

Machine Learning Tutorial With Multipler Classifiers

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```
[1]: import pandas as pd
import numpy as np
```

```
[2]: iris=pd.read_csv(r"Iris.csv")
```

```
[3]: iris
iris.head()
```

```
[3]:      150      4  setosa  versicolor  disorder
0  5.1  3.5    1.4         0.2    Normal
1  4.9  3.0    1.4         0.2    Normal
2  4.7  3.2    1.3         0.2    Normal
3  4.6  3.1    1.5         0.2    Normal
4  5.0  3.6    1.4         0.2    Normal
```

```
[4]: X = iris.iloc[:, 0:4].values
X
```

```
[4]: array([[5.1, 3.5, 1.4, 0.2],
          [4.9, 3. , 1.4, 0.2],
          [4.7, 3.2, 1.3, 0.2],
          [4.6, 3.1, 1.5, 0.2],
          [5. , 3.6, 1.4, 0.2],
          [5.4, 3.9, 1.7, 0.4],
          [4.6, 3.4, 1.4, 0.3],
          [5. , 3.4, 1.5, 0.2],
          [4.4, 2.9, 1.4, 0.2],
          [4.9, 3.1, 1.5, 0.1],
          [5.4, 3.7, 1.5, 0.2],
          [4.8, 3.4, 1.6, 0.2],
          [4.8, 3. , 1.4, 0.1],
          [4.3, 3. , 1.1, 0.1],
          [5.8, 4. , 1.2, 0.2],
          [5.7, 4.4, 1.5, 0.4],
          [5.4, 3.9, 1.3, 0.4],
          [5.1, 3.5, 1.4, 0.3],
          [5.7, 3.8, 1.7, 0.3],
```

[5.1, 3.8, 1.5, 0.3],
[5.4, 3.4, 1.7, 0.2],
[5.1, 3.7, 1.5, 0.4],
[4.6, 3.6, 1. , 0.2],
[5.1, 3.3, 1.7, 0.5],
[4.8, 3.4, 1.9, 0.2],
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[5. , 3.4, 1.6, 0.4],
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[5.5, 4.2, 1.4, 0.2],
[4.9, 3.1, 1.5, 0.2],
[5. , 3.2, 1.2, 0.2],
[5.5, 3.5, 1.3, 0.2],
[4.9, 3.6, 1.4, 0.1],
[4.4, 3. , 1.3, 0.2],
[5.1, 3.4, 1.5, 0.2],
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[5.1, 3.8, 1.9, 0.4],
[4.8, 3. , 1.4, 0.3],
[5.1, 3.8, 1.6, 0.2],
[4.6, 3.2, 1.4, 0.2],
[5.3, 3.7, 1.5, 0.2],
[5. , 3.3, 1.4, 0.2],
[7. , 3.2, 4.7, 1.4],
[6.4, 3.2, 4.5, 1.5],
[6.9, 3.1, 4.9, 1.5],
[5.5, 2.3, 4. , 1.3],
[6.5, 2.8, 4.6, 1.5],
[5.7, 2.8, 4.5, 1.3],
[6.3, 3.3, 4.7, 1.6],
[4.9, 2.4, 3.3, 1.],
[6.6, 2.9, 4.6, 1.3],
[5.2, 2.7, 3.9, 1.4],
[5. , 2. , 3.5, 1.],
[5.9, 3. , 4.2, 1.5],
[6. , 2.2, 4. , 1.],
[6.1, 2.9, 4.7, 1.4],
[5.6, 2.9, 3.6, 1.3],
[6.7, 3.1, 4.4, 1.4],

[5.6, 3. , 4.5, 1.5],
 [5.8, 2.7, 4.1, 1.],
 [6.2, 2.2, 4.5, 1.5],
 [5.6, 2.5, 3.9, 1.1],
 [5.9, 3.2, 4.8, 1.8],
 [6.1, 2.8, 4. , 1.3],
 [6.3, 2.5, 4.9, 1.5],
 [6.1, 2.8, 4.7, 1.2],
 [6.4, 2.9, 4.3, 1.3],
 [6.6, 3. , 4.4, 1.4],
 [6.8, 2.8, 4.8, 1.4],
 [6.7, 3. , 5. , 1.7],
 [6. , 2.9, 4.5, 1.5],
 [5.7, 2.6, 3.5, 1.],
 [5.5, 2.4, 3.8, 1.1],
 [5.5, 2.4, 3.7, 1.],
 [5.8, 2.7, 3.9, 1.2],
 [6. , 2.7, 5.1, 1.6],
 [5.4, 3. , 4.5, 1.5],
 [6. , 3.4, 4.5, 1.6],
 [6.7, 3.1, 4.7, 1.5],
 [6.3, 2.3, 4.4, 1.3],
 [5.6, 3. , 4.1, 1.3],
 [5.5, 2.5, 4. , 1.3],
 [5.5, 2.6, 4.4, 1.2],
 [6.1, 3. , 4.6, 1.4],
 [5.8, 2.6, 4. , 1.2],
 [5. , 2.3, 3.3, 1.],
 [5.6, 2.7, 4.2, 1.3],
 [5.7, 3. , 4.2, 1.2],
 [5.7, 2.9, 4.2, 1.3],
 [6.2, 2.9, 4.3, 1.3],
 [5.1, 2.5, 3. , 1.1],
 [5.7, 2.8, 4.1, 1.3],
 [6.3, 3.3, 6. , 2.5],
 [5.8, 2.7, 5.1, 1.9],
 [7.1, 3. , 5.9, 2.1],
 [6.3, 2.9, 5.6, 1.8],
 [6.5, 3. , 5.8, 2.2],
 [7.6, 3. , 6.6, 2.1],
 [4.9, 2.5, 4.5, 1.7],
 [7.3, 2.9, 6.3, 1.8],
 [6.7, 2.5, 5.8, 1.8],
 [7.2, 3.6, 6.1, 2.5],
 [6.5, 3.2, 5.1, 2.],
 [6.4, 2.7, 5.3, 1.9],
 [6.8, 3. , 5.5, 2.1],

[5.7, 2.5, 5. , 2.],
[5.8, 2.8, 5.1, 2.4],
[6.4, 3.2, 5.3, 2.3],
[6.5, 3. , 5.5, 1.8],
[7.7, 3.8, 6.7, 2.2],
[7.7, 2.6, 6.9, 2.3],
[6. , 2.2, 5. , 1.5],
[6.9, 3.2, 5.7, 2.3],
[5.6, 2.8, 4.9, 2.],
[7.7, 2.8, 6.7, 2.],
[6.3, 2.7, 4.9, 1.8],
[6.7, 3.3, 5.7, 2.1],
[7.2, 3.2, 6. , 1.8],
[6.2, 2.8, 4.8, 1.8],
[6.1, 3. , 4.9, 1.8],
[6.4, 2.8, 5.6, 2.1],
[7.2, 3. , 5.8, 1.6],
[7.4, 2.8, 6.1, 1.9],
[7.9, 3.8, 6.4, 2.],
[6.4, 2.8, 5.6, 2.2],
[6.3, 2.8, 5.1, 1.5],
[6.1, 2.6, 5.6, 1.4],
[7.7, 3. , 6.1, 2.3],
[6.3, 3.4, 5.6, 2.4],
[6.4, 3.1, 5.5, 1.8],
[6. , 3. , 4.8, 1.8],
[6.9, 3.1, 5.4, 2.1],
[6.7, 3.1, 5.6, 2.4],
[6.9, 3.1, 5.1, 2.3],
[5.8, 2.7, 5.1, 1.9],
[6.8, 3.2, 5.9, 2.3],
[6.7, 3.3, 5.7, 2.5],
[6.7, 3. , 5.2, 2.3],
[6.3, 2.5, 5. , 1.9],
[6.5, 3. , 5.2, 2.],
[6.2, 3.4, 5.4, 2.3],
[5.9, 3. , 5.1, 1.8]]

