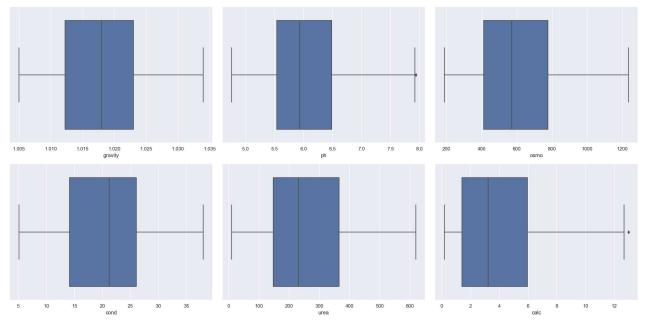
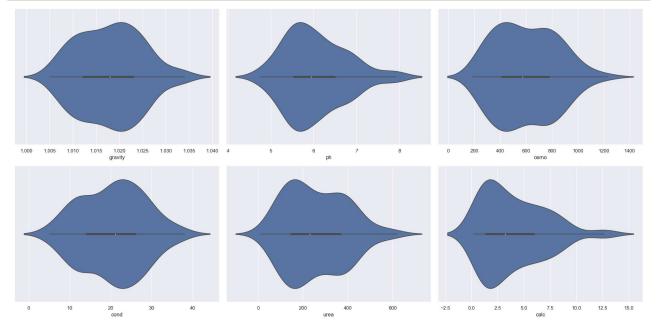
```
In [1]: import pandas as pd
         import seaborn as sns
         import matplotlib.pyplot as plt
         import numpy as np
         sns.set_theme(color_codes=True)
In [2]: df = pd.read_csv('kidney-stone-dataset.csv')
         df.head()
Out[2]:
             Unnamed: 0 gravity
                                 ph osmo cond urea calc target
          0
                      0
                         1.021 4.91
                                                  443
                                                      2.45
                                                               0
                                       725
                                            14.0
          1
                      1
                          1.017 5.74
                                       577
                                            20.0
                                                  296 4.49
                                                               0
          2
                         1.008 7.20
                      2
                                       321
                                            14.9
                                                  101
                                                      2.36
                                                               0
          3
                      3
                          1.011 5.51
                                       408
                                            12.6
                                                  224
                                                      2.15
                                                               0
                      4
                          1.005 6.52
                                             7.5
                                                               0
                                       187
                                                   91 1.16
In [3]: df.drop(columns='Unnamed: 0', inplace=True)
In [4]: | df.head()
Out[4]:
             gravity
                     ph osmo cond urea calc target
          0
              1.021
                    4.91
                                           2.45
                                                    0
                           725
                                14.0
                                      443
          1
              1.017 5.74
                           577
                                20.0
                                      296 4.49
                                                    0
          2
              1.008 7.20
                           321
                                14.9
                                      101
                                           2.36
                                                    0
          3
              1.011
                    5.51
                           408
                                12.6
                                      224 2.15
                                                    0
              1.005 6.52
                           187
                                 7.5
                                       91 1.16
                                                    0
```

Exploratory Data Analysis

```
In [6]: num_vars = ['gravity', 'ph', 'osmo', 'cond', 'urea', 'calc']
fig, axs = plt.subplots(nrows=2, ncols=3, figsize=(20, 10))
axs = axs.flatten()
for i, var in enumerate(num_vars):
    sns.boxplot(x=var, data=df, ax=axs[i])
fig.tight_layout()
plt.show()
```





