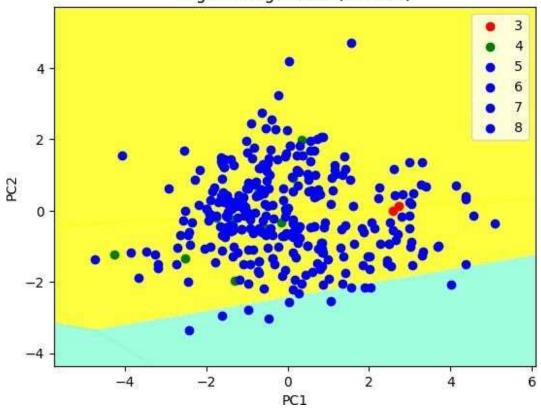
```
# Visualising the Test set results through scatter plot
from matplotlib.colors import ListedColormap
X set, y set = X test red, y test red
X1, X2 = np.meshgrid(np.arange(start = X_set[:, 0].min() - 1,
                     stop = X set[:, 0].max() + 1, step = 0.01),
                     np.arange(start = X set[:, 1].min() - 1,
                     stop = X set[:, 1].max() + 1, step = 0.01))
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(),
             X2.ravel()]).T).reshape(X1.shape), alpha = 0.75,
             cmap = ListedColormap(('yellow', 'white', 'aquamarine')))
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y set)):
    plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
                c = ListedColormap(('red', 'green', 'blue'))(i), label
= j)
```

```
# title for scatter plot
plt.title('Logistic Regression (Test set)')
plt.xlabel('PC1') # for Xlabel
plt.ylabel('PC2') # for Ylabel
plt.legend()

# show scatter plot
plt.show()

C:\Users\nithi\AppData\Local\Temp\ipykernel_4776\3957206111.py:19:
UserWarning: *c* argument looks like a single numeric RGB or RGBA
sequence, which should be avoided as value-mapping will have
precedence in case its length matches with *x* & *y*. Please use the
*color* keyword-argument or provide a 2D array with a single row if
you intend to specify the same RGB or RGBA value for all points.
plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1],
```

#### Logistic Regression (Test set)



# **Building LDA model**

from sklearn.discriminant\_analysis import LinearDiscriminantAnalysis
as LDA

```
lda = LDA(n components = 2)
X train red = lda.fit transform(X train red, y train red)
X test red = lda.transform(X test red)
from sklearn.linear model import LogisticRegression
from sklearn.metrics import accuracy score
classifier = LogisticRegression(random state = 0)
classifier.fit(X train red, y train red)
LogisticRegression(random state=0)
y pred = classifier.predict(X test red)
print(y pred)
[5\ 5\ 6\ 6\ 7\ 5\ 6\ 5\ 6\ 6\ 6\ 6\ 5\ 5\ 7\ 6\ 6\ 5\ 6\ 6\ 6\ 5\ 5\ 5\ 5\ 5\ 6\ 6\ 6\ 5\ 5\ 5\ 5
5 6 5 6 5 6 6 6 6 5 6 5 5 5 6 6 5 6 6 6 6 6 5 6 5 6 6 6 5 6 5 6 6
6 7
5 7
5 6
5 5 5 5 5 6 6 5 6 5 5 5 5 5 5 5 5 6 5 7 6 6 6 7]
accuracy = accuracy score(y test red, y pred)
print("Accuracy of LDA model:", accuracy)
Accuracy of LDA model: 0.5375
```