```
[5]: y = iris.iloc[:, 4].values
y
```

```
[5]: array(['Normal', 'Normal', 'Normal', 'Normal', 'Normal', 'Normal',  
          'Normal', 'Normal', 'Normal', 'Normal', 'Normal', 'Normal',  
          'Normal', 'Normal', 'Normal', 'Normal', 'Normal', 'Normal',  
          'Normal', 'Normal', 'Normal', 'Normal', 'Normal', 'Normal',  
          'Normal', 'Normal', 'Normal', 'Normal', 'Normal', 'Normal']
```

[illegible]

```
[6]: import seaborn as sns
import matplotlib.pyplot as plt
plot = sns.countplot(data=iris, x="disorder")

# Get the count of each category
value_counts = iris["disorder"].value_counts()

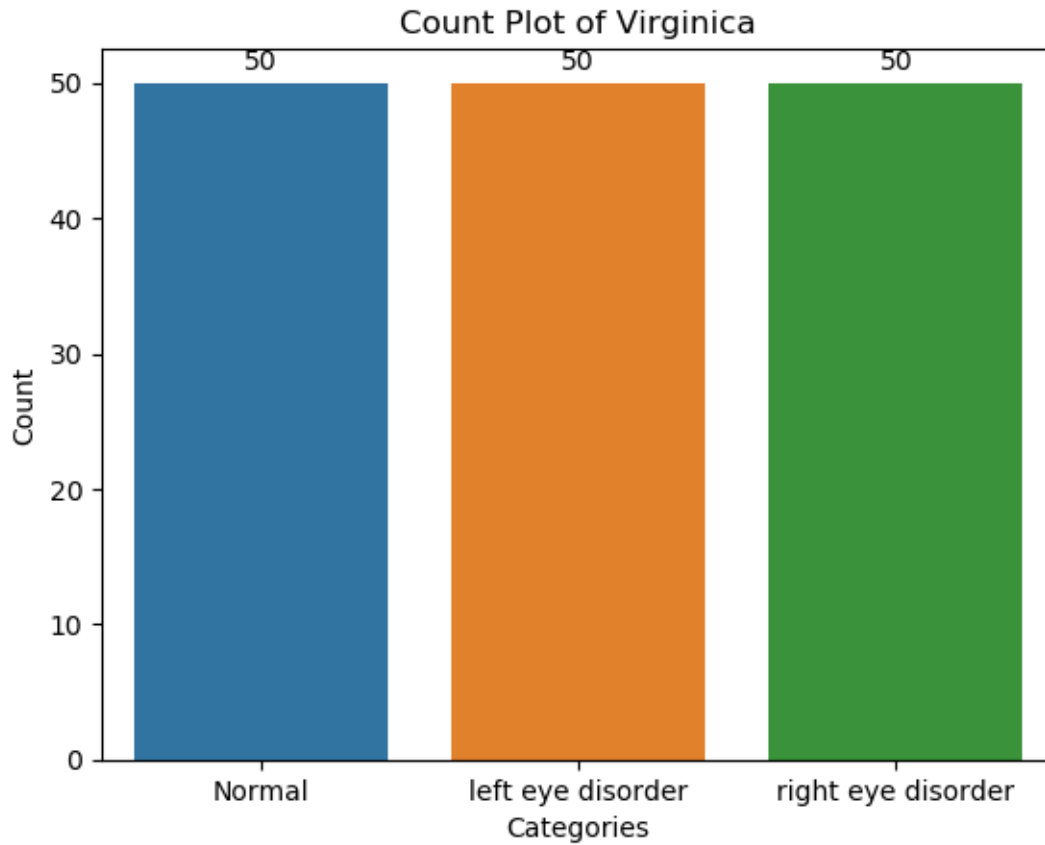
# Display the count values on top of the bars
for i, count in enumerate(value_counts):
    plot.text(x=i, y=count + 1, s=str(count), ha="center")
```

```

# Customize the plot
plot.set_xlabel("Categories")
plot.set_ylabel("Count")
plot.set_title("Count Plot of Virginica")

# Show the plot
plt.show()

```



```

[7]: labels = set(y)
      labels

```

```

[7]: {'Normal', 'left eye disorder', 'right eye disorder'}

```

```

[8]: from sklearn import preprocessing
      le = preprocessing.LabelEncoder()

```

```

[9]: y = le.fit_transform(y)
      y

```

```
[9]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
          0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
          0, 0, 0, 0, 0, 0, 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, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
          2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
          2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

```
[ ]:
```

```
[10]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
↳random_state=0)
print("X shape is",X.shape)
print("Y shape is",y.shape)
print("X_test shape is",X_test.shape)
print("X_train shape is",X_train.shape)
print("y_train shape is",y_train.shape)
print("y_test shape is",y_test.shape)
```

```
X shape is (150, 4)
Y shape is (150,)
X_test shape is (30, 4)
X_train shape is (120, 4)
y_train shape is (120,)
y_test shape is (30,)
```

```
[11]: from sklearn.linear_model import LogisticRegression
lr = LogisticRegression(solver='liblinear')
lr.fit(X_train,y_train)
y_pred = lr.predict(X_test)
print("Original y_test values are ",y_test)
print("predicted y_test values are",y_pred)
```

```
Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 2 0 0 2 0
0 1 1 0]
```

```
[12]: from sklearn.metrics import accuracy_score
from sklearn.metrics import precision_score
from sklearn.metrics import recall_score
from sklearn.metrics import f1_score
# Target is multiclass but average='binary'. Please choose another average
↳setting, one of [None, 'micro', 'macro', 'weighted'].
accuracy = accuracy_score(y_test, y_pred)
# Precision
```

```
precision = precision_score(y_test, y_pred, average='weighted')
# Recall
recall = recall_score(y_test, y_pred, average='weighted')
# F1 score
f1 = f1_score(y_test, y_pred, average='weighted')
```