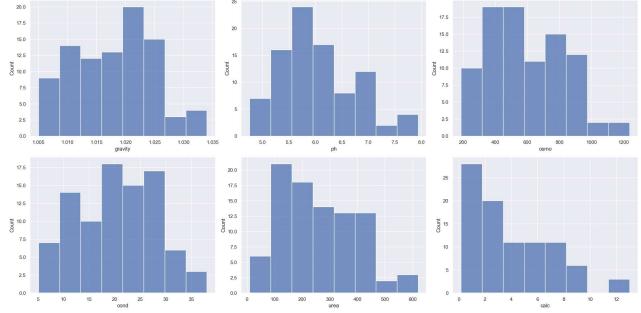
```
In [13]: num_vars = ['gravity', 'ph', 'osmo', 'cond', 'urea', 'calc']

fig, axs = plt.subplots(nrows=2, ncols=3, figsize=(20, 10))
axs = axs.flatten()

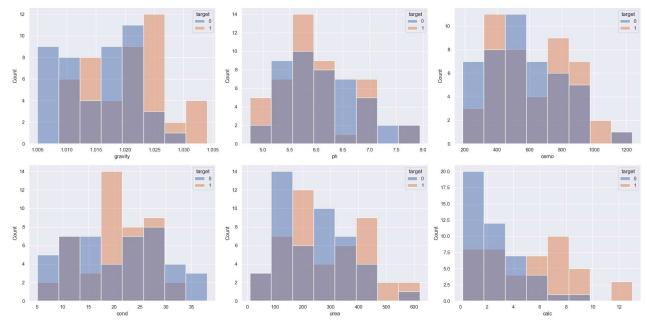
for i, var in enumerate(num_vars):
    sns.histplot(x=var, data=df, ax=axs[i])

fig.tight_layout()

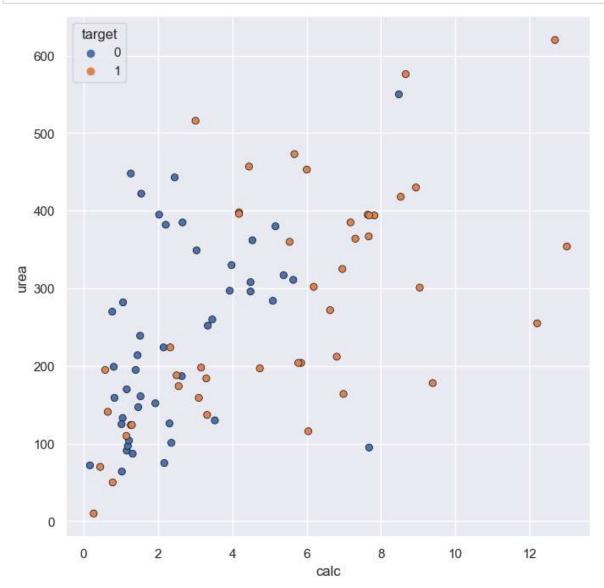
plt.show()
```



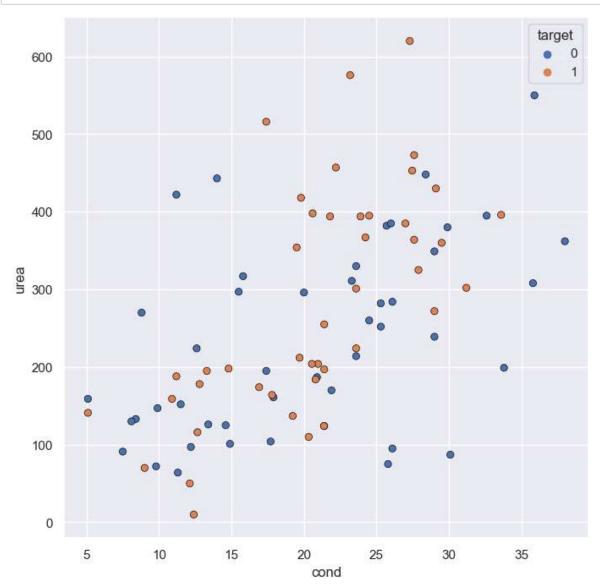
```
In [14]: num_vars = ['gravity', 'ph', 'osmo', 'cond', 'urea', 'calc']
fig, axs = plt.subplots(nrows=2, ncols=3, figsize=(20, 10))
axs = axs.flatten()
for i, var in enumerate(num_vars):
    sns.histplot(x=var, data=df, hue='target', ax=axs[i])
fig.tight_layout()
plt.show()
```



```
In [16]: plt.figure(figsize=(8,8),dpi=100)
    sns.scatterplot(x="calc", y="urea", hue="target", data=df, edgecolor="black")
    plt.show()
```



```
In [17]: plt.figure(figsize=(8,8),dpi=100)
    sns.scatterplot(x="cond", y="urea", hue="target", data=df, edgecolor="black")
    plt.show()
```



