CMSE428 Task 1

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1 Task 1

[149]:
```
%matplotlib inline

import pandas as pd
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

1.1 Dataset Loaded in

[150]:
```
dataset = pd.read_csv('dataset.csv')
```

1.2 Original Dataset

[151]:
```
dataset
```

[151]:

	ClassLabel	Gender	Age	Race	Education	FamilyHistory	HighBloodSugar	\
0	Negative	1	44	3	3.0	1	1	
1	Negative	2	44	1	2.0	2	1	
2	Negative	1	22	3	5.0	1	1	
3	Negative	2	35	3	4.0	1	1	
4	Negative	1	54	2	5.0	2	1	
…	…	…	…	…	…	…	…	
1498	Positive	2	59	3	3.0	2	1	
1499	Positive	2	74	3	NaN	2	1	
1500	Positive	2	80	3	2.0	1	1	
1501	Positive	2	78	3	4.0	2	1	
1502	Positive	1	80	2	3.0	1	1	

	BMI	WaistCircumference	SystolicBP	DiastolicBP	LDL	HDL	\
0	29.10	106.6	119	88	113	37	
1	25.06	82.3	130	86	138	63	
2	28.07	94.5	120	87	96	57	
3	20.29	71.9	110	64	93	89	

```
4     36.32                116.4           92            42  110   62
…     …                  …          …           …   …   …
1498  26.67                103.0          110            60  132   47
1499  25.08                 95.5          135            60  114   36
1500    NaN                119.3          121            40  111   38
1501  46.98                129.5          146            78  136   56
1502  27.14                 96.7          154            30  128   65

      Triglycerides
0               235
1                61
2               111
3                58
4                71
…               …
1498            190
1499            270
1500            131
1501            144
1502            108

[1503 rows x 14 columns]
```

1.3 Checking which columns have missing Data

By running the below code, we can see that Education and BMI has missing data.

[152]:
```
dataset.isnull().sum()
```

[152]:
```
ClassLabel            0
Gender                0
Age                   0
Race                  0
Education            12
FamilyHistory         0
HighBloodSugar        0
BMI                   9
WaistCircumference    0
SystolicBP            0
DiastolicBP           0
LDL                   0
HDL                   0
Triglycerides         0
dtype: int64
```

1.4 For example, 150th row of BMI has no data

[153]:
```
dataset.loc[150,'BMI']
```

[153]:
```
nan
```

1.5 Filling in missing BMI data by Mean and Education by Most Frequent Value

BMI value can be a float, so we can use mean to impute missing data.By running the print command, we can see that the missing values are filled in with mean.

[154]:
```
dataset['BMI'].fillna(dataset['BMI'].mean(), inplace=True)
dataset['Education'].fillna(dataset['Education'].value_counts().index[0],␣
↪inplace=True)
dataset.isnull().sum()
```

[154]:
```
ClassLabel            0
Gender                0
Age                   0
Race                  0
Education             0
FamilyHistory         0
HighBloodSugar        0
BMI                   0
WaistCircumference    0
SystolicBP            0
DiastolicBP           0
LDL                   0
HDL                   0
Triglycerides         0
dtype: int64
```

1.6 Checking the 150th row again

[155]:
```
dataset.loc[150,'BMI']
```

[155]:
```
28.41636546184739
```

1.7 Number of Positives and Negatives

[156]:
```
dataset['ClassLabel'].value_counts()
```

[156]:
```
Negative    775
Positive    728
Name: ClassLabel, dtype: int64
```

1.8 Number of Positives and Gender==1 and Race==2

[157]:
```
len(dataset[(dataset['ClassLabel']=='Positive') & (dataset['Gender']==1) &␣
↪(dataset['Race']==2)])
```

[157]:
```
40
```

1.9 Race that is most frequent in Negative samples

[158]:
```
negativeData = dataset[(dataset['ClassLabel']=='Negative')]
negativeData
```

[158]:
```
    ClassLabel  Gender  Age  Race  Education  FamilyHistory  HighBloodSugar  \
0     Negative       1   44     3        3.0              1               1
1     Negative       2   44     1        2.0              2               1
2     Negative       1   22     3        5.0              1               1
3     Negative       2   35     3        4.0              1               1
4     Negative       1   54     2        5.0              2               1
..         …     …   …   …   …         …              …               …
770   Negative       2   53     2        5.0              2               1
771   Negative       1   56     3        5.0              1               1
772   Negative       2   67     1        4.0              2               1
773   Negative       2   34     2        1.0              1               1
774   Negative       2   48     3        5.0              1               1

       BMI  WaistCircumference  SystolicBP  DiastolicBP  LDL  HDL  \
0    29.10                106.6         119           88  113   37
1    25.06                 82.3         130           86  138   63
2    28.07                 94.5         120           87   96   57
3    20.29                 71.9         110           64   93   89
4    36.32                116.4          92           42  110   62
..     …                    …           …            …   …   …
770  20.09                 75.0          94           60  148   67
771  30.89                113.1         120           83  139   65
772  20.37                 71.6         150           80   95   60
773  38.74                117.4         107           67   91   34
774  21.66                 71.5         102           61   88   72

     Triglycerides
0              235
1               61
2              111
3               58
4               71
..              …
770             72
771             84
772            138
```

