```
1 import pandas as pd
2 import matplotlib.pyplot as plt
3 import seaborn as sns
4 import numpy as np
5 sns.set_theme(color_codes=True)
1 df = pd.read_csv('dataR2.csv')
2 df
Гэ
        Age
                 BMI Glucose Insulin
                                       HOMA Leptin Adiponectin Resistin MCP.1 Classifica
                       70
     0
        48 23.500000
                              2.707 0.467409 8.8071
                                                         9.702400 7.99585 417.114
                              3.115 0.706897 8.8438
                                                       5.429285 4.06405 468.786
     1
        83 20.690495
                          92
         82 23.124670
                               4.498 1.009651 17.9393
                                                       22.432040 9.27715 554.697
         68 21.367521
                          77
                               3.226 0.612725 9.8827
                                                        7.169560 12.76600 928.220
        86 21.111111
                                                       4.819240 10.57635 773.920
     4
                         92 3.549 0.805386 6.6994
    111 45 26.850000 92 3.330 0.755688 54.6800
                                                       12.100000 10.96000 268.230
    112 62 26.840000 100 4.530 1.117400 12.4500 21.420000 7.32000 330.160
                         97 5.730 1.370998 61.4800
                                                       22.540000 10.33000 314.050
    113 65 32.050000
    114 72 25.590000
                        82
                              2.820 0.570392 24.9600
                                                       33.750000 3.27000 392.460
    115 86 27.180000
                         138
                               19.910 6.777364 90.2800
                                                       14.110000 4.35000 90.090
    116 rows × 10 columns
```

Checking if there is null value

```
1 df.isnull().sum()
    Age
BMI
    Glucose
Insulin
     HOMA
     Leptin
     Adiponectin
     Resistin
     MCP.1
     Classification
     dtype: int64
1 df_copy = df.copy(deep = True)
2 df_copy[['Age','BMI','Glucose','Insulin','HOMA','Leptin','Adiponectin','Resistin','MCP.1']] = df_copy[['Age','BMI','Glucose','Insulin','HOMA','Leptin','Adiponectin','Resistin','MCP.1']]
4 # Showing the Count of NANs
5 print(df_copy.isnull().sum())
    Age
BMI
Glucose
     Insulin
                          0
     HOMA
Leptin
    Adiponectin
Resistin
MCP.1
                          0
     Classification
```

Checking if the class value is balanced or not

```
1 sns.countplot(df['Classification'])
2 print(df.Classification.value_counts())

2 64
1 52
Name: Classification, dtype: int64
/usr/local/lib/python3.8/dist-packages/seaborn/_decorators.py:36: FutureWarning: Pass the followi warnings.warn(

60
50
10
Classification
2
Classification
```

→ Outlier Detection using Boxplot and Outlier Cleansing using Z-Score

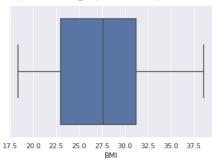
```
1 sns.boxplot(x=df["Age"])
```

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae25feb0>



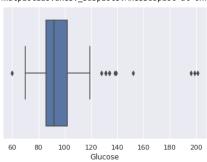
1 sns.boxplot(x=df["BMI"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae204880>



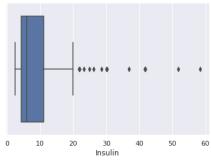
1 sns.boxplot(x=df["Glucose"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae19f610>



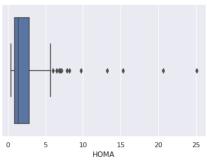
1 sns.boxplot(x=df["Insulin"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5af6aa370>



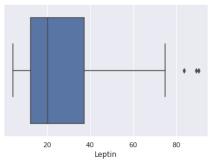
1 sns.boxplot(x=df["HOMA"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae0d2460>



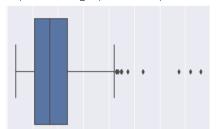
1 sns.boxplot(x=df["Leptin"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae09d4c0>



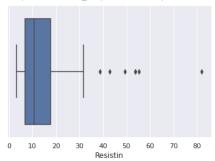
1 sns.boxplot(x=df["Adiponectin"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5ae0318e0>



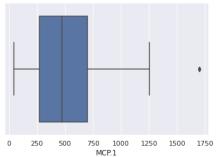
1 sns.boxplot(x=df["Resistin"])

<matplotlib.axes._subplots.AxesSubplot at 0x7fb5adfdebe0>



1 sns.boxplot(x=df["MCP.1"])

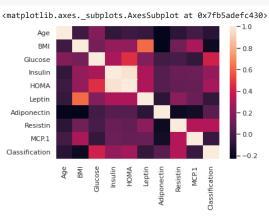
<matplotlib.axes._subplots.AxesSubplot at 0x7fb5adf35b20>



```
1 import scipy.stats as stats
2 z = np.abs(stats.zscore(df))
3 data_clean = df[(z<3).all(axis = 1)]
4 data_clean.shape</pre>
```

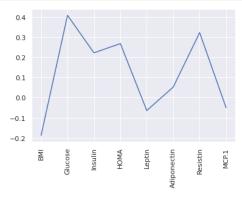
(102, 10)

1 sns.heatmap(data_clean.corr())



Correlation between Class and other attributes

```
1 corr = data_clean[data_clean.columns[1:]].corr()['Classification'][:-1]
2 plt.plot(corr)
3 plt.xticks(rotation=90)
4 plt.show()
```



Machine Learning

```
1 X = data_clean.drop('Classification', axis=1)
2 y = data_clean['Classification']

1 from sklearn.model_selection import train_test_split
2 from sklearn.metrics import accuracy_score
3 X_train, X_test, y_train, y_test = train_test_split(X,y, test_size=0.1,random_state=0)
```

Decision Tree

```
1 from sklearn.tree import DecisionTreeClassifier
2 dtree = DecisionTreeClassifier(random_state = 0)
3 dtree.fit(X_train, y_train)

DecisionTreeClassifier(random_state=0)

1 y_pred = dtree.predict(X_test)
2 print("Accuracy Score :", round(accuracy_score(y_test, y_pred)*100 ,2), "%")

Accuracy Score : 90.91 %

1 from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score
2 print('F-1 Score : ',(f1_score(y_test, y_pred)))
3 print('Precision Score : ',(precision_score(y_test, y_pred)))
4 print('Recall Score : ',(recall_score(y_test, y_pred)))

F-1 Score : 0.8571428571428571
Precision Score : 0.75
Recall Score : 1.0
```

→ Random Forest

Support Vector Machine

Logistic Regression

```
1 from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score
2 print('F-1 Score : ',(f1_score(y_test, y_pred)))
3 print('Precision Score : ',(precision_score(y_test, y_pred)))
4 print('Recall Score : ',(recall_score(y_test, y_pred)))
```

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