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Year of study: 4th Year

Semester: 1st Sem

Subject name: Machine Learning Lab (Subject Code: IT/PC/B/S/411)

Git link: https://github.com/GargiGhosal/MachineLearning_Lab.git

Solutions::

Q1. Datasets used :

- a. Wine Dataset: <https://archive.ics.uci.edu/ml/datasets/wine>
- b. Ionosphere Dataset: <https://archive.ics.uci.edu/ml/datasets/Ionosphere>

Q1. Answers::

```
# WINE DATASET
```

```
# SVM(Without Tuning)[70-30 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("wine.data",header=None)
```

```
col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total  
phenols','Flavanoids',
```

```
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of  
diluted wines','Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.svm import SVC
```

```
classifier = SVC()
```

```
classifier.fit(X_train,y_train)
```

```
y_pred = classifier.predict(X_test)
```

```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, y_pred))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, y_pred))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, y_pred))
```

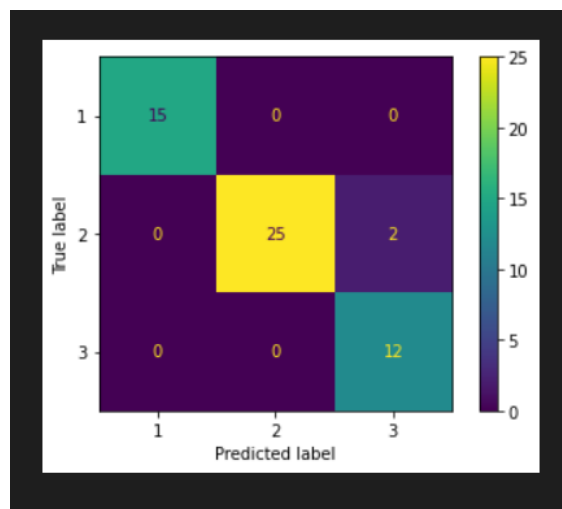
```
import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()
```

```
Confusion Matrix:
[[15  0  0]
 [ 0 25  2]
 [ 0  0 12]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     1       1.00      1.00      1.00        15
     2       1.00      0.93      0.96        27
     3       0.86      1.00      0.92        12

 accuracy          0.96        54
 macro avg          0.95      0.98      0.96        54
weighted avg          0.97      0.96      0.96        54

-----
Accuracy:
0.9629629629629629
```



WINE DATASET

SVM(Without Tuning)[60-40 split]

```

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv("wine.data",header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of
diluted wines','Proline']

df.columns = col_name

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)

```

```
X_test = sc.transform(X_test)

# Classification
from sklearn.svm import SVC

classifier = SVC()
classifier.fit(X_train,y_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
```

```

from sklearn.metrics import plot_confusion_matrix

plot_confusion_matrix(classifier, X_test, y_test)

plt.show()

```

```

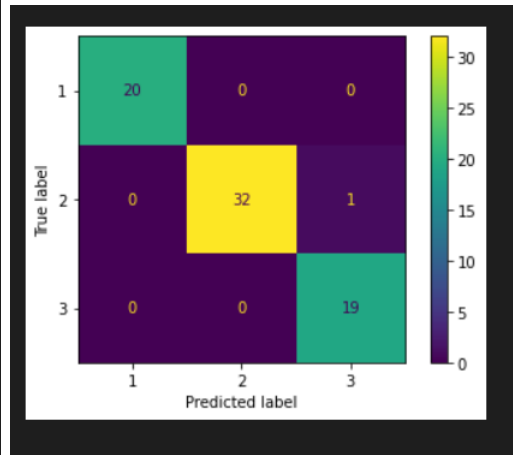
Confusion Matrix:
[[20  0  0]
 [ 0 32  1]
 [ 0  0 19]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     1         1.00      1.00      1.00        20
     2         1.00      0.97      0.98        33
     3         0.95      1.00      0.97        19

 accuracy          0.99          0.99          0.99          72
 macro avg          0.98          0.99          0.99          72
weighted avg          0.99          0.99          0.99          72

-----
Accuracy:
0.9861111111111112

```



WINE DATASET

SVM(Without Tuning)[40-60 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("wine.data",header=None)
```

```
col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',
```

```
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of
diluted wines','Proline']
```

```
df.columns = col_name
```

```

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from sklearn.svm import SVC

classifier = SVC()
classifier.fit(X_train,y_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

```

```

print("Performance Evaluation")

print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()

```

```

Confusion Matrix:
[[34  0  0]
 [ 0 42  0]
 [ 0  0 31]]
-----

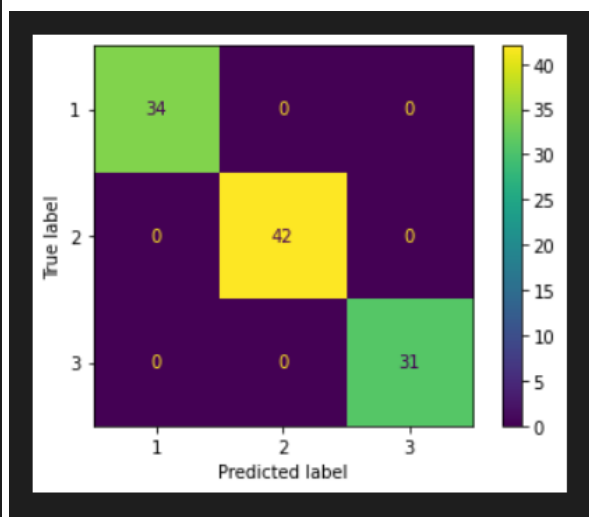
Performance Evaluation
      precision    recall  f1-score   support

     1       1.00      1.00      1.00        34
     2       1.00      1.00      1.00        42
     3       1.00      1.00      1.00        31

   accuracy          1.00          107
  macro avg          1.00          107
weighted avg          1.00          107
-----

Accuracy:
1.0

```



WINE DATASET

SVM(With Tuning)[40-60 split]


```

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv("wine.data",header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of
diluted wines','Proline']

df.columns = col_name

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

```

```

# Classification
from sklearn.svm import SVC

classifier = SVC()

#####
#
# Showing all the parameters

from pprint import pprint
# Look at parameters used by our current forest
print('Parameters currently in use:\n')
pprint(classifier.get_params())

#####
#
# Creating a set of important sample features

param_grid = {'C': [0.1,1, 10, 100], 'gamma': [1,0.1,0.01,0.001], 'kernel': ['rbf', 'poly',
'sigmoid']}

pprint(param_grid)

#####
#

from sklearn.model_selection import GridSearchCV

# Use the random grid to search for best hyperparameters
# First create the base model to tune

```

```

classifier = SVC()

# Random search of parameters, using 3 fold cross validation,
# search across 100 different combinations, and use all available cores

rf_random = GridSearchCV(SVC(), param_grid, refit=True, verbose=2)
rf_random.fit(X_train, y_train)

y_pred = rf_random.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

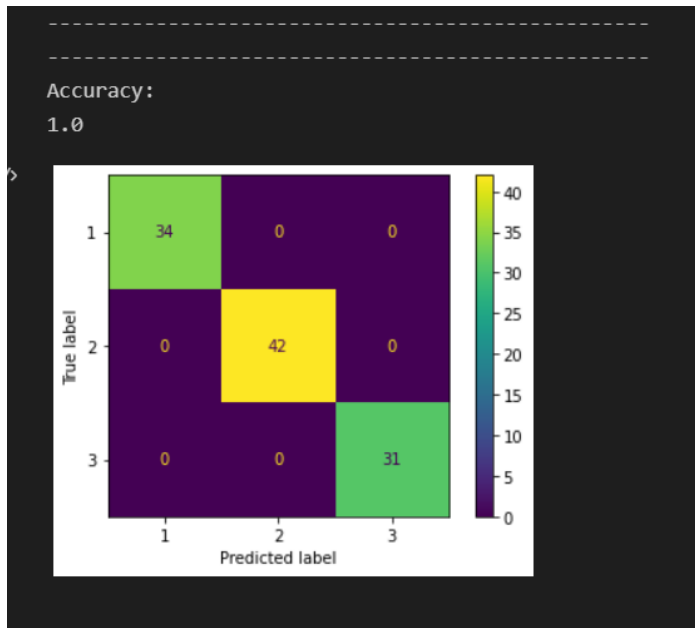
import matplotlib.pyplot as plt

```

```

from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(rf_random, X_test, y_test)
plt.show()

```



WINE DATASET

Decision Tree (Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("wine.data", header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of ash', 'Magnesium', 'Total phenols', 'Flavanoids',
```

```
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from sklearn.tree import DecisionTreeClassifier

classifier = DecisionTreeClassifier().fit(X_train,y_train)
classifier.fit(X_train,y_train)

y_pred=classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")

```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
from sklearn.metrics import plot_confusion_matrix
```

```
plot_confusion_matrix(classifier, X_test, y_test)
```

```
plt.show()
```

```
from sklearn import tree
```

```
text_representation = tree.export_text(classifier)
```

```
print(text_representation)
```

Confusion Matrix:

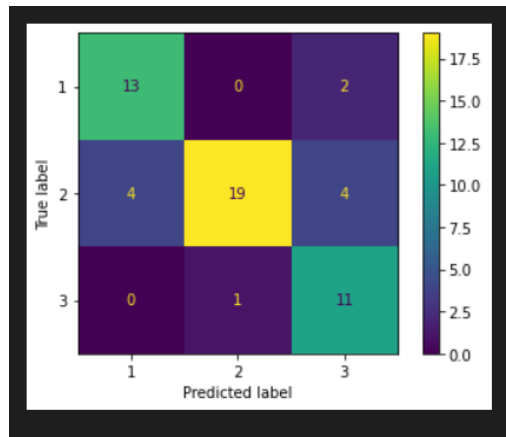
```
[[13  0  2]
 [ 4 19  4]
 [ 0  1 11]]
```

Performance Evaluation

	precision	recall	f1-score	support
1	0.76	0.87	0.81	15
2	0.95	0.70	0.81	27
3	0.65	0.92	0.76	12
accuracy			0.80	54
macro avg	0.79	0.83	0.79	54
weighted avg	0.83	0.80	0.80	54

Accuracy:

0.7962962962962963



```
|--- feature_12 <= 0.06
| |--- feature_9 <= -0.47
| | |--- class: 2
| |--- feature_9 > -0.47
| | |--- feature_10 <= 0.17
| | | |--- class: 3
| | |--- feature_10 > 0.17
| | | |--- feature_11 <= 0.38
| | | | |--- class: 2
| | | |--- feature_11 > 0.38
| | | | |--- class: 1
|--- feature_12 > 0.06
| |--- feature_6 <= 0.16
| | |--- feature_7 <= -0.76
| | | |--- class: 2
| | |--- feature_7 > -0.76
| | | |--- class: 3
| |--- feature_6 > 0.16
| | |--- feature_3 <= 2.32
| | | |--- class: 1
| | |--- feature_3 > 2.32
| | | |--- class: 2
```

WINE DATASET

Decision Tree (With Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv("wine.data",header=None)