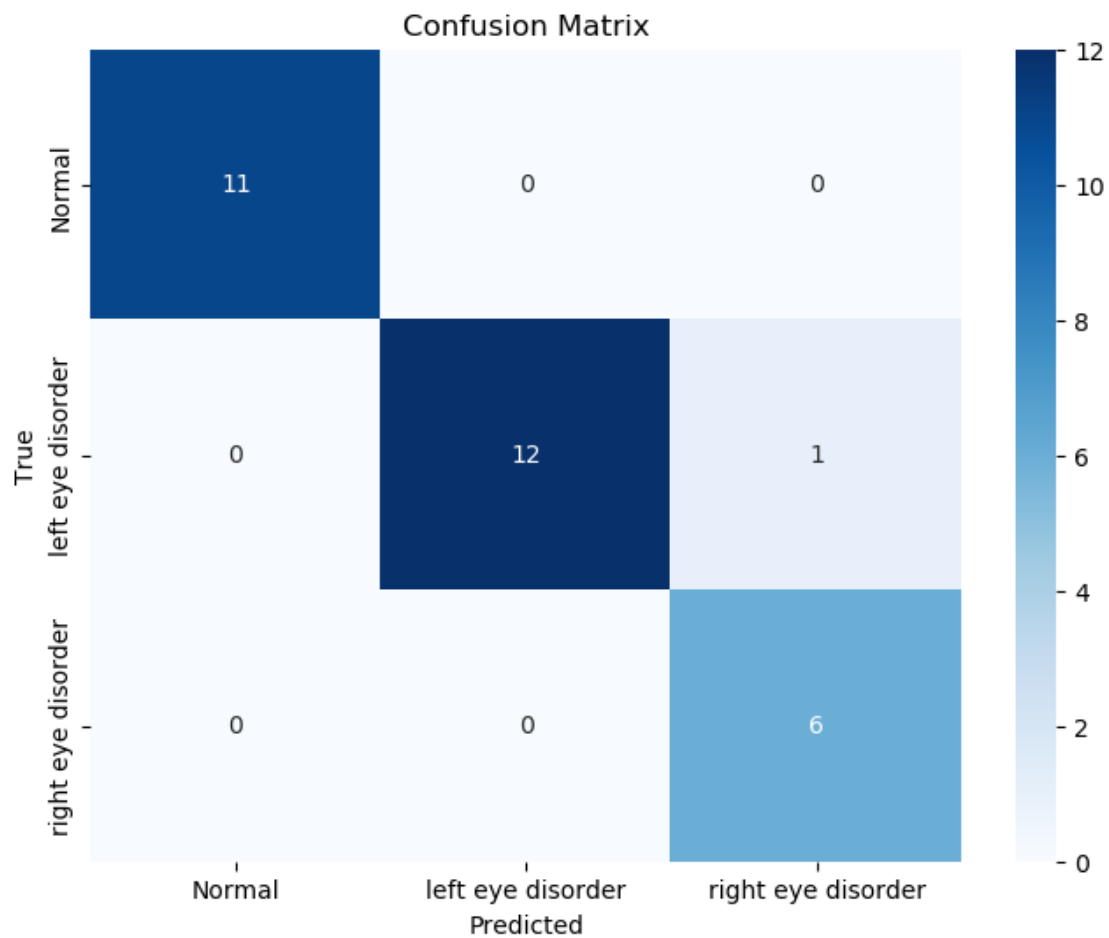
```
[13]: print(f'Accuracy: {accuracy:.2f}')
      print(f'Precision: {precision:.2f}')
      print(f'Recall: {recall:.2f}')
      print(f'F1 Score: {f1:.2f}')
```

```
Accuracy: 0.97
Precision: 0.97
Recall: 0.97
F1 Score: 0.97
```

```
[14]: from sklearn.metrics import confusion_matrix

      # Assuming y_test and y_pred are your true labels and predicted labels,
      ↪ respectively
      conf_matrix = confusion_matrix(y_test, y_pred)

      # Create a heatmap of the confusion matrix
      plt.figure(figsize=(8, 6))
      sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
      ↪ yticklabels=labels)
      plt.xlabel('Predicted')
      plt.ylabel('True')
      plt.title('Confusion Matrix')
      plt.show()
```

```
[15]: from sklearn.metrics import classification_report

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report:')
print(class_report)
```

Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	0.92	0.96	13
right eye disorder	0.86	1.00	0.92	6
accuracy			0.97	30
macro avg	0.95	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

```
[16]: from sklearn.tree import DecisionTreeClassifier
dt = DecisionTreeClassifier()
dt.fit(X_train,y_train)
y_pred = lr.predict(X_test)

print("Original y_test values are ",y_test)
print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix of of LRC')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report of LRC:')
print(class_report)
```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

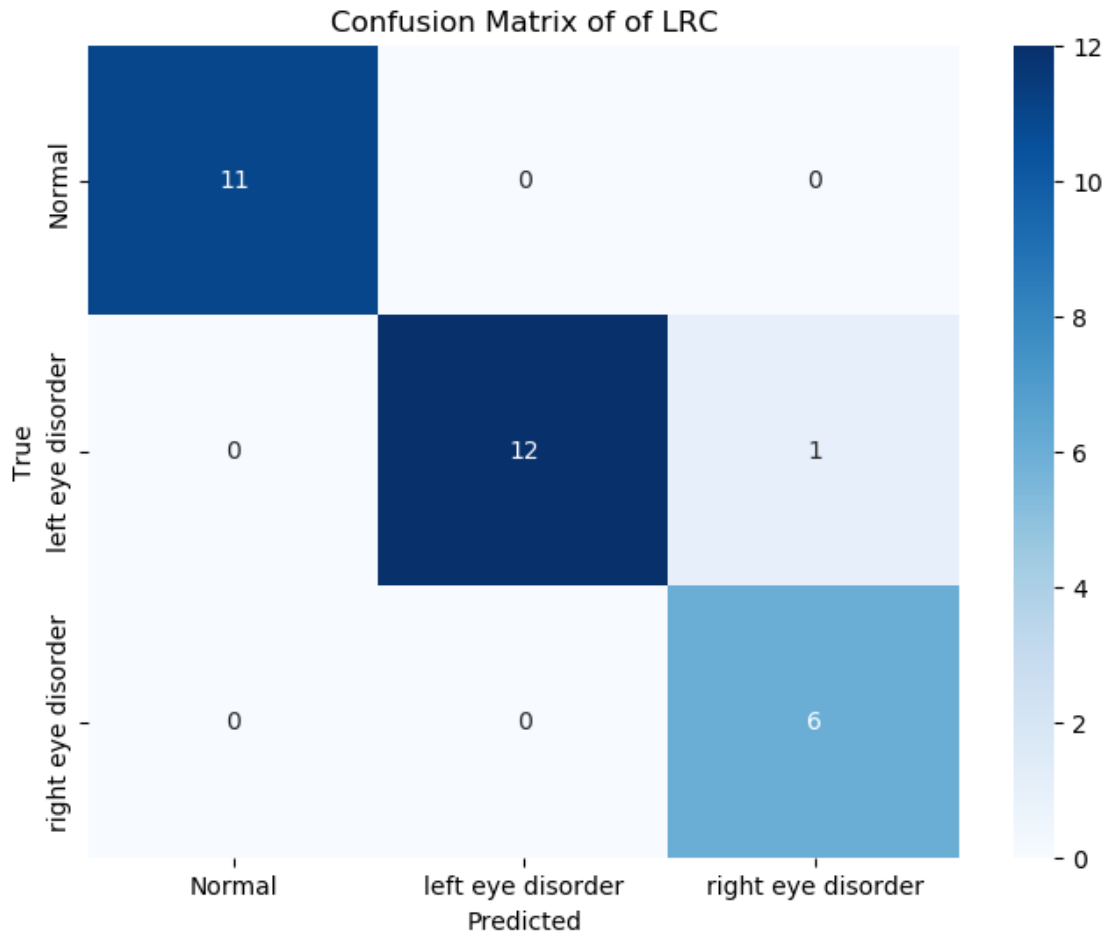
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 2 0 0 2 0
0 1 1 0]