Data Preproceessing

Check 'Target' value is it balanced or not

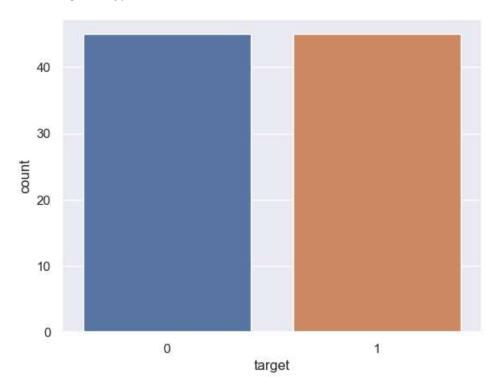
```
In [19]: sns.countplot(df['target'])
df['target'].value_counts()
```

D:\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following varia ble as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[19]: 0 45 1 45

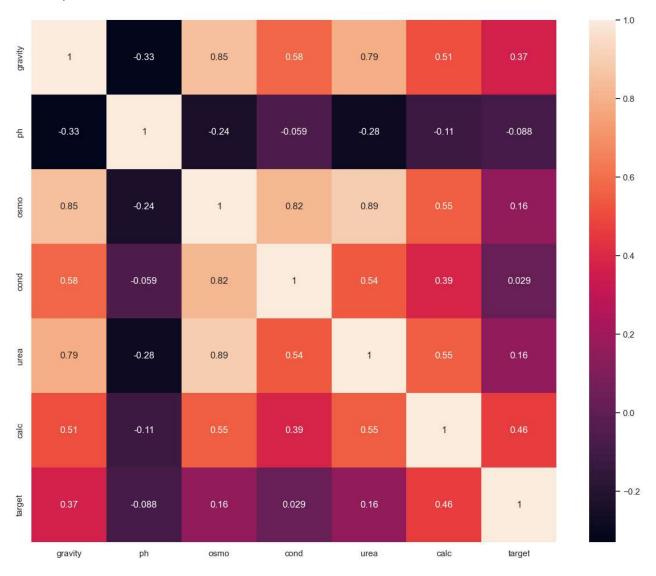
Name: target, dtype: int64



There's no outlier, so we don't have to remove it

```
In [21]: plt.figure(figsize=(15,12))
sns.heatmap(df.corr(), fmt='.2g', annot=True)
```

Out[21]: <AxesSubplot:>



Train test split

```
In [22]: X = df.drop('target', axis=1)
y = df['target']
In [23]: #test size 20% and train size 80%
```

```
In [23]: #test size 20% and train size 80%
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
X_train, X_test, y_train, y_test = train_test_split(X,y, test_size=0.2,random_state=0)
```