```
col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total  
phenols','Flavanoids',  
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of  
diluted wines','Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.tree import DecisionTreeClassifier
```

```
classifier = DecisionTreeClassifier(criterion="entropy",max_depth=20).fit(X_train,y_train)
```

```
classifier.fit(X_train,y_train)
```



```

y_pred=classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()

from sklearn import tree
text_representation = tree.export_text(classifier)
print(text_representation)

```

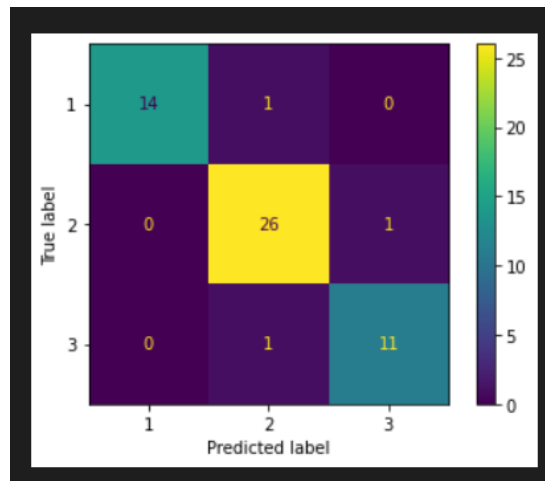
```

Confusion Matrix:
[[14  1  0]
 [ 0 26  1]
 [ 0  1 11]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     1         1.00      0.93      0.97        15
     2         0.93      0.96      0.95        27
     3         0.92      0.92      0.92        12

 accuracy          0.94          54
 macro avg         0.95          54
weighted avg         0.95          54
-----
Accuracy:
0.9444444444444444

```



```

|--- feature_6 <= -0.60
| |--- feature_9 <= -0.61
| | |--- class: 2
| |--- feature_9 > -0.61
| | |--- class: 3
|--- feature_6 > -0.60
| |--- feature_12 <= -0.14
| | |--- class: 2
| |--- feature_12 > -0.14
| | |--- feature_9 <= -0.71
| | | |--- class: 2
| | |--- feature_9 > -0.71
| | | |--- class: 1

```

WINE DATASET

Random Forest Classifier(Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("wine.data",header=None)
```

```
col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of
diluted wines','Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.ensemble import RandomForestClassifier
```

```
classifier = RandomForestClassifier(n_estimators=20, random_state=0)
```

```
classifier.fit(X_train,y_train)
```

```
y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()
```

Confusion Matrix:

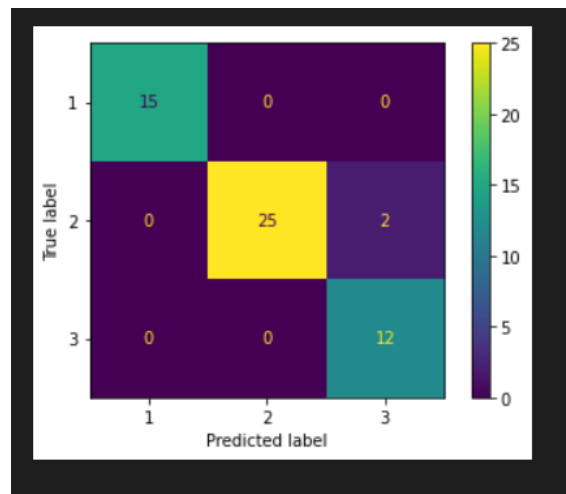
```
[[15  0  0]
 [ 0 25  2]
 [ 0  0 12]]
```

Performance Evaluation

	precision	recall	f1-score	support
1	1.00	1.00	1.00	15
2	1.00	0.93	0.96	27
3	0.86	1.00	0.92	12
accuracy			0.96	54
macro avg	0.95	0.98	0.96	54
weighted avg	0.97	0.96	0.96	54

Accuracy:

0.9629629629629629



WINE DATASET

Random Forest Classifier(With Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv("wine.data",header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',

'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of
diluted wines','Proline']

df.columns = col_name

```

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.30,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from sklearn.ensemble import RandomForestClassifier

classifier = RandomForestClassifier()

#####
#
# Showing all the parameters

from pprint import pprint
# Look at parameters used by our current forest
print('Parameters currently in use:\n')
pprint(classifier.get_params())

```

```
#####
#
# Creating a set of important sample features

from sklearn.model_selection import RandomizedSearchCV

# Number of trees in random forest
n_estimators = [int(x) for x in np.linspace(start = 200, stop = 2000, num = 10)]

# Number of features to consider at every split
max_features = ['auto', 'sqrt']

# Maximum number of levels in tree
max_depth = [int(x) for x in np.linspace(10, 110, num = 11)]
max_depth.append(None)

# Minimum number of samples required to split a node
min_samples_split = [2, 5, 10]

# Minimum number of samples required at each leaf node
min_samples_leaf = [1, 2, 4]

# Method of selecting samples for training each tree
bootstrap = [True, False]

# Create the random grid
random_grid = {'n_estimators': n_estimators,
               'max_features': max_features,
               'max_depth': max_depth,
               'min_samples_split': min_samples_split,
               'min_samples_leaf': min_samples_leaf,
               'bootstrap': bootstrap}

pprint(random_grid)

#####
#

# Use the random grid to search for best hyperparameters
```

```

# First create the base model to tune
classifier = RandomForestClassifier()

# Random search of parameters, using 3 fold cross validation,
# search across 100 different combinations, and use all available cores
rf_random = RandomizedSearchCV(estimator = classifier, param_distributions =
random_grid, n_iter = 100, cv = 3, verbose=2, random_state=42, n_jobs = -1)

# Fit the random search model
rf_random.fit(X_train, y_train)

y_pred = rf_random.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt

```



```

from sklearn.metrics import plot_confusion_matrix

plot_confusion_matrix(rf_random, X_test, y_test)

plt.show()

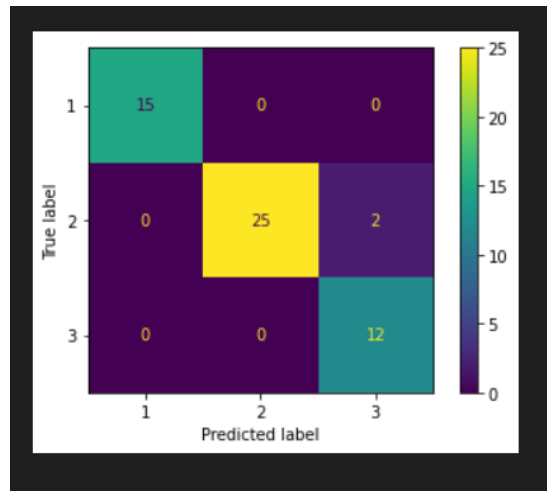
```

```

{'bootstrap': True,
 'ccp_alpha': 0.0,
 'class_weight': None,
 'criterion': 'gini',
 'max_depth': None,
 'max_features': 'auto',
 'max_leaf_nodes': None,
 'max_samples': None,
 'min_impurity_decrease': 0.0,
 'min_impurity_split': None,
 'min_samples_leaf': 1,
 'min_samples_split': 2,
 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 100,
 'n_jobs': None,
 'oob_score': False,
 'random_state': None,
 'verbose': 0,
 'warm_start': False}
{'bootstrap': [True, False],
 'max_depth': [10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, None],
 'max_features': ['auto', 'sqrt'],
 'min_samples_leaf': [1, 2, 4],
show more (open the raw output data in a text editor) ...

-----
Accuracy:
0.9629629629629629

```



WINE DATASET

Gaussian Naive Bayes (Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("wine.data", header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of ash', 'Magnesium', 'Total phenols', 'Flavanoids',
```

```
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.naive_bayes import GaussianNB
```

```
classifier = GaussianNB().fit(X_train,y_train)
```

```
classifier.fit(X_train,y_train)
```

```
y_pred=classifier.predict(X_test)
```

```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
from sklearn.metrics import plot_confusion_matrix
```

```
plot_confusion_matrix(classifier, X_test, y_test)
```

```
plt.show()
```

Confusion Matrix:

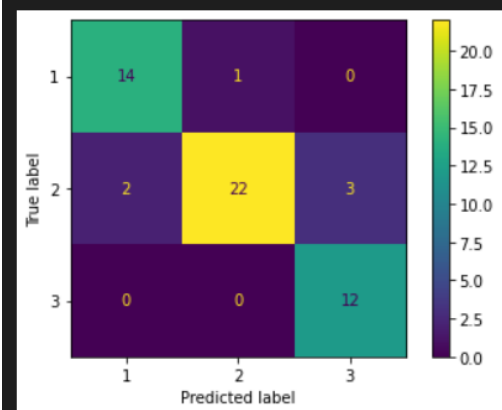
```
[[14  1  0]
 [ 2 22  3]
 [ 0  0 12]]
```

Performance Evaluation

	precision	recall	f1-score	support
1	0.88	0.93	0.90	15
2	0.96	0.81	0.88	27
3	0.80	1.00	0.89	12
accuracy			0.89	54
macro avg	0.88	0.92	0.89	54
weighted avg	0.90	0.89	0.89	54

Accuracy:

0.8888888888888888



WINE DATASET

Gaussian Naive Bayes (With Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv("wine.data",header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of ash','Magnesium','Total
phenols','Flavanoids',

```
    'Nonflavanoid phenols','Proanthocyanins','Color intensity','Hue','OD280/OD315 of  
diluted wines','Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.naive_bayes import GaussianNB
```

```
classifier = GaussianNB(priors=None,var_smoothing=1e-5).fit(X_train,y_train)
```

```
classifier.fit(X_train,y_train)
```

```
y_pred=classifier.predict(X_test)
```

```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
from sklearn.metrics import plot_confusion_matrix
```