Accuracy: 0.97

Precision: 0.97

Recall: 0.97

F1 Score: 0.97



Classification Report of LRC:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	0.92	0.96	13
right eye disorder	0.86	1.00	0.92	6
accuracy			0.97	30
macro avg	0.95	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

```
[17]: from sklearn.tree import DecisionTreeClassifier
dt = DecisionTreeClassifier()
dt.fit(X_train,y_train)
y_pred = dt.predict(X_test)

print("Original y_test values are ",y_test)
```

```

print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix of DTC')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report of DTC:')
print(class_report)

```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

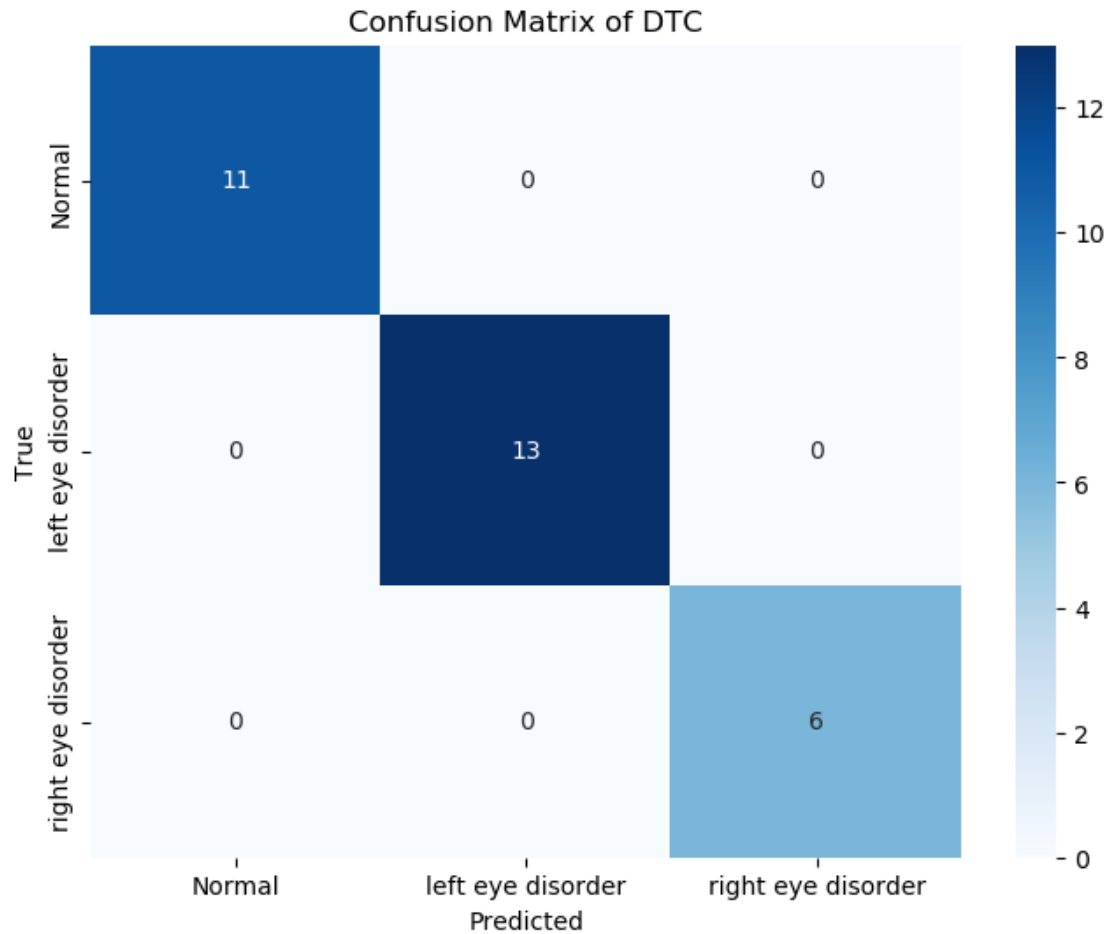
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

Accuracy: 1.00

Precision: 1.00

Recall: 1.00

F1 Score: 1.00



Classification Report of DTC:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[18]: from sklearn.ensemble import RandomForestClassifier
rf = RandomForestClassifier()
rf.fit(X_train,y_train)
y_pred = rf.predict(X_test)