Machine Learrning Model Building

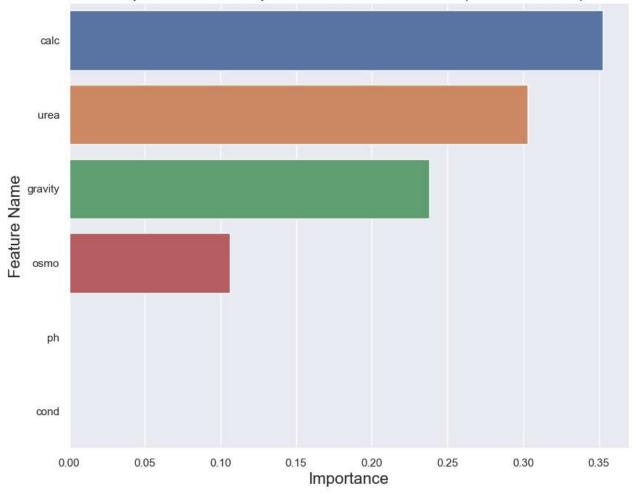
Decision Tree

```
In [24]: | from sklearn.tree import DecisionTreeClassifier
         from sklearn.model selection import GridSearchCV
         dtree = DecisionTreeClassifier()
         param_grid = {
              'max_depth': [3, 4, 5, 6, 7, 8],
              'min_samples_split': [2, 3, 4],
              'min_samples_leaf': [1, 2, 3, 4]
         # Perform a grid search with cross-validation to find the best hyperparameters
         grid search = GridSearchCV(dtree, param grid, cv=5)
         grid search.fit(X train, y train)
         # Print the best hyperparameters
         print(grid_search.best_params_)
         {'max_depth': 7, 'min_samples_leaf': 1, 'min_samples_split': 4}
In [25]: from sklearn.tree import DecisionTreeClassifier
         dtree = DecisionTreeClassifier(random_state=0, max_depth=7, min_samples_leaf=1, min_samples_split=
         dtree.fit(X_train, y_train)
Out[25]: DecisionTreeClassifier(max depth=7, min samples split=4, random state=0)
In [26]: y pred = dtree.predict(X test)
         print("Accuracy Score :", round(accuracy_score(y_test, y_pred)*100 ,2), "%")
         Accuracy Score: 66.67 %
In [27]: from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score, jaccard_score
         print('F-1 Score : ',(f1_score(y_test, y_pred, average='micro')))
         print('Precision Score : ',(precision_score(y_test, y_pred, average='micro')))
         print('Recall Score : ',(recall_score(y_test, y_pred, average='micro')))
print('Jaccard Score : ',(jaccard_score(y_test, y_pred, average='micro')))
         print('Log Loss : ',(log_loss(y_test, y_pred)))
         F-1 Score: 0.666666666666666
         Recall Score : 0.666666666666666
         Jaccard Score : 0.5
         Log Loss: 11.513014309129138
```

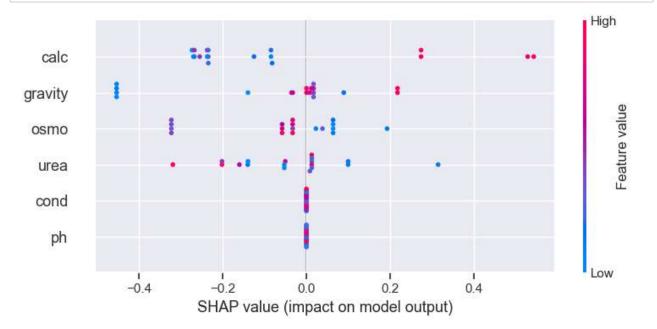
```
In [28]: imp_df = pd.DataFrame({
    "Feature Name": X_train.columns,
    "Importance": dtree.feature_importances_
})
fi = imp_df.sort_values(by="Importance", ascending=False)

fi2 = fi.head(10)
plt.figure(figsize=(10,8))
sns.barplot(data=fi2, x='Importance', y='Feature Name')
plt.title('Top 10 Feature Importance Each Attributes (Decision Tree)', fontsize=18)
plt.xlabel ('Importance', fontsize=16)
plt.ylabel ('Feature Name', fontsize=16)
plt.show()
```

Top 10 Feature Importance Each Attributes (Decision Tree)



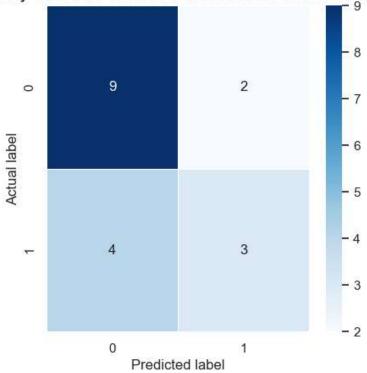
```
In [29]: # compute SHAP values
import shap
explainer = shap.TreeExplainer(dtree)
shap_values = explainer.shap_values(X_test)
shap.summary_plot(shap_values[1], X_test.values, feature_names = X_test.columns)
```



```
In [47]: from sklearn.metrics import confusion_matrix
    cm = confusion_matrix(y_test, y_pred)
    plt.figure(figsize=(5,5))
    sns.heatmap(data=cm,linewidths=.5, annot=True, cmap = 'Blues')
    plt.ylabel('Actual label')
    plt.xlabel('Predicted label')
    all_sample_title = 'Accuracy Score for Decision Tree: {0}'.format(dtree.score(X_test, y_test))
    plt.title(all_sample_title, size = 15)
```