```
plot_confusion_matrix(classifier, X_test, y_test)
```

```
plt.show()
```

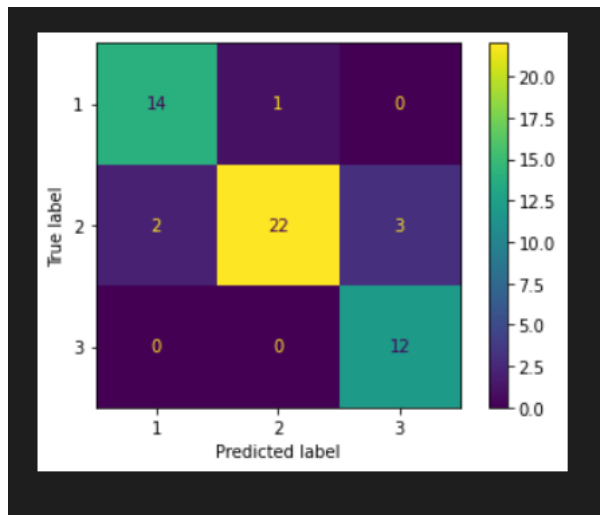
```

Confusion Matrix:
[[14  1  0]
 [ 2 22  3]
 [ 0  0 12]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     1       0.88       0.93       0.90        15
     2       0.96       0.81       0.88        27
     3       0.80       1.00       0.89        12

 accuracy          0.89        54
 macro avg          0.88        54
 weighted avg       0.90        54
-----
Accuracy:
0.8888888888888888

```



IONOSPHERE DATASET

SVM(Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)


# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)


# Classification
from sklearn.svm import SVC

classifier = SVC()
classifier.fit(X_train,y_train)

y_pred = classifier.predict(X_test)


from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")


print("Performance Evaluation")
print(classification_report(y_test, y_pred))

```



```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
from sklearn.metrics import plot_confusion_matrix
```

```
plot_confusion_matrix(classifier, X_test, y_test)
```

```
plt.show()
```

```

Confusion Matrix:
[[34  6]
 [ 0 66]]
-----

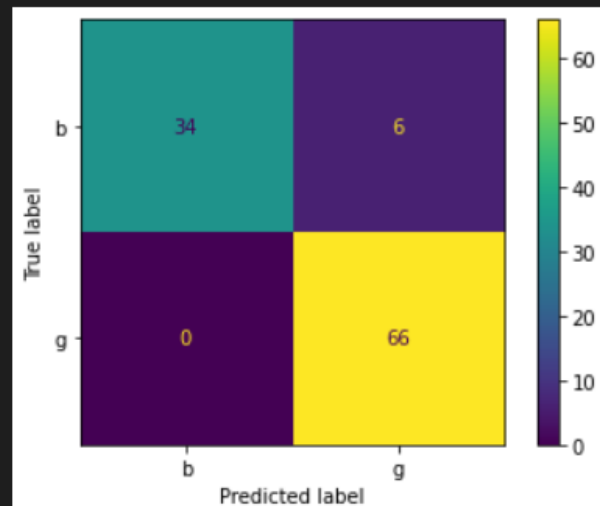
Performance Evaluation
      precision    recall  f1-score   support

     b         1.00      0.85      0.92         40
     g         0.92      1.00      0.96         66

 accuracy          0.94         106
 macro avg         0.96      0.93      0.94         106
weighted avg         0.95      0.94      0.94         106
-----

Accuracy:
0.9433962264150944

```



```
# IONOSPHERE DATASET
```

```
# SVM(With Tuning)[70-30 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.svm import SVC
```

```
classifier = SVC()
```

```
#####
#
# Showing all the parameters

from pprint import pprint
# Look at parameters used by our current forest
print('Parameters currently in use:\n')
pprint(classifier.get_params())

#####
#
# Creating a set of important sample features

param_grid = {'C': [0.1,1, 10, 100], 'gamma': [1,0.1,0.01,0.001], 'kernel': ['rbf', 'poly',
'sigmoid']}

pprint(param_grid)

#####
#

from sklearn.model_selection import GridSearchCV

# Use the random grid to search for best hyperparameters
# First create the base model to tune
classifier = SVC()

# Random search of parameters, using 3 fold cross validation,
# search across 100 different combinations, and use all available cores

rf_random = GridSearchCV(SVC(), param_grid, refit=True, verbose=2)
rf_random.fit(X_train, y_train)
```

```

y_pred = rf_random.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

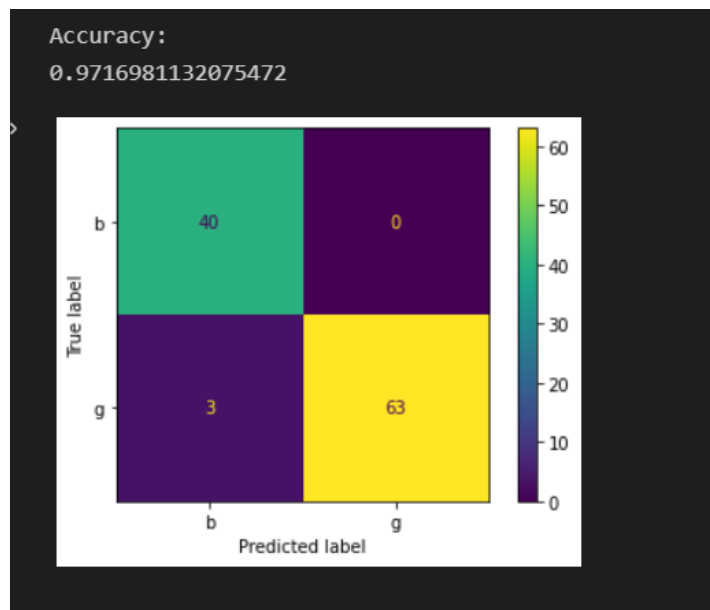
print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(rf_random, X_test, y_test)
plt.show()

```



Ionosphere DATASET

Decision Tree (Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from sklearn.tree import DecisionTreeClassifier

classifier = DecisionTreeClassifier().fit(X_train,y_train)
classifier.fit(X_train,y_train)

y_pred=classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

```

```
print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()

from sklearn import tree
text_representation = tree.export_text(classifier)
print(text_representation)
```



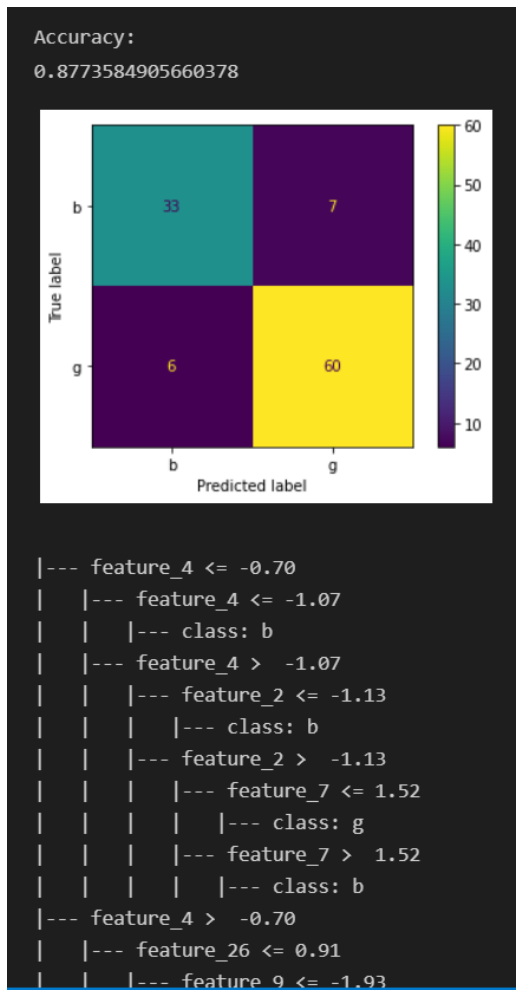
```

Confusion Matrix:
[[33  7]
 [ 6 60]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     b         0.85     0.82     0.84         40
     g         0.90     0.91     0.90         66

 accuracy          0.88         106
 macro avg         0.87     0.87     0.87         106
weighted avg         0.88     0.88     0.88         106
-----
Accuracy:
0.8773584905660378

```



Ionosphere DATASET

Decision Tree (With Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from sklearn.tree import DecisionTreeClassifier
```

```
classifier = DecisionTreeClassifier(criterion="entropy",max_depth=20).fit(X_train,y_train)
```

```
classifier.fit(X_train,y_train)
```

```
y_pred=classifier.predict(X_test)
```

```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
from sklearn.metrics import plot_confusion_matrix
```

```
plot_confusion_matrix(classifier, X_test, y_test)
```

```
plt.show()
```

```
from sklearn import tree
```

```
text_representation = tree.export_text(classifier)
```

```
print(text_representation)
```

Confusion Matrix:

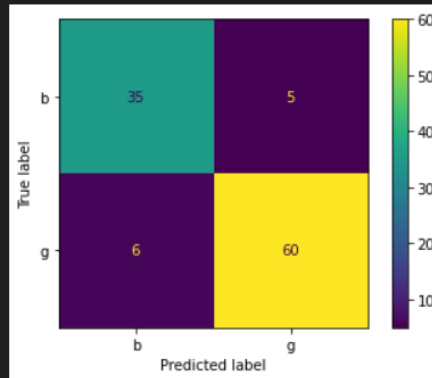
```
[[35  5]
 [ 6 60]]
```

Performance Evaluation

	precision	recall	f1-score	support
b	0.85	0.88	0.86	40
g	0.92	0.91	0.92	66
accuracy			0.90	106
macro avg	0.89	0.89	0.89	106
weighted avg	0.90	0.90	0.90	106

Accuracy:

0.8962264150943396



```
|--- feature_4 <= -1.07
| |--- class: b
|--- feature_4 > -1.07
| |--- feature_26 <= 0.91
| | |--- feature_2 <= 0.13
| | | |--- feature_2 <= -1.13
| | | |--- class: b
| | | |--- feature_2 > -1.13
| | | |--- feature_10 <= 0.63
| | | |--- feature_29 <= -0.58
| | | | |--- class: b
| | | |--- feature_29 > -0.58
| | | | |--- feature_25 <= 0.48
| | | | |--- class: g
```

IONOSPHERE DATASET

Random Forest Classifier(Without Tuning)[70-30 split]

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.30,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from sklearn.ensemble import RandomForestClassifier

classifier = RandomForestClassifier(n_estimators=20, random_state=0)
classifier.fit(X_train,y_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, y_pred))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, y_pred))
```

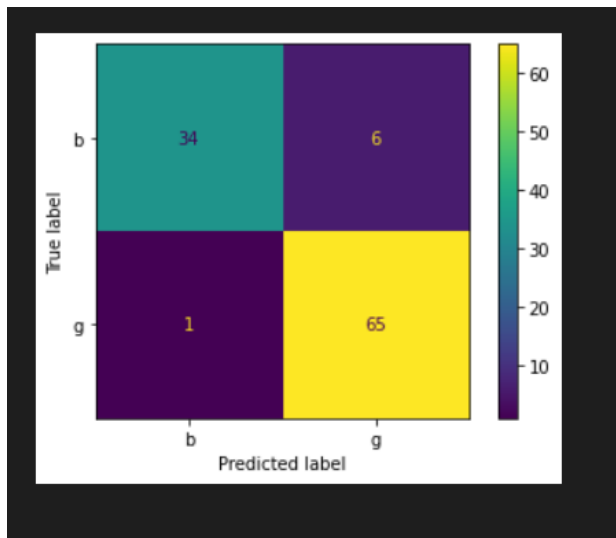
```
import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()
```