print("Original y_test values are ",y_test)
```

```

print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix of RFC')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report of RFC:')
print(class_report)

```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

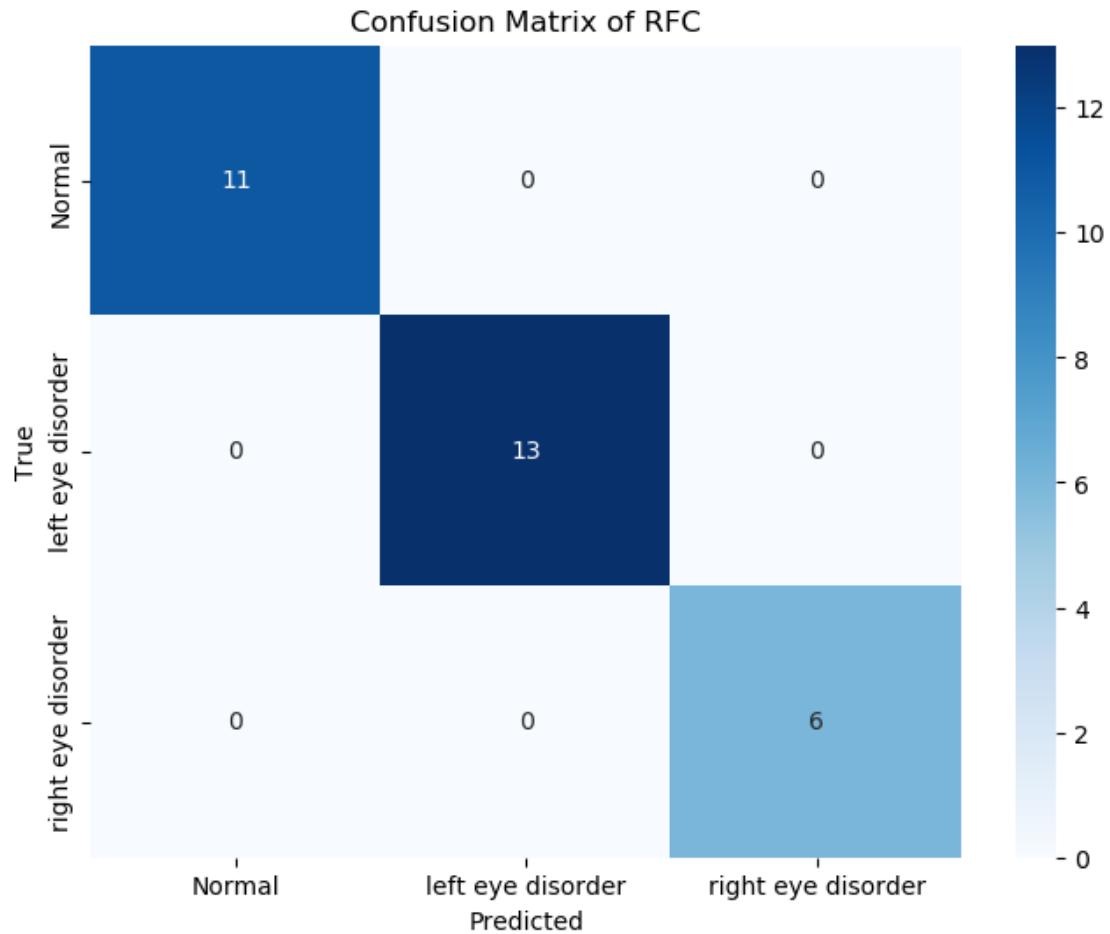
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

Accuracy: 1.00

Precision: 1.00

Recall: 1.00

F1 Score: 1.00



Classification Report of RFC:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[19]: from sklearn.svm import SVC
svm = SVC()
svm.fit(X_train, y_train)
y_pred = svm.predict(X_test)

print("Original y_test values are ",y_test)
```

```

print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix of SVM')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

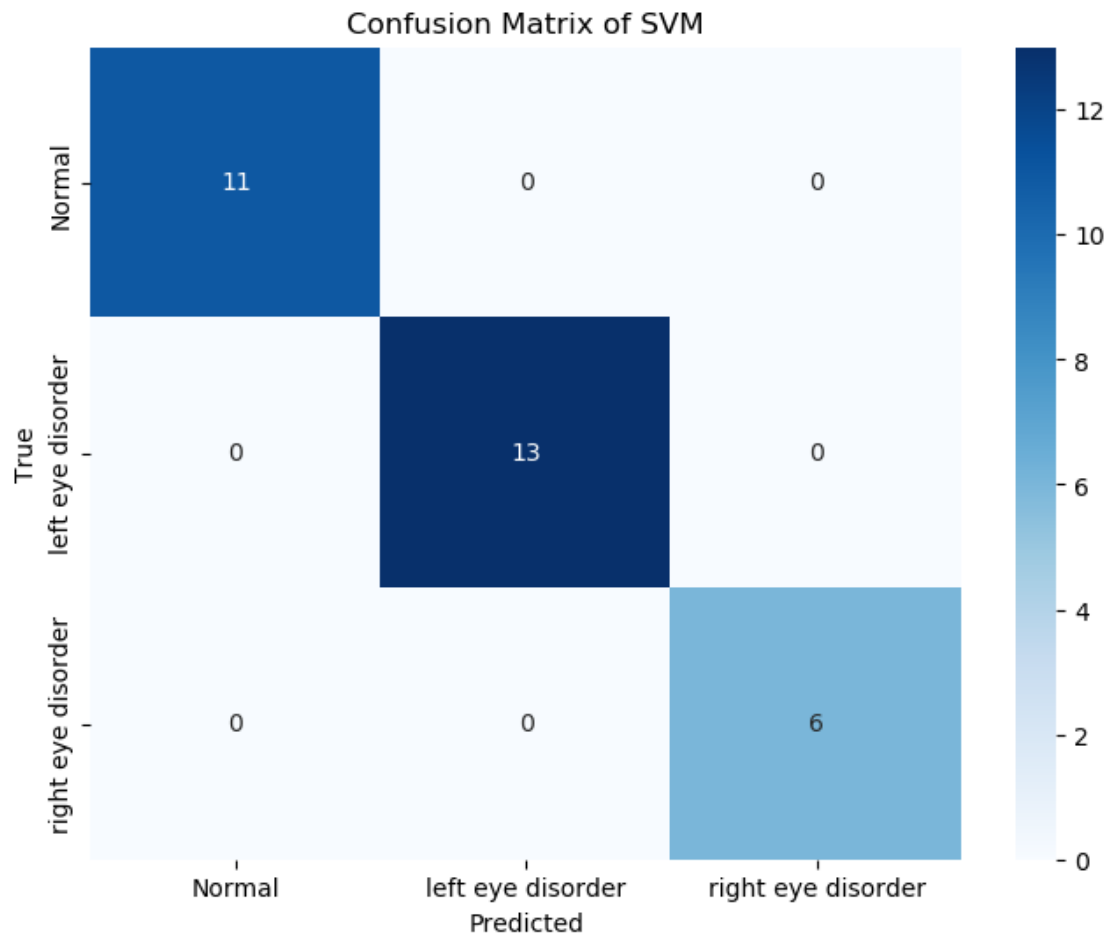
# Print the classification report with labels
print('Classification Report of SVM:')
print(class_report)

```

```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
Accuracy: 1.00
Precision: 1.00
Recall: 1.00
F1 Score: 1.00

```

Classification Report of SVM:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[20]: from sklearn.neighbors import KNeighborsClassifier

knn = KNeighborsClassifier()
knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
```

```

# The rest of your code remains the same
print("Original y_test values are ",y_test)
print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