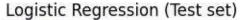
#### Visualisation of LDA model

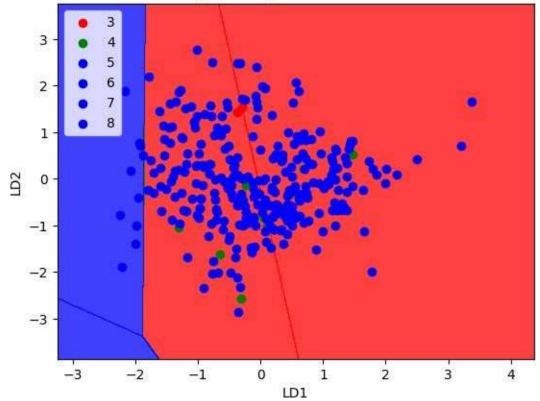
```
from matplotlib.colors import ListedColormap
X_set, y_set = X_test_red, y_test_red

X1, X2 = np.meshgrid(np.arange(start = X_set[:, 0].min() - 1, stop = X_set[:, 0].max() + 1, step = 0.01), np.arange(start = X_set[:, 1].min() - 1, stop = X_set[:, 1].max() + 1, step = 0.01))

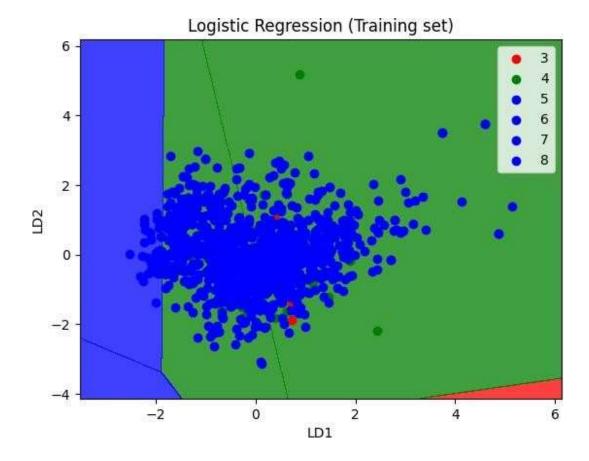
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(), X2.ravel()]).T).reshape(X1.shape), alpha = 0.75, cmap = ListedColormap(('red', 'green', 'blue')))
```

```
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y set)):
    plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
                c = ListedColormap(('red', 'green', 'blue'))(i), label
= j)
plt.title('Logistic Regression (Test set)')
plt.xlabel('LD1')
plt.ylabel('LD2')
plt.legend()
plt.show()
C:\Users\nithi\AppData\Local\Temp\ipykernel 4776\531035640.py:12:
UserWarning: *c* argument looks like a single numeric RGB or RGBA
sequence, which should be avoided as value-mapping will have
precedence in case its length matches with *x* & *y*. Please use the
*color* keyword-argument or provide a 2D array with a single row if
you intend to specify the same RGB or RGBA value for all points.
  plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
```





```
from matplotlib.colors import ListedColormap
X set, y set = X train red, y train red
X1, X2 = np.meshgrid(np.arange(start = X set[:, 0].min() - 1, stop =
X = [:, 0].max() + 1, step = [0.01], np.arange(start = [x] set[:,
1].min() - 1, stop = X set[:, 1].max() + 1, step = 0.01))
plt.contourf(X1, X2, classifier.predict(np.array([X1.ravel(), X2.ravel()]
).T).reshape(X1.shape), alpha = 0.75, cmap = ListedColormap(('red',
'green', 'blue')))
plt.xlim(X1.min(), X1.max())
plt.ylim(X2.min(), X2.max())
for i, j in enumerate(np.unique(y set)):
   plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
                c = ListedColormap(('red', 'green', 'blue'))(i), label
= j)
plt.title('Logistic Regression (Training set)')
plt.xlabel('LD1')
plt.ylabel('LD2')
plt.legend()
plt.show()
C:\Users\nithi\AppData\Local\Temp\ipykernel 4776\171321940.py:12:
UserWarning: *c* argument looks like a single numeric RGB or RGBA
sequence, which should be avoided as value-mapping will have
precedence in case its length matches with *x* & *y*. Please use the
*color* keyword-argument or provide a 2D array with a single row if
you intend to specify the same RGB or RGBA value for all points.
 plt.scatter(X set[y set == j, 0], X set[y set == j, 1],
```



### Inference

- 1. After applying PCA, you can analyze the principal components to understand which original features contribute the most to the variance in the data. You can also visualize the data in reduced dimensions to explore patterns or clusters.
- 2. After applying LDA, you can interpret the learned linear discriminants to understand how the classes are separated in the reduced-dimensional space. LDA provides insight into which features are most discriminative for class separation.

# **Learning Outcomes**

- 1. Implementation of Pre-processing, EDA and feature selection.
- 2. Implementation of PCA nad LDA models and visualising it.
- 3. Displaying the confusion matrix.
- 4. Understanding the techniques of dimentionality reduction.