```
Confusion Matrix:
[[34  6]
 [ 1 65]]
-----
-----
Performance Evaluation
              precision    recall  f1-score   support

      b         0.97        0.85        0.91         40
      g         0.92        0.98        0.95         66

   accuracy              0.93         106
  macro avg         0.94        0.92        0.93         106
weighted avg         0.94        0.93        0.93         106

-----
-----
Accuracy:
0.9339622641509434
```



IONOSPHERE DATASET

Random Forest Classifier(With Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv("ionosphere.data",header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
, '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['Class'], axis=1)

y = df['Class']

```

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)


# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)


# Classification
from sklearn.ensemble import RandomForestClassifier

classifier = RandomForestClassifier()


#####
#
# Showing all the parameters

from pprint import pprint
# Look at parameters used by our current forest
print('Parameters currently in use:\n')
pprint(classifier.get_params())


#####
#
# Creating a set of important sample features

```



```

from sklearn.model_selection import RandomizedSearchCV

# Number of trees in random forest
n_estimators = [int(x) for x in np.linspace(start = 200, stop = 2000, num = 10)]

# Number of features to consider at every split
max_features = ['auto', 'sqrt']

# Maximum number of levels in tree
max_depth = [int(x) for x in np.linspace(10, 110, num = 11)]
max_depth.append(None)

# Minimum number of samples required to split a node
min_samples_split = [2, 5, 10]

# Minimum number of samples required at each leaf node
min_samples_leaf = [1, 2, 4]

# Method of selecting samples for training each tree
bootstrap = [True, False]

# Create the random grid
random_grid = {'n_estimators': n_estimators,
               'max_features': max_features,
               'max_depth': max_depth,
               'min_samples_split': min_samples_split,
               'min_samples_leaf': min_samples_leaf,
               'bootstrap': bootstrap}

pprint(random_grid)

#####
#

# Use the random grid to search for best hyperparameters
# First create the base model to tune
classifier = RandomForestClassifier()

# Random search of parameters, using 3 fold cross validation,
# search across 100 different combinations, and use all available cores

```

```

rf_random = RandomizedSearchCV(estimator = classifier, param_distributions =
random_grid, n_iter = 100, cv = 3, verbose=2, random_state=42, n_jobs = -1)

# Fit the random search model
rf_random.fit(X_train, y_train)

y_pred = rf_random.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

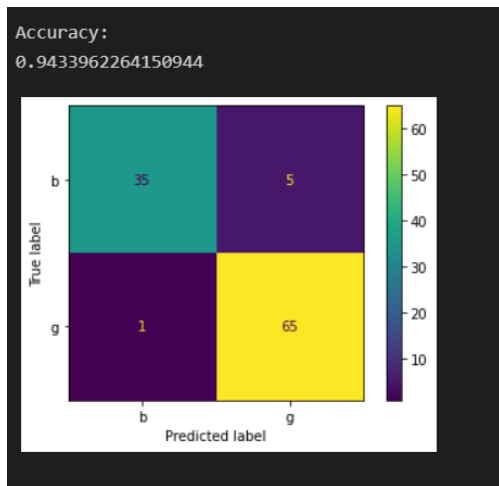
print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(rf_random, X_test, y_test)
plt.show()

```



```
# Ionosphere DATASET
```

```
# Gaussian Naive Bayes (Without Tuning)[70-30 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)


# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)


# Classification
from sklearn.naive_bayes import GaussianNB

classifier = GaussianNB().fit(X_train,y_train)
classifier.fit(X_train,y_train)

y_pred=classifier.predict(X_test)


from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

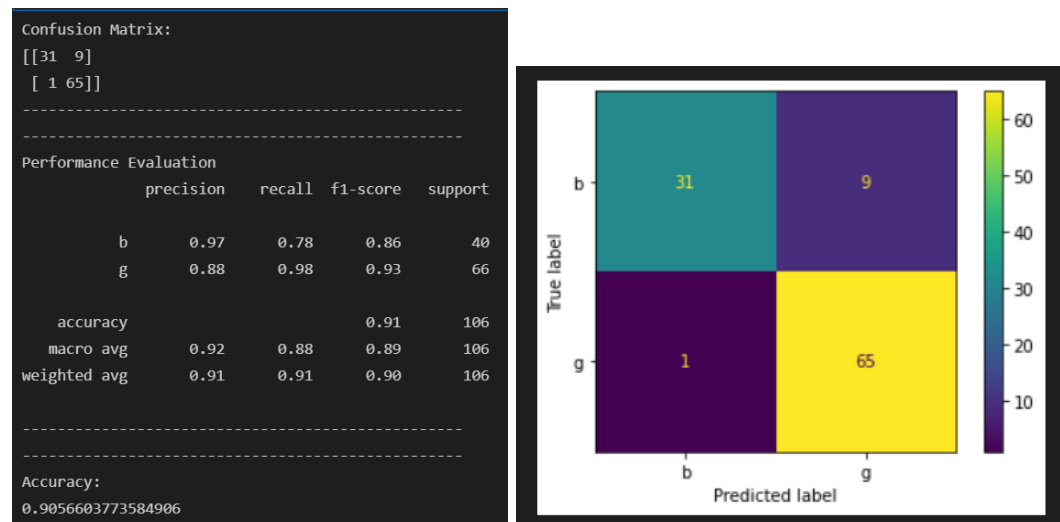

print("Performance Evaluation")
print(classification_report(y_test, y_pred))

```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, y_pred))
```

```
import matplotlib.pyplot as plt
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(classifier, X_test, y_test)
plt.show()
```



Q3. Used datasets

- Iris plants dataset: <https://archive.ics.uci.edu/ml/datasets/Iris/>
- Diabetes dataset: <https://www4.stat.ncsu.edu/~boos/var.select/diabetes.html>

IONOSPHERE DATASET

GaussianHMM(Without Tuning)[70-30 split]

```
print("Split: 70-30")
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
```

```

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")

strings

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, strings))

```

```

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()

```

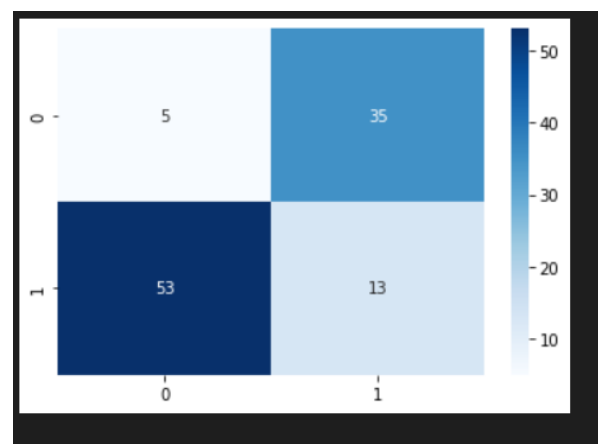
```

Split: 70-30
Confusion Matrix:
[[ 5 35]
 [53 13]]
-----
-----
Performance Evaluation
      precision    recall  f1-score   support

      b         0.09      0.12      0.10         40
      g         0.27      0.20      0.23         66

 accuracy          0.17         106
 macro avg         0.18      0.16      0.17         106
 weighted avg      0.20      0.17      0.18         106
-----
-----
Accuracy:
0.16981132075471697

```




```

# IONOSPHERE DATASET
# GaussianHMM(With Tuning)[70-30 split]

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv("ionosphere.data",header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
            , '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)

```

```

X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2,
covariance_type="full",n_iter=5,algorithm='viterbi',verbose=False)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")

strings

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

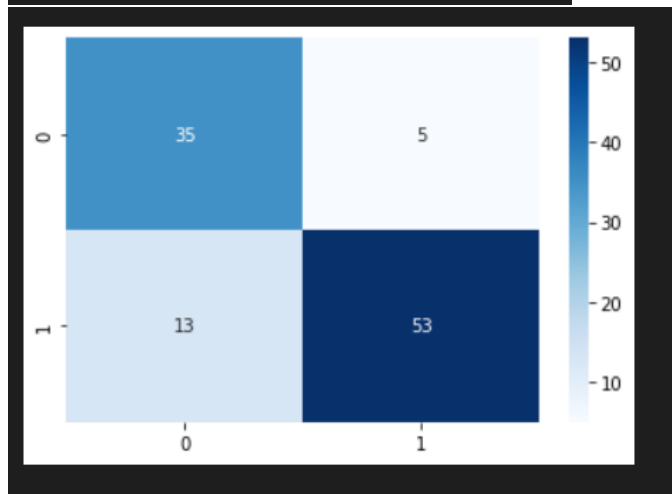
```

Confusion Matrix:
[[35  5]
 [13 53]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     b       0.73       0.88       0.80        40
     g       0.91       0.80       0.85        66

 accuracy              0.83        106
  macro avg       0.82       0.84       0.83        106
 weighted avg       0.84       0.83       0.83        106
-----
Accuracy:
0.8301886792452831

```



```
# IONOSPHERE DATASET
```

```
# GMMHMM(Without Tuning)[60-40 split]
```

```
print("Split: 60-40")
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("ionosphere.data",header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("g")
```

```
    else:
```

```
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```

import matplotlib.pyplot as plt

import seaborn as sns

cm = confusion_matrix(y_test, strings)

sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')

plt.show()

```

```

Split: 60-40
Confusion Matrix:
[[ 7 48]
 [70 16]]
-----

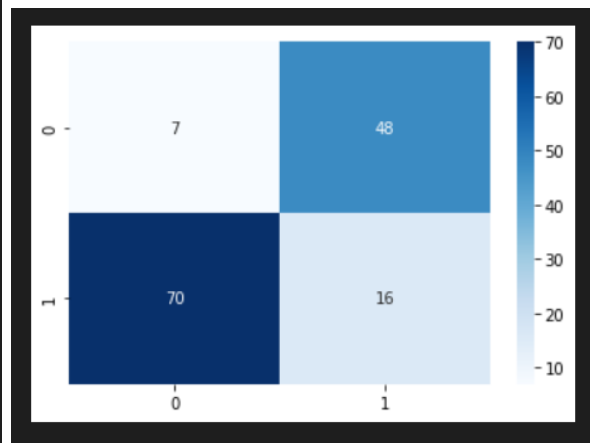
Performance Evaluation
      precision    recall  f1-score   support

     b         0.09     0.13     0.11         55
     g         0.25     0.19     0.21         86

 accuracy          0.16         141
 macro avg         0.17     0.16     0.16         141
 weighted avg      0.19     0.16     0.17         141
-----