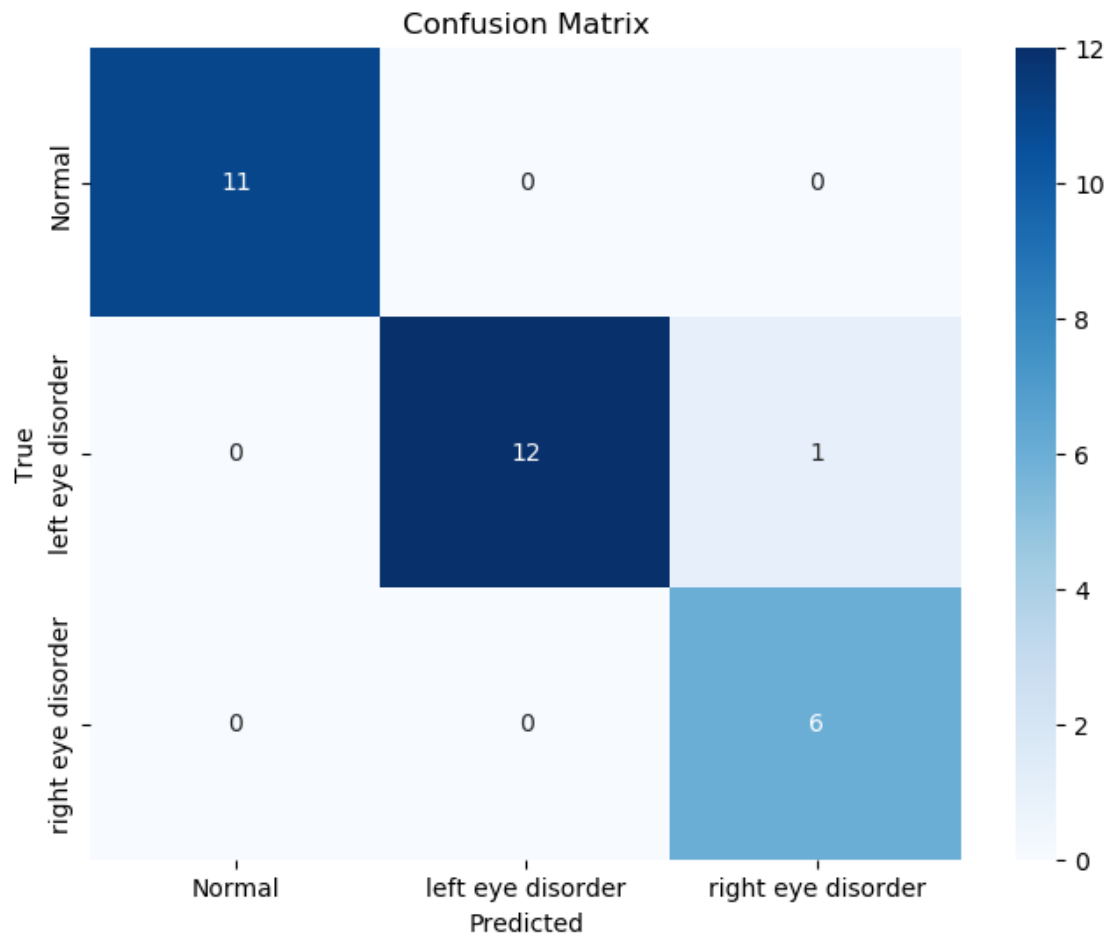
# Print the classification report with labels
print('Classification Report:')
print(class_report)

```

```

Original y_test values are  [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 2 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
Accuracy: 0.97
Precision: 0.97
Recall: 0.97
F1 Score: 0.97

```



Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	0.92	0.96	13
right eye disorder	0.86	1.00	0.92	6
accuracy			0.97	30
macro avg	0.95	0.97	0.96	30
weighted avg	0.97	0.97	0.97	30

```
[21]: from sklearn.neural_network import MLPClassifier

mlp = MLPClassifier()
mlp.fit(X_train, y_train)
y_pred = mlp.predict(X_test)
```

```

# The rest of your code remains the same

# The rest of your code remains the same
print("Original y_test values are ",y_test)
print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report:')
print(class_report)

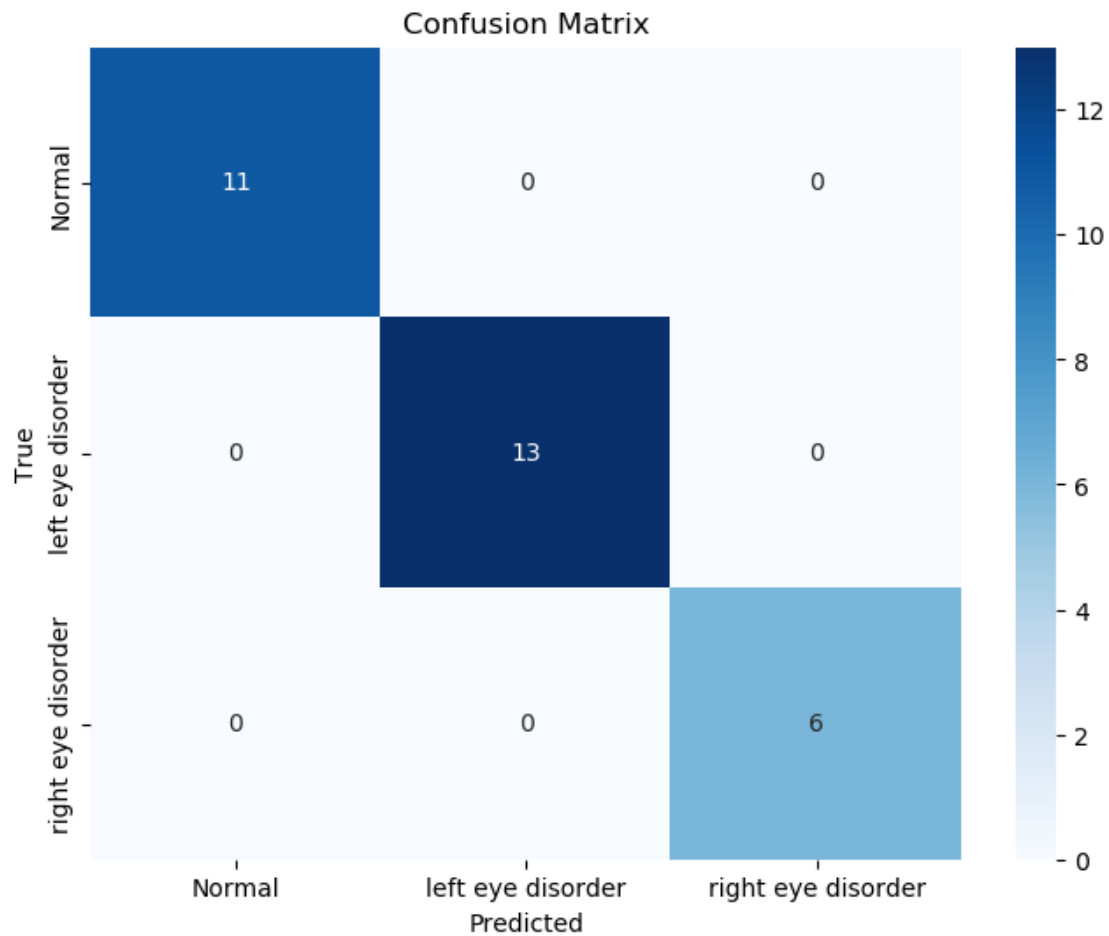
```

```

c:\users\asus\appdata\local\programs\python\python37\lib\site-
packages\sklearn\normalization\_multilayer_perceptron.py:585:
ConvergenceWarning: Stochastic Optimizer: Maximum iterations (200) reached and
the optimization hasn't converged yet.
  % self.max_iter, ConvergenceWarning)