Out[47]: Text(0.5, 1.0, 'Accuracy Score for Decision Tree: 0.66666666666666666')

Accuracy Score for Decision Tree: 0.666666666666666



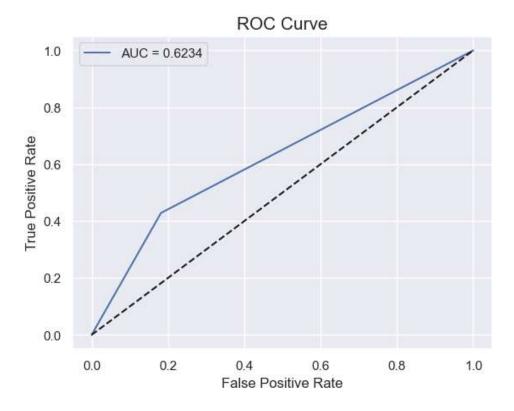
```
In [48]: from sklearn.metrics import roc_curve, roc_auc_score
    y_pred_proba = dtree.predict_proba(X_test)[:][:,1]

df_actual_predicted = pd.concat([pd.DataFrame(np.array(y_test), columns=['y_actual']), pd.DataFrame
    df_actual_predicted.index = y_test.index

fpr, tpr, tr = roc_curve(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])
    auc = roc_auc_score(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])

plt.plot(fpr, tpr, label='AUC = %0.4f' %auc)
    plt.plot(fpr, fpr, linestyle = '--', color='k')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plt.title('ROC Curve', size = 15)
    plt.legend()
```

Out[48]: <matplotlib.legend.Legend at 0x1aecbd067f0>



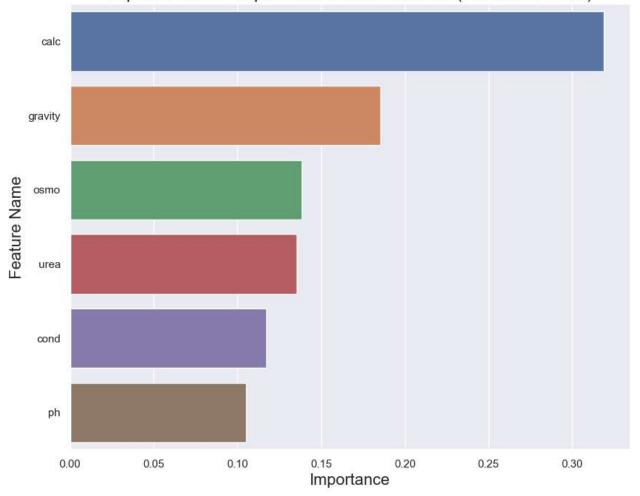
Random Forest

```
In [49]: | from sklearn.ensemble import RandomForestClassifier
         from sklearn.model selection import GridSearchCV
         rfc = RandomForestClassifier()
         param_grid = {
             'n_estimators': [100, 200],
             'max_depth': [None, 5, 10],
             'max_features': ['sqrt', 'log2', None]
         # Perform a grid search with cross-validation to find the best hyperparameters
         grid search = GridSearchCV(rfc, param grid, cv=5)
         grid search.fit(X train, y train)
         # Print the best hyperparameters
         print(grid_search.best_params_)
         {'max_depth': None, 'max_features': None, 'n_estimators': 200}
In [51]: | from sklearn.ensemble import RandomForestClassifier
         rfc = RandomForestClassifier(random state=0)
         rfc.fit(X_train, y_train)
Out[51]: RandomForestClassifier(random state=0)
In [52]: y_pred = rfc.predict(X_test)
         print("Accuracy Score :", round(accuracy_score(y_test, y_pred)*100 ,2), "%")
         Accuracy Score: 88.89 %
In [53]: from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score, jaccard_score
         print('F-1 Score : ',(f1_score(y_test, y_pred, average='micro')))
         print('Precision Score : ',(precision_score(y_test, y_pred, average='micro')))
         print('Recall Score : ',(recall_score(y_test, y_pred, average='micro')))
print('Jaccard Score : ',(jaccard_score(y_test, y_pred, average='micro')))
         print('Log Loss : ',(log_loss(y_test, y_pred)))
         Jaccard Score: 0.8
         Log Loss: 3.837686243736199
```