Accuracy:
0.16312056737588654

```



IONOSPHERE DATASET

GMMHMM(With Tuning)[60-40 split]

```
print("Split size: 60-40")
```

```
import pandas as pd
```

```
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv("ionosphere.data", header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):

```



```

    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")

strings

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, strings))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns

```

```

cm = confusion_matrix(y_test, strings)

sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')

plt.show()

```

```

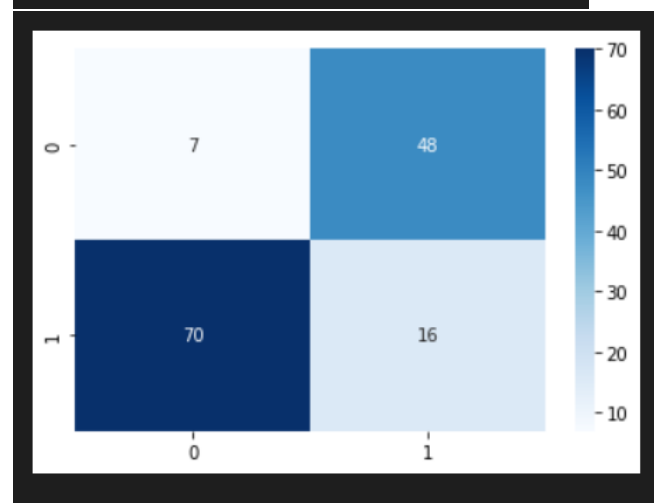
Split size: 60-40
Confusion Matrix:
[[ 7 48]
 [70 16]]
-----
-----
Performance Evaluation
      precision    recall  f1-score   support

     b         0.09      0.13      0.11         55
     g         0.25      0.19      0.21         86

 accuracy          0.16         141
 macro avg         0.17      0.16      0.16         141
weighted avg         0.19      0.16      0.17         141

-----
-----
Accuracy:
0.16312056737588654

```



```
# IONOSPHERE DATASET
```

```
# MultinomialHMM(Without Tuning)[40-60 split]
```

```
print("Split: 40-60")
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm  
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math  
row = len(X_train)  
col = len(X_train[0])  
new = [1] * 33  
for i in range(row):  
    for j in range(col):  
        X_train[i][j] = X_train[i][j]*10  
        X_train[i][j] = math.floor(X_train[i][j])  
    x = X_train[i].astype(int)  
    new = np.vstack([new,x])  
  
y = new  
y = np.absolute(y)  
X_train = y
```

```
import math  
row = len(X_test)  
col = len(X_test[0])  
new  
for i in range(row):  
    for j in range(col):  
        X_test[i][j] = X_test[i][j]*10  
        X_test[i][j] = math.floor(X_test[i][j])  
    x = X_test[i].astype(int)  
    new = np.vstack([new,x])  
  
y = new
```

```

y = np.absolute(y)
X_test = y

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("b")
    else:
        strings[i] = ("g")

strings
strings = strings[0:211]

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```

Split: 40-60
Confusion Matrix:
[[ 25  51]
 [ 21 114]]
-----

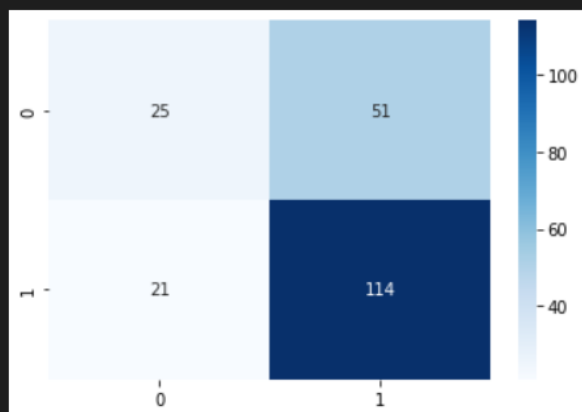
Performance Evaluation
      precision    recall  f1-score   support

     b         0.54         0.33         0.41         76
     g         0.69         0.84         0.76        135

 accuracy          0.66         0.66        211
 macro avg         0.62         0.59         0.58        211
weighted avg         0.64         0.66         0.63        211
-----

Accuracy:
0.6587677725118484

```



```
# Irish DATASET
```

```
# GaussianHMM(Without Tuning)[30-70 split]
```

```
print("Split: 30-70")
```

```
import pandas as pd
```

```
import numpy as np
```

```
import sklearn
```

```
from sklearn import datasets
```

```
iris = datasets.load_iris()
```

```

X=iris.data
y=iris.target

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

```



```
print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```

Split: 30-70
Confusion Matrix:
[[33  0  0]
 [ 0 36  0]
 [ 0 36  0]]
-----

Performance Evaluation
      precision    recall  f1-score   support

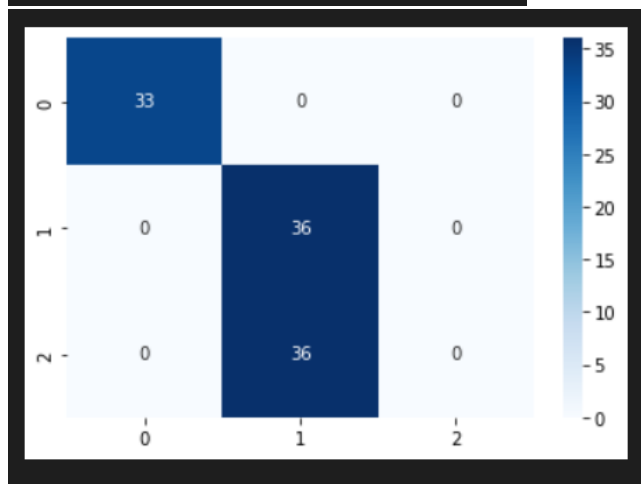
      0         1.00      1.00      1.00        33
      1         0.50      1.00      0.67        36
      2         0.00      0.00      0.00        36

 accuracy         0.66      0.66      0.66       105
 macro avg         0.50      0.67      0.56       105
weighted avg         0.49      0.66      0.54       105

-----

Accuracy:
0.6571428571428571

```



```
# Irish DATASET
```

```
# GaussianHMM(With Tuning)[70-30 split]
```

```
print("Split: 70-30")
```

```
import pandas as pd
```

```
import numpy as np
```

```
import sklearn
```

```
from sklearn import datasets
```

```
iris = datasets.load_iris()
```

```

X=iris.data
y=iris.target

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2,
covariance_type="full",n_iter=5,algorithm='viterbi',verbose=False)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")

```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, y_pred))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, y_pred))
```

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

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, y_pred)
```

```
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

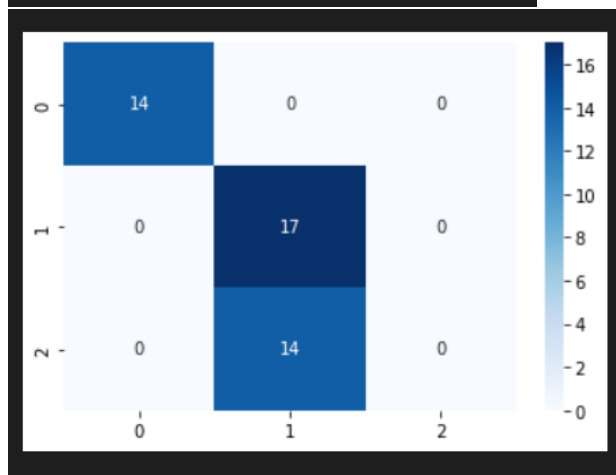
```

Split: 70-30
Confusion Matrix:
[[14  0  0]
 [ 0 17  0]
 [ 0 14  0]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     0       1.00      1.00      1.00        14
     1       0.55      1.00      0.71        17
     2       0.00      0.00      0.00        14

 accuracy          0.69      0.69      0.69      45
 macro avg          0.52      0.67      0.57      45
weighted avg          0.52      0.69      0.58      45
-----
Accuracy:
0.6888888888888889

```



Irish DATASET

GMMHMM(With Tuning)[30-70 split]

```
print("Split: 30-70")
```

```
import pandas as pd
```

```
import numpy as np
```

```
import sklearn
```

```
from sklearn import datasets
```

```
iris = datasets.load_iris()
```

```

X=iris.data
y=iris.target

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

print("-----")
print("-----")

```

```
print("Performance Evaluation")
print(classification_report(y_test, y_pred))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, y_pred))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

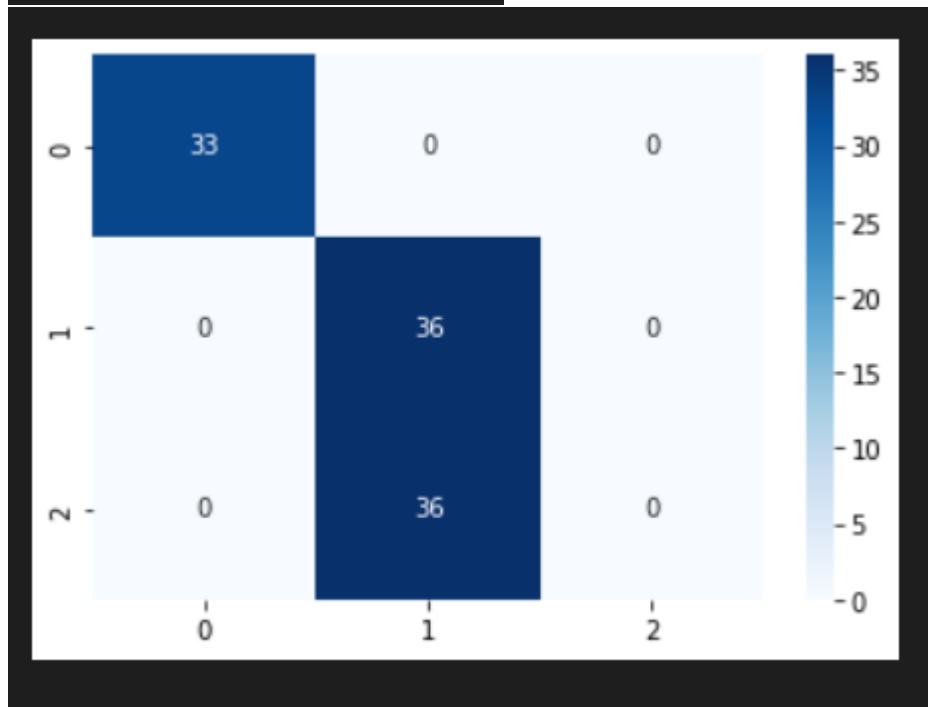
```