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
Accuracy: 1.00
Precision: 1.00
Recall: 1.00
F1 Score: 1.00

```



Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[22]: from sklearn.linear_model import LinearRegression
```

```
lin_reg = LinearRegression()
lin_reg.fit(X_train, y_train)
y_pred = lin_reg.predict(X_test)
y_pred=np.round(y_pred)
```

```

print("Original y_test values are ",y_test)
print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

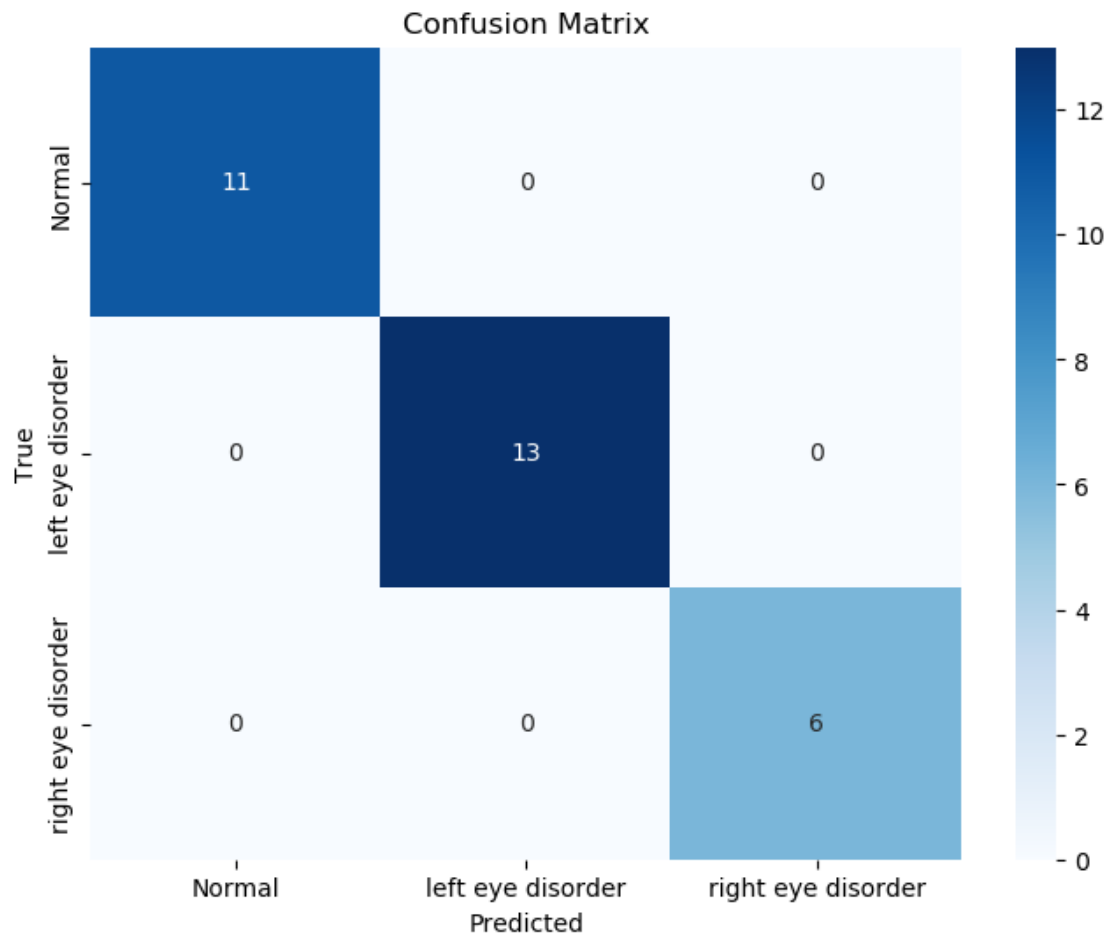
# Print the classification report with labels
print('Classification Report:')
print(class_report)

```

```

Original y_test values are  [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]
predicted y_test values are [ 2.  1. -0.  2. -0.  2. -0.  1.  1.  1.  2.  1.  1.
 1.  1. -0.  1.  1.
 0. -0.  2.  1.  0.  0.  2. -0.  0.  1.  1.  0.]
Accuracy: 1.00
Precision: 1.00
Recall: 1.00
F1 Score: 1.00

```



Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[23]: from sklearn.ensemble import AdaBoostClassifier
```

```
ada_boost = AdaBoostClassifier()
ada_boost.fit(X_train, y_train)
y_pred = ada_boost.predict(X_test)
print("Original y_test values are ",y_test)
```

```

print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report:')
print(class_report)

```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

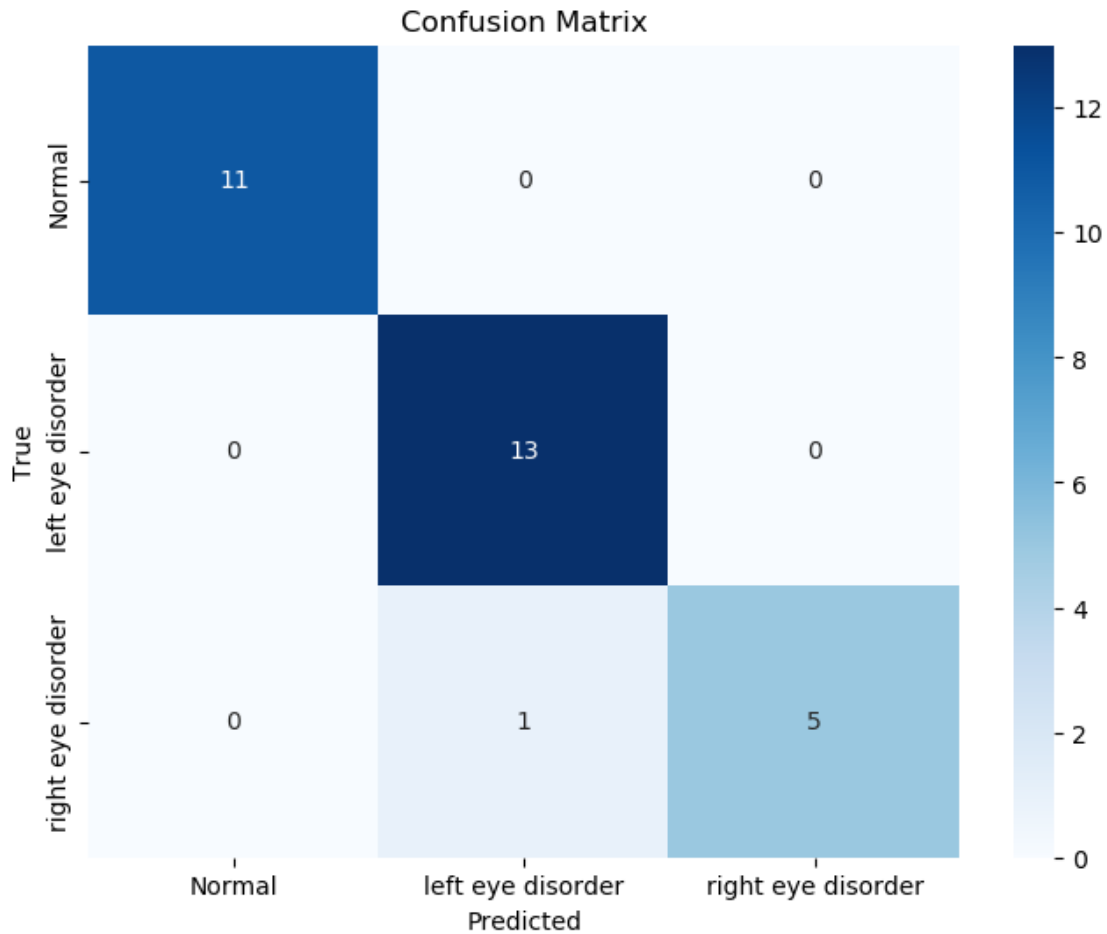
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 1 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

Accuracy: 0.97

Precision: 0.97

Recall: 0.97

F1 Score: 0.97



Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	0.93	1.00	0.96	13
right eye disorder	1.00	0.83	0.91	6
accuracy			0.97	30
macro avg	0.98	0.94	0.96	30
weighted avg	0.97	0.97	0.97	30