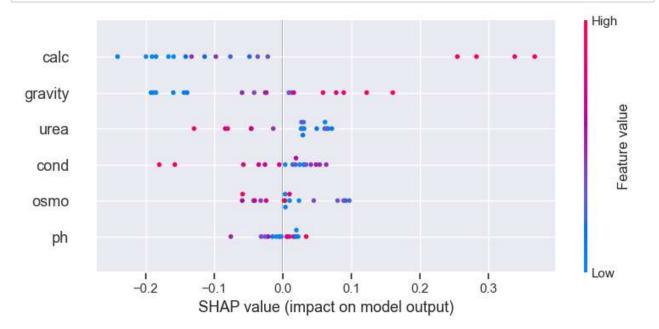
```
In [54]: imp_df = pd.DataFrame({
    "Feature Name": X_train.columns,
    "Importance": rfc.feature_importances_
})
fi = imp_df.sort_values(by="Importance", ascending=False)

fi2 = fi.head(10)
plt.figure(figsize=(10,8))
sns.barplot(data=fi2, x='Importance', y='Feature Name')
plt.title('Top 10 Feature Importance Each Attributes (Random Forest)', fontsize=18)
plt.xlabel ('Importance', fontsize=16)
plt.ylabel ('Feature Name', fontsize=16)
plt.show()
```

Top 10 Feature Importance Each Attributes (Random Forest)

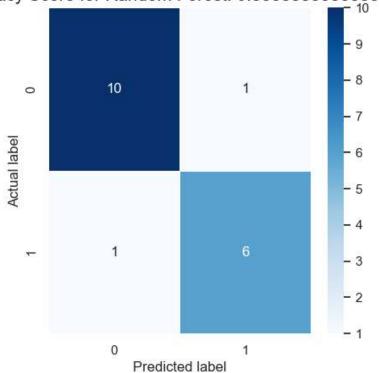


```
In [55]: # compute SHAP values
import shap
explainer = shap.TreeExplainer(rfc)
shap_values = explainer.shap_values(X_test)
shap.summary_plot(shap_values[1], X_test.values, feature_names = X_test.columns)
```



```
In [57]: from sklearn.metrics import confusion_matrix
    cm = confusion_matrix(y_test, y_pred)
    plt.figure(figsize=(5,5))
    sns.heatmap(data=cm,linewidths=.5, annot=True, cmap = 'Blues')
    plt.ylabel('Actual label')
    plt.xlabel('Predicted label')
    all_sample_title = 'Accuracy Score for Random Forest: {0}'.format(rfc.score(X_test, y_test))
    plt.title(all_sample_title, size = 15)
```





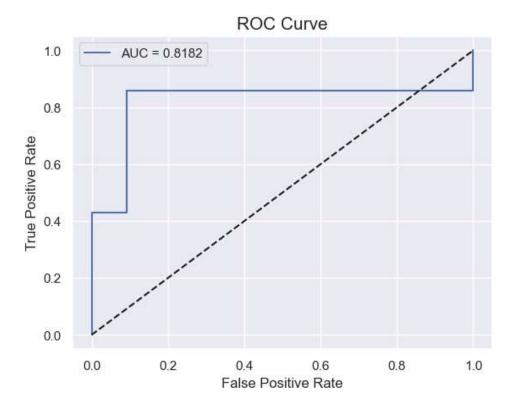
```
In [58]: from sklearn.metrics import roc_curve, roc_auc_score
    y_pred_proba = rfc.predict_proba(X_test)[:][:,1]

df_actual_predicted = pd.concat([pd.DataFrame(np.array(y_test), columns=['y_actual']), pd.DataFrame
    df_actual_predicted.index = y_test.index

fpr, tpr, tr = roc_curve(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])
    auc = roc_auc_score(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])

plt.plot(fpr, tpr, label='AUC = %0.4f' %auc)
    plt.plot(fpr, fpr, linestyle = '--', color='k')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plt.title('ROC Curve', size = 15)
    plt.legend()
```