Split: 30-70
Confusion Matrix:
[[33  0  0]
 [ 0 36  0]
 [ 0 36  0]]
-----
Performance Evaluation
      precision    recall  f1-score   support

     0       1.00      1.00      1.00        33
     1       0.50      1.00      0.67        36
     2       0.00      0.00      0.00        36

 accuracy          0.66        105
 macro avg       0.50      0.67      0.56        105
weighted avg       0.49      0.66      0.54        105
-----
Accuracy:
0.6571428571428571

```



Q5. Used Dataset

Wine Dataset: <https://archive.ics.uci.edu/ml/datasets/wine>

Kmeans vs Kmedoids

#importing libraries

import numpy as np

import pandas as pd

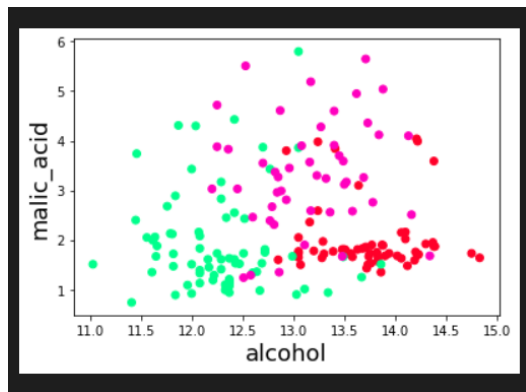
import sklearn as sk


```

import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.cluster import KMeans
from sklearn.datasets import load_wine
wine=load_wine() #loading iris dataset from sklearn.datasets
x=wine.data
df=pd.DataFrame(data=wine.data, columns=['alcohol',
    'malic_acid',
    'ash',
    'alcalinity_of_ash',
    'magnesium',
    'total_phenols',
    'flavanoids',
    'nonflavanoid_phenols',
    'proanthocyanins',
    'color_intensity',
    'hue',
    'od280/od315_of_diluted_wines',
    'proline'])
plt.scatter(x=df['alcohol'], y=df['malic_acid'],c=wine.target,
    cmap='gist_rainbow') #try using cmap='rainbow'

plt.xlabel('alcohol', fontsize=18)
plt.ylabel('malic_acid', fontsize=18)

```



```
kmeans = KMeans(init="random", n_clusters=3, n_init=10, max_iter=300,
random_state=42)
```

```
y = kmeans.fit_predict(x)
```

```
print("K-Means Cluster Centers")
```

```
print(kmeans.cluster_centers_)
```

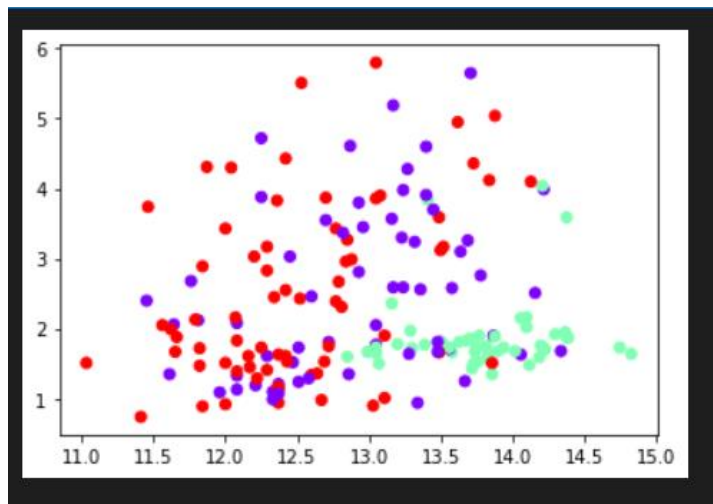
```
print("Cluster Labels")
```

```
print(kmeans.labels_)
```

```
K-Means Cluster Centers
[[1.29298387e+01 2.50403226e+00 2.40806452e+00 1.98903226e+01
 1.03596774e+02 2.11112903e+00 1.58403226e+00 3.88387097e-01
 1.50338710e+00 5.65032258e+00 8.83967742e-01 2.36548387e+00
 7.28338710e+02]
 [1.38044681e+01 1.88340426e+00 2.42617021e+00 1.70234043e+01
 1.05510638e+02 2.86723404e+00 3.01425532e+00 2.85319149e-01
 1.91042553e+00 5.70255319e+00 1.07829787e+00 3.11404255e+00
 1.19514894e+03]
 [1.25166667e+01 2.49420290e+00 2.28855072e+00 2.08231884e+01
 9.23478261e+01 2.07072464e+00 1.75840580e+00 3.90144928e-01
 1.45188406e+00 4.08695651e+00 9.41159420e-01 2.49072464e+00
 4.58231884e+02]]
Cluster Labels
[1 1 1 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 0 0 0 1 1 0 0 1 1 1 1 1 1 0 0
 1 1 0 0 1 1 0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 2 0 2 0 2 2 0 2 2 0 0 0 2 2 1
 0 2 2 2 0 2 2 0 0 2 2 2 2 2 0 0 2 2 2 2 0 0 2 0 2 0 2 2 2 0 2 2 2 0 2
 2 0 2 2 2 2 2 0 2 2 2 2 2 2 2 2 0 2 2 0 0 0 2 2 2 0 0 2 2 0 0 2 0
 0 2 2 2 2 0 0 0 2 0 0 0 2 0 2 0 0 2 0 0 0 2 2 0 0 0 2 2 0 0 2 2]
```

```
plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=kmeans.labels_,
cmap='rainbow') #try using cmap='rainbow'
```

```
plt.show()
```



```
fig, axes = plt.subplots(1, 2, figsize=(14,6))

axes[0].scatter(x=df['alcohol'], y=df['malic_acid'], c=y,
               cmap='rainbow',edgecolor='k', s=150) #you can also try cmap='rainbow'

axes[1].scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target,
               cmap='jet',edgecolor='k', s=150)

axes[0].set_xlabel('alcohol', fontsize=18)

axes[0].set_ylabel('malic_acid', fontsize=18)

axes[1].set_xlabel('alcohol', fontsize=18)

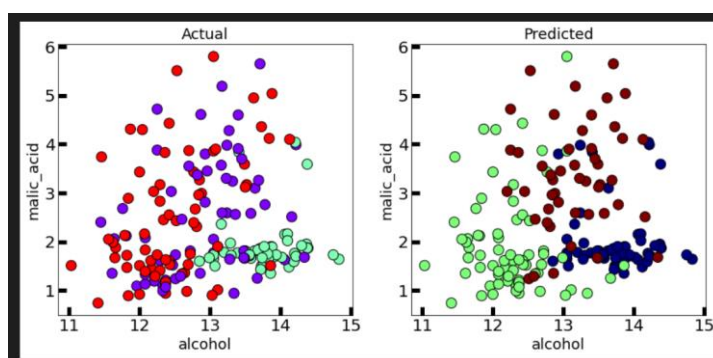
axes[1].set_ylabel('malic_acid', fontsize=18)

axes[0].tick_params(direction='in', length=10, width=5, colors='k',
                    labels=20)

axes[1].tick_params(direction='in', length=10, width=5, colors='k',
                    labels=20)

axes[0].set_title('Actual', fontsize=18)

axes[1].set_title('Predicted', fontsize=18)
```



```
from sklearn.metrics import silhouette_score
print("The silhouette score is :")
silhouette_score(x, kmeans.labels_)
```

```
The silhouette score is :
0.5711381937868838
```

```
from sklearn.metrics import calinski_harabasz_score
print("The calinski harabasz score is :")
calinski_harabasz_score(x, kmeans.labels_)
```

```
The calinski harabasz score is :
561.815657860671
```

```
from sklearn.metrics import davies_bouldin_score
print("The davies bouldin score is :")
davies_bouldin_score(x, kmeans.labels_)
```

```
The davies bouldin score is :
0.5342431775436286
```

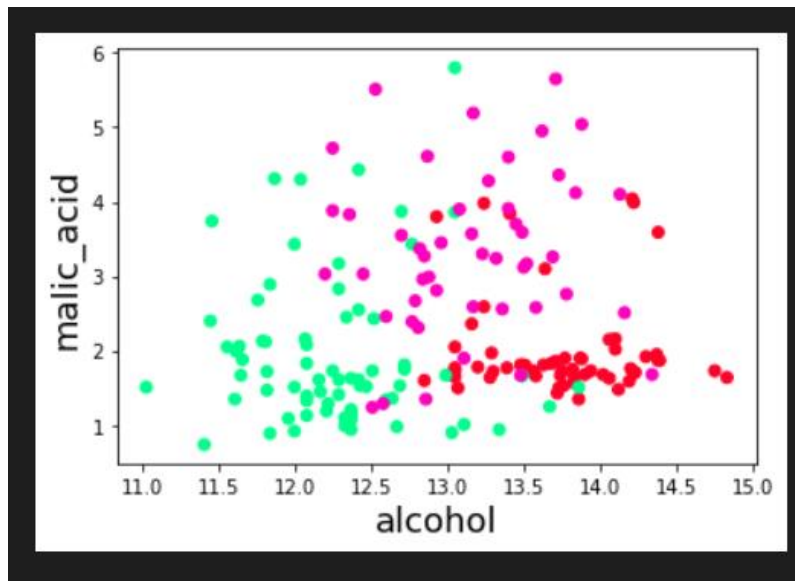
```
#importing libraries
import numpy as np
import pandas as pd
import sklearn as sk
import matplotlib.pyplot as plt
%matplotlib inline

from sklearn_extra.cluster import KMedoids
from sklearn.datasets import load_wine
wine=load_wine() #loading iris dataset from sklearn.datasets
```

```
x=wine.data
df=pd.DataFrame(data=wine.data, columns=['alcohol',
    'malic_acid',
    'ash',
    'alcalinity_of_ash',
    'magnesium',
    'total_phenols',
    'flavanoids',
    'nonflavanoid_phenols',
    'proanthocyanins',
    'color_intensity',
    'hue',
    'od280/od315_of_diluted_wines',
    'proline'])

plt.scatter(x=df['alcohol'], y=df['malic_acid'],c=wine.target,
cmap='gist_rainbow') #try using cmap='rainbow'

plt.xlabel('alcohol', fontsize=18)
plt.ylabel('malic_acid', fontsize=18)
```



```
kmedoid = KMedoids(init="heuristic", n_clusters=3, max_iter=300,
random_state=42)
```