```
[24]: from sklearn.ensemble import ExtraTreesClassifier

extra_tree = ExtraTreesClassifier()
extra_tree.fit(X_train, y_train)
y_pred = extra_tree.predict(X_test)
print("Original y_test values are ",y_test)
```

```

print("predicted y_test values are",y_pred)

accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print(f'Accuracy: {accuracy:.2f}')
print(f'Precision: {precision:.2f}')
print(f'Recall: {recall:.2f}')
print(f'F1 Score: {f1:.2f}')

conf_matrix = confusion_matrix(y_test, y_pred)
# Create a heatmap of the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=labels,
            yticklabels=labels)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()

class_report = classification_report(y_test, y_pred, target_names=labels)

# Print the classification report with labels
print('Classification Report:')
print(class_report)

```

Original y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

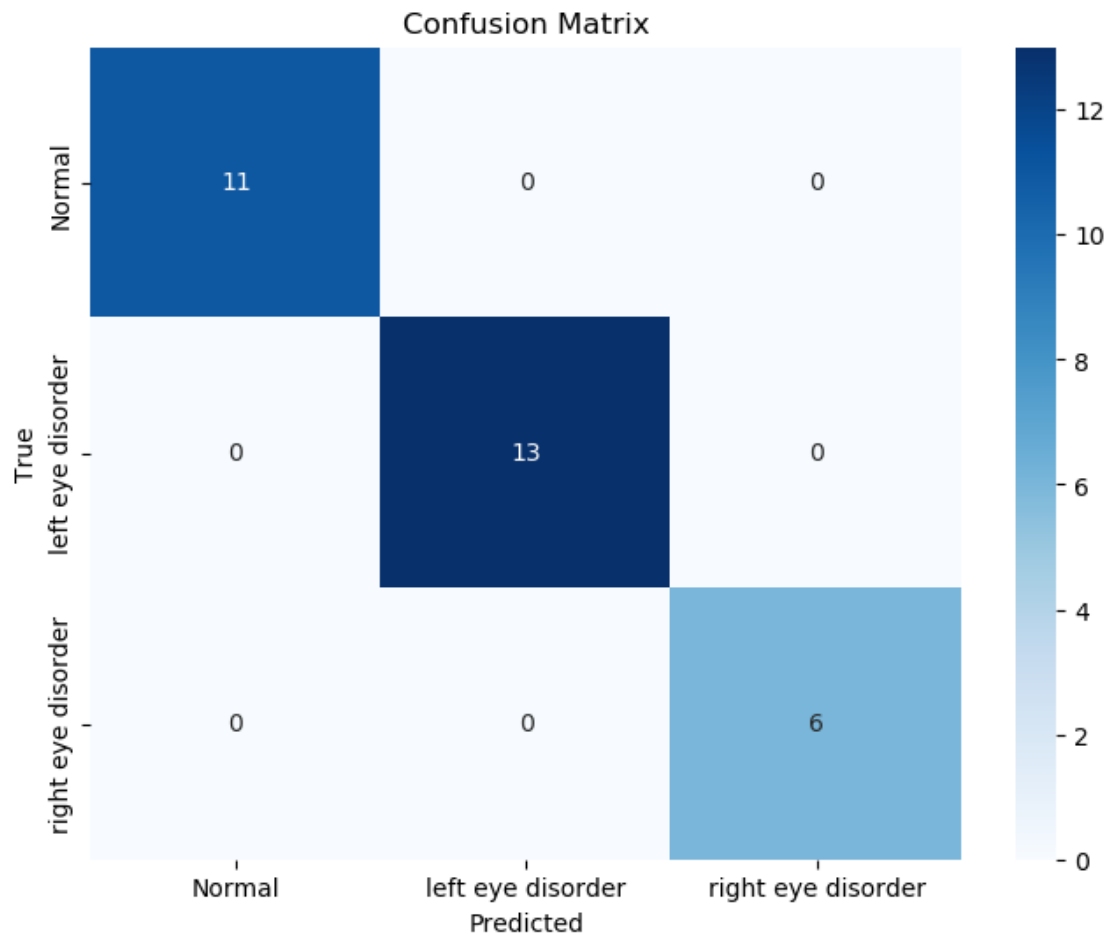
predicted y_test values are [2 1 0 2 0 2 0 1 1 1 2 1 1 1 1 0 1 1 0 0 2 1 0 0 2 0
0 1 1 0]

Accuracy: 1.00

Precision: 1.00

Recall: 1.00

F1 Score: 1.00



Classification Report:

	precision	recall	f1-score	support
Normal	1.00	1.00	1.00	11
left eye disorder	1.00	1.00	1.00	13
right eye disorder	1.00	1.00	1.00	6
accuracy			1.00	30
macro avg	1.00	1.00	1.00	30
weighted avg	1.00	1.00	1.00	30

```
[25]: # all classifiers available in SKlearn
# Linear Models
from sklearn.linear_model import LogisticRegression

# Support Vector Machines
from sklearn.svm import SVC
```

```

# Decision Trees
from sklearn.tree import DecisionTreeClassifier

# Ensemble Methods
from sklearn.ensemble import RandomForestClassifier, BaggingClassifier,
↳ GradientBoostingClassifier, AdaBoostClassifier, ExtraTreesClassifier

# Naive Bayes
from sklearn.naive_bayes import GaussianNB, MultinomialNB, ComplementNB,
↳ BernoulliNB

# Nearest Neighbors
from sklearn.neighbors import KNeighborsClassifier

```

```

[26]: # all regressors available in SKlearn

# Linear Models
from sklearn.linear_model import LinearRegression, Ridge, Lasso, ElasticNet

# Support Vector Machines
from sklearn.svm import SVR

# Decision Trees
from sklearn.tree import DecisionTreeRegressor

# Ensemble Methods
from sklearn.ensemble import RandomForestRegressor, BaggingRegressor,
↳ GradientBoostingRegressor, AdaBoostRegressor, ExtraTreesRegressor

# Nearest Neighbors
from sklearn.neighbors import KNeighborsRegressor

```

```

[27]: # all unsupervised machine learning models available in SKlearn

# Clustering
from sklearn.cluster import KMeans, DBSCAN, AgglomerativeClustering

# Support Vector Machines for Outliers Detection
from sklearn.svm import OneClassSVM

```

```

[28]: # all feature reduction models available in SKlearn

# Principal Component Analysis (PCA)
from sklearn.decomposition import PCA

# Dimensionality Reduction

```

```
from sklearn.decomposition import TruncatedSVD
```