Out[58]: <matplotlib.legend.Legend at 0x1aecca7cfd0>



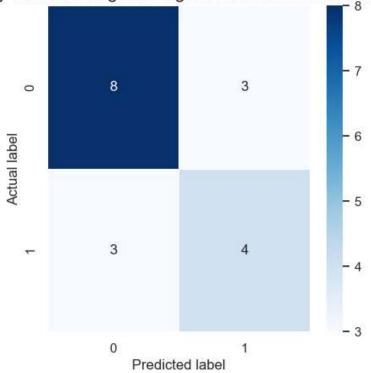
Logistic Regression

```
In [61]: | from sklearn.linear_model import LogisticRegression
         from sklearn.model selection import GridSearchCV
         # Create a Logistic Regression model
         logreg = LogisticRegression(solver='liblinear', max_iter=10000)
         # Define the parameter grid
         param_grid = {
              'penalty': ['l1', 'l2'],
             'C': [0.01, 0.1, 1, 10]
         # Perform a grid search with cross-validation to find the best hyperparameters
         grid search = GridSearchCV(logreg, param grid, cv=5)
         grid_search.fit(X_train, y_train)
         # Print the best hyperparameters
         print(grid_search.best_params_)
         {'C': 0.1, 'penalty': 'l1'}
In [62]: from sklearn.ensemble import RandomForestClassifier
         logreg = LogisticRegression(solver='liblinear', max iter=10000, C=0.1, penalty='l1')
         logreg.fit(X train, y train)
Out[62]: LogisticRegression(C=0.1, max iter=10000, penalty='l1', solver='liblinear')
In [63]: |y_pred = logreg.predict(X_test)
         print("Accuracy Score :", round(accuracy_score(y_test, y_pred)*100 ,2), "%")
         Accuracy Score : 66.67 %
In [65]: from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score, jaccard_score
         print('F-1 Score : ',(f1_score(y_test, y_pred, average='micro')))
         print('Precision Score : ',(precision_score(y_test, y_pred, average='micro')))
         print('Recall Score : ',(recall_score(y_test, y_pred, average='micro')))
print('Jaccard Score : ',(jaccard_score(y_test, y_pred, average='micro')))
         print('Log Loss : ',(log_loss(y_test, y_pred)))
         F-1 Score: 0.66666666666666
         Precision Score : 0.666666666666666
         Jaccard Score: 0.5
         Log Loss: 11.513058731208593
```

```
In [67]: from sklearn.metrics import confusion_matrix
    cm = confusion_matrix(y_test, y_pred)
    plt.figure(figsize=(5,5))
    sns.heatmap(data=cm,linewidths=.5, annot=True, cmap = 'Blues')
    plt.ylabel('Actual label')
    plt.xlabel('Predicted label')
    all_sample_title = 'Accuracy Score for Logistic Regression: {0}'.format(logreg.score(X_test, y_test))
    plt.title(all_sample_title, size = 15)
```

Out[67]: Text(0.5, 1.0, 'Accuracy Score for Logistic Regression: 0.66666666666666666)





```
In [68]:
    from sklearn.metrics import roc_curve, roc_auc_score
    y_pred_proba = logreg.predict_proba(X_test)[:][:,1]

    df_actual_predicted = pd.concat([pd.DataFrame(np.array(y_test), columns=['y_actual']), pd.DataFramedf_actual_predicted.index = y_test.index

    fpr, tpr, tr = roc_curve(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])
    auc = roc_auc_score(df_actual_predicted['y_actual'], df_actual_predicted['y_pred_proba'])

    plt.plot(fpr, tpr, label='AUC = %0.4f' %auc)
    plt.plot(fpr, fpr, linestyle = '--', color='k')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plt.title('ROC Curve', size = 15)
    plt.legend()
```

Out[68]: <matplotlib.legend.Legend at 0x1aecc934190>