```
y = kmedoid.fit_predict(x)
```

```
print("K-Medoids Cluster Centers")
```

```
print(kmedoid.cluster_centers_)
```

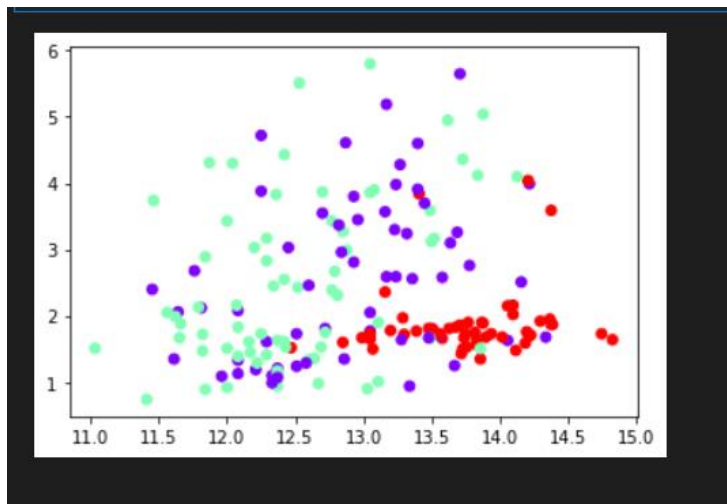
```
print("Cluster Labels")
```

```
print(kmedoid.labels_)
```

```
K-Medoids Cluster Centers
[[1.260e+01 2.460e+00 2.200e+00 1.850e+01 9.400e+01 1.620e+00 6.600e-01
 6.300e-01 9.400e-01 7.100e+00 7.300e-01 1.580e+00 6.950e+02]
 [1.349e+01 1.660e+00 2.240e+00 2.400e+01 8.700e+01 1.880e+00 1.840e+00
 2.700e-01 1.030e+00 3.740e+00 9.800e-01 2.780e+00 4.720e+02]
 [1.383e+01 1.570e+00 2.620e+00 2.000e+01 1.150e+02 2.950e+00 3.400e+00
 4.000e-01 1.720e+00 6.600e+00 1.130e+00 2.570e+00 1.130e+03]]
Cluster Labels
[2 2 2 2 0 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 0 0 0 2 2 0 0 2 2 2 2 2 2 2 2 0
 2 2 0 0 2 2 0 0 2 2 2 2 2 2 2 2 2 2 2 2 2 2 1 0 1 0 1 1 0 1 1 0 0 0 1 1 2
 0 1 1 1 0 1 1 0 0 1 1 1 1 1 0 0 1 1 1 1 2 0 1 0 1 0 1 1 1 0 1 1 1 1 0 1
 1 0 1 1 1 1 1 1 1 0 1 1 1 1 1 1 1 1 1 0 1 1 0 0 0 0 1 1 0 0 0 1 1 0 0 1 0
 0 1 1 1 1 0 0 0 1 0 0 0 1 0 1 0 0 1 0 0 0 0 1 1 0 0 0 0 0 1]
```

```
plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=kmedoid.labels_, cmap='rainbow') #try using
cmap='rainbow'
```

```
plt.show()
```



```
fig, axes = plt.subplots(1, 2, figsize=(14,6))
```

```
axes[0].scatter(x=df['alcohol'], y=df['malic_acid'], c=y, cmap='rainbow',edgecolor='k', s=150)
#you can also try cmap='rainbow'
```

```
axes[1].scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target, cmap='jet',edgecolor='k',
s=150)
```

```
axes[0].set_xlabel('alcohol', fontsize=18)
```

```
axes[0].set_ylabel('malic_acid', fontsize=18)
```

```
axes[1].set_xlabel('alcohol', fontsize=18)
```

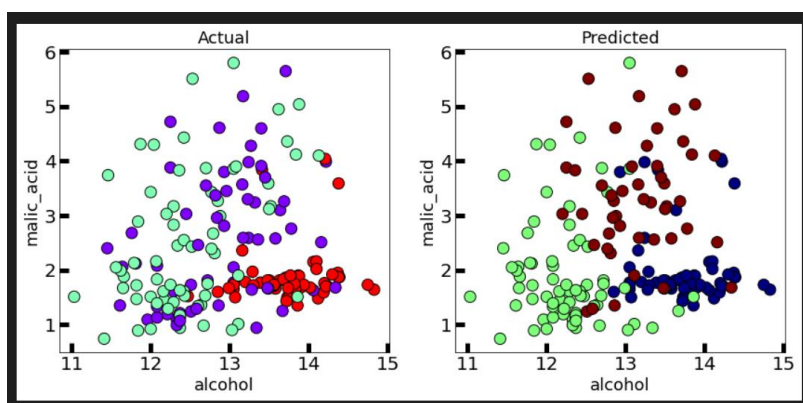
```
axes[1].set_ylabel('malic_acid', fontsize=18)
```

```
axes[0].tick_params(direction='in', length=10, width=5, colors='k', labelsiz=20)
```

```
axes[1].tick_params(direction='in', length=10, width=5, colors='k', labelsiz=20)
```

```
axes[0].set_title('Actual', fontsize=18)
```

```
axes[1].set_title('Predicted', fontsize=18)
```



```
from sklearn.metrics import silhouette_score
print("The silhouette score is :")
```

```
silhouette_score(x, kmedoid.labels_)
```

The silhouette score is :

0.566648040863657

```
from sklearn.metrics import calinski_harabasz_score
print("The calinski harabasz score is :")
calinski_harabasz_score(x, kmedoid.labels_)
```

The calinski harabasz score is :

539.3792353535451

```
from sklearn.metrics import davies_bouldin_score
print("The davies bouldin score is :")
davies_bouldin_score(x, kmedoid.labels_)
```

The davies bouldin score is :

0.5292394126003174

```
#importing libraries
import numpy as np
import pandas as pd
import sklearn as sk
import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.cluster import KMeans
from sklearn.datasets import load_wine

wine=load_wine() #loading iris dataset from sklearn.datasets
x=wine.data

df=pd.DataFrame(data=wine.data, columns=['alcohol',
    'malic_acid',
    'ash',
    'alcalinity_of_ash',
    'magnesium',
    'total_phenols',
    'flavanoids',
    'nonflavanoid_phenols',
    'proanthocyanins',
    'color_intensity',
    'hue',
    'od280/od315_of_diluted_wines',
```



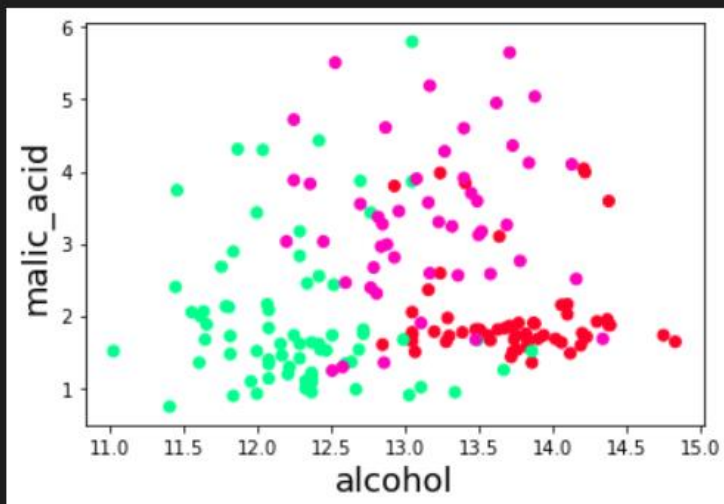
```

        'proline'])

plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target,
            cmap='gist_rainbow') #try using cmap='rainbow'

plt.xlabel('alcohol', fontsize=18)
plt.ylabel('malic_acid', fontsize=18)

```



```

from sklearn.cluster import DBSCAN

dbscan = DBSCAN(eps=35, algorithm='auto', metric='euclidean')
y = dbscan.fit_predict(x)

print("Cluster Labels")
print(dbscan.labels_)

```

```

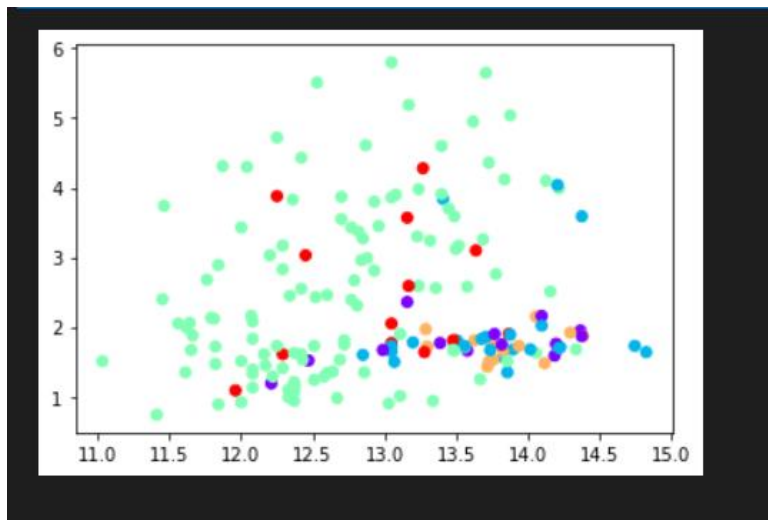
Cluster Labels
[ 0  0 -1 -1  1 -1  2  2  0  0 -1  2  2  0 -1  2  2  0 -1  3  1  1  0  0
  3  3 -1  2  3  0  2 -1  0  2  0  3  3  0  0  1  1  0  0  1  3  0  0  0
  0  2  0  2 -1 -1  0  0  0  2  2  1  1  1  1  1  1  1  1  1  1 -1  3  1
  1 -1  3  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1 -1
  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1
  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1  1
  3  3  1  1  1  1  1  1  1  1  1  1  1  3  1  1  1  1  1  1  1  1  1
  1  1  1  1  1  1  1  3  3  1]

```

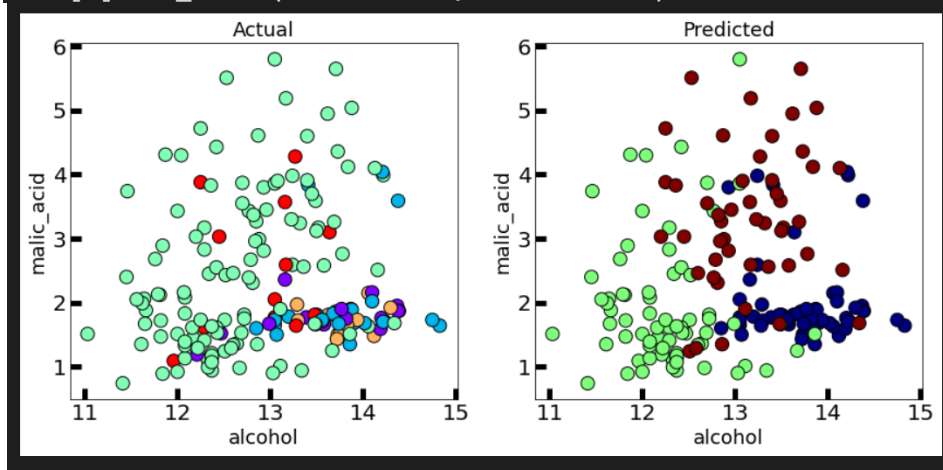
```

plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=dbscan.labels_,
            cmap='rainbow') #try using cmap='rainbow'
plt.show()

```



```
fig, axes = plt.subplots(1, 2, figsize=(14,6))
axes[0].scatter(x=df['alcohol'], y=df['malic_acid'], c=y,
               cmap='rainbow',edgecolor='k', s=150) #you can also try cmap='rainbow'
axes[1].scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target,
               cmap='jet',edgecolor='k', s=150)
axes[0].set_xlabel('alcohol', fontsize=18)
axes[0].set_ylabel('malic_acid', fontsize=18)
axes[1].set_xlabel('alcohol', fontsize=18)
axes[1].set_ylabel('malic_acid', fontsize=18)
axes[0].tick_params(direction='in', length=10, width=5, colors='k',
                    labels=20)
axes[1].tick_params(direction='in', length=10, width=5, colors='k',
                    labels=20)
axes[0].set_title('Actual', fontsize=18)
axes[1].set_title('Predicted', fontsize=18)
```



```
from sklearn.metrics import silhouette_score
print("The silhouette score is :")
silhouette_score(x, dbscan.labels_)
```

```
The silhouette score is :  
0.4413295944891921
```

```
from sklearn.metrics import calinski_harabasz_score  
print("The calinski harabasz score is :")  
calinski_harabasz_score(x, dbscan.labels_)
```

```
The calinski harabasz score is :  
208.9449395725058
```

```
from sklearn.metrics import davies_bouldin_score  
print("The davies bouldin score is :")  
davies_bouldin_score(x, dbscan.labels_)
```

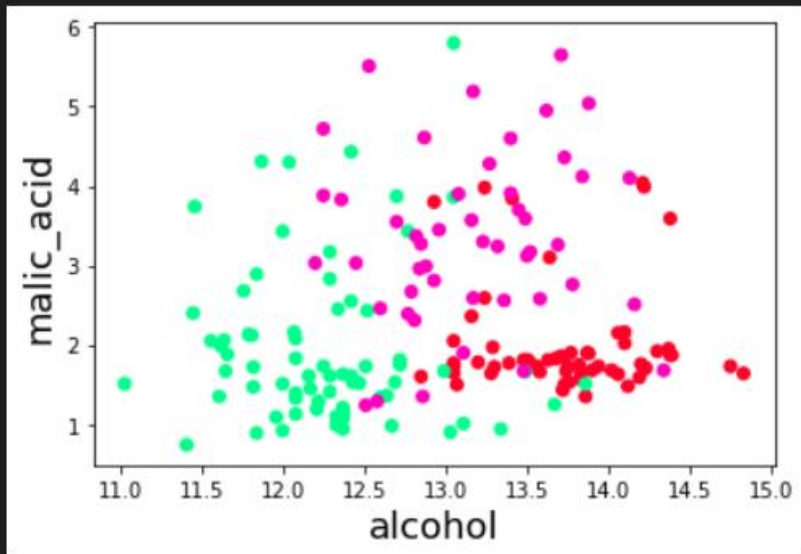
```
The davies bouldin score is :  
7.812129203046089
```

```
#importing libraries  
import numpy as np  
import pandas as pd  
import sklearn as sk  
import matplotlib.pyplot as plt  
%matplotlib inline  
from sklearn.cluster import KMeans  
from sklearn.datasets import load_wine  
  
wine=load_wine() #loading iris dataset from sklearn.datasets  
x=wine.data  
  
df=pd.DataFrame(data=wine.data, columns=['alcohol',  
    'malic_acid',  
    'ash',  
    'alcalinity_of_ash',  
    'magnesium',  
    'total_phenols',  
    'flavanoids',  
    'nonflavanoid_phenols',  
    'proanthocyanins',  
    'color_intensity',  
    'hue',
```

```
'od280/od315_of_diluted_wines',
'proline']])
```

```
plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target,
cmap='gist_rainbow') #try using cmap='rainbow'
```

```
plt.xlabel('alcohol', fontsize=18)
plt.ylabel('malic_acid', fontsize=18)
Text(0, 0.5, 'malic_acid')
```



```
from sklearn.cluster import OPTICS
```

```
optics = OPTICS(min_samples=13)
y = optics.fit_predict(x)
```

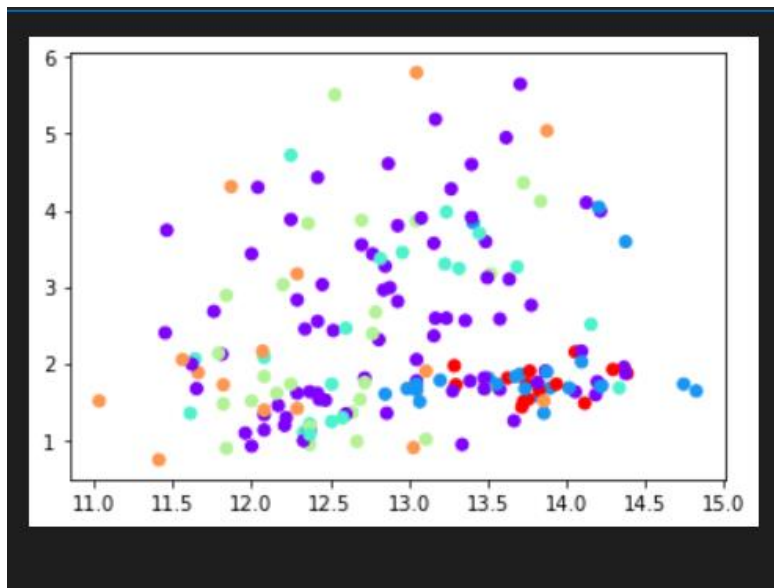
```
print("Cluster Labels")
print(optics.labels_)
```

```
Cluster Labels
```

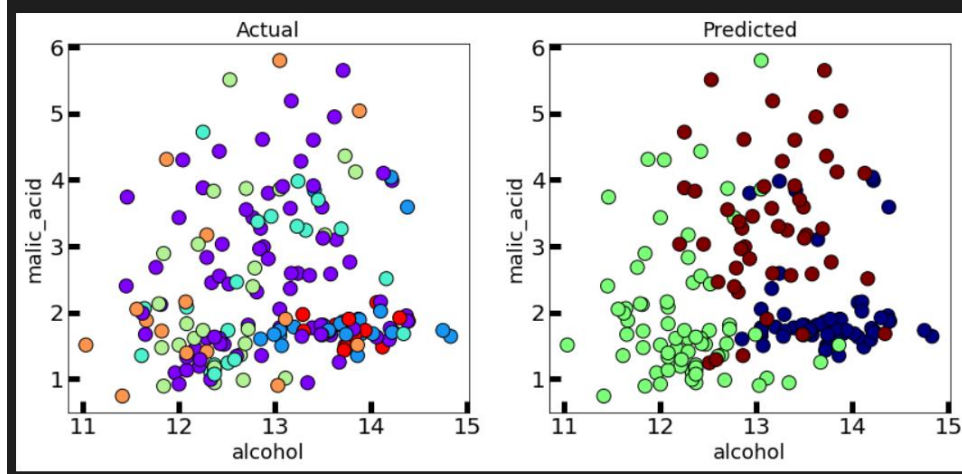
```
[ 0  0 -1 -1 -1 -1  4  4  0  0 -1  4  4  0 -1  4  4  0 -1 -1 -1 -1  0  0
-1 -1 -1  4 -1  0  4 -1  0  4  0 -1 -1  0  0 -1 -1  0  0  1 -1  0  0  0
 0  4  0  4 -1  4  0  0  0  4  4  2  1  2 -1 -1 -1  1  2  2 -1 -1 -1  3
 2  0 -1  3  3  2 -1  2 -1 -1 -1  2  2  2  2 -1  1 -1  2  2  2 -1 -1 -1
-1  3  1  3  1 -1 -1  3  1 -1  2  2 -1  1 -1 -1 -1  3  3  3  2 -1 -1 -1
-1  3 -1  3  3  3 -1  2 -1 -1 -1 -1 -1 -1  1  1  1  2 -1 -1 -1 -1  2 -1
-1 -1  3 -1  1 -1 -1  2  3  1  1 -1  2 -1  1 -1  2  1 -1  1 -1  2  1  1
-1 -1  2  2  1 -1 -1 -1 -1 -1]
```

[+ Code](#)

```
plt.scatter(x=df['alcohol'], y=df['malic_acid'], c=optics.labels_,
cmap='rainbow') #try using cmap='rainbow'
plt.show()
```



```
fig, axes = plt.subplots(1, 2, figsize=(14,6))
axes[0].scatter(x=df['alcohol'], y=df['malic_acid'], c=y,
cmap='rainbow',edgecolor='k', s=150) #you can also try cmap='rainbow'
axes[1].scatter(x=df['alcohol'], y=df['malic_acid'], c=wine.target,
cmap='jet',edgecolor='k', s=150)
axes[0].set_xlabel('alcohol', fontsize=18)
axes[0].set_ylabel('malic_acid', fontsize=18)
axes[1].set_xlabel('alcohol', fontsize=18)
axes[1].set_ylabel('malic_acid', fontsize=18)
axes[0].tick_params(direction='in', length=10, width=5, colors='k',
labels=20)
axes[1].tick_params(direction='in', length=10, width=5, colors='k',
labels=20)
axes[0].set_title('Actual', fontsize=18)
axes[1].set_title('Predicted', fontsize=18)
```



```
from sklearn.metrics import silhouette_score
print("The silhouette score is :")
silhouette_score(x, optics.labels_)
```

```
The silhouette score is :  
0.08708464008252535
```

```
from sklearn.metrics import calinski_harabasz_score  
print("The calinski harabasz score is :")  
calinski_harabasz_score(x, optics.labels_)
```

```
The calinski harabasz score is :  
44.955612959266325
```

```
from sklearn.metrics import davies_bouldin_score  
print("The davies bouldin score is :")  
davies_bouldin_score(x, optics.labels_)
```

```
The davies bouldin score is :  
2.275571431762288
```