```
773             60
774             50

[775 rows x 14 columns]
```

[159]:
```
negativeData['Race'].value_counts()
```

[159]:
```
3    404
1    127
4    117
2     86
5     41
Name: Race, dtype: int64
```

1.10 Average BMI of samples with Education==3

[160]:
```
education_ft = dataset[(dataset['Education']==3)]
education_ft['BMI'].mean()
```

[160]:
```
29.276166255235008
```

1.11 Average BMI of Gender==1

[161]:
```
gender_filter = dataset[(dataset['Gender']==1)]
gender_filter['BMI'].mean()
```

[161]:
```
28.49773850117451
```

1.12 Average BMI of Gender==2

[162]:
```
gender_filter = dataset[(dataset['Gender']==2)]
gender_filter['BMI'].mean()
```

[162]:
```
28.347488282367802
```

2 Task 2

2.1 Seperating Categorical and Numerical

Because that datatypes of columns are given not correctly, the columns are seperated manually.

Seperating Categorical Columns

[163]:
```
categorical_dataset =␣
↪dataset[['ClassLabel','Gender','Race','Education','FamilyHistory','HighBloodSugar']]
categorical_dataset
```

[163]:
```
      ClassLabel  Gender  Race  Education  FamilyHistory  HighBloodSugar
0       Negative       1     3        3.0              1               1
1       Negative       2     1        2.0              2               1
2       Negative       1     3        5.0              1               1
3       Negative       2     3        4.0              1               1
4       Negative       1     2        5.0              2               1
…          …     …   …        …             …              …
1498    Positive       2     3        3.0              2               1
1499    Positive       2     3        4.0              2               1
1500    Positive       2     3        2.0              1               1
1501    Positive       2     3        4.0              2               1
1502    Positive       1     2        3.0              1               1

[1503 rows x 6 columns]
```

Seperating Numerical Values

[164]:
```
numerical_dataset =␣
↪dataset[['Age','BMI','WaistCircumference','SystolicBP','DiastolicBP','LDL','HDL','Triglycer
numerical_dataset
```

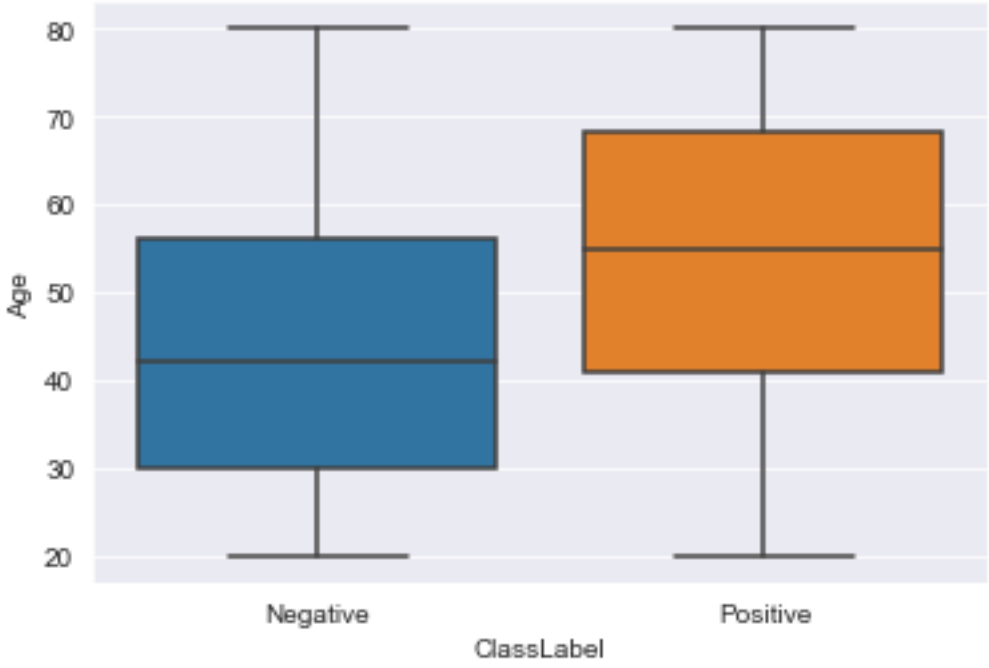
[164]:
```
       Age        BMI  WaistCircumference  SystolicBP  DiastolicBP  LDL  HDL  \
0       44  29.100000               106.6         119           88  113   37
1       44  25.060000                82.3         130           86  138   63
2       22  28.070000                94.5         120           87   96   57
3       35  20.290000                71.9         110           64   93   89
4       54  36.320000               116.4          92           42  110   62
…   …         …                   …          …            …  …   …
1498    59  26.670000               103.0         110           60  132   47
1499    74  25.080000                95.5         135           60  114   36
1500    80  28.416365               119.3         121           40  111   38
1501    78  46.980000               129.5         146           78  136   56
1502    80  27.140000                96.7         154           30  128   65

      Triglycerides
0               235
1                61
2               111
3                58
4                71
…               …
1498            190
1499            270
1500            131
1501            144
1502            108

[1503 rows x 8 columns]
```

2.2 Boxplots for Classes with respect to each Variable

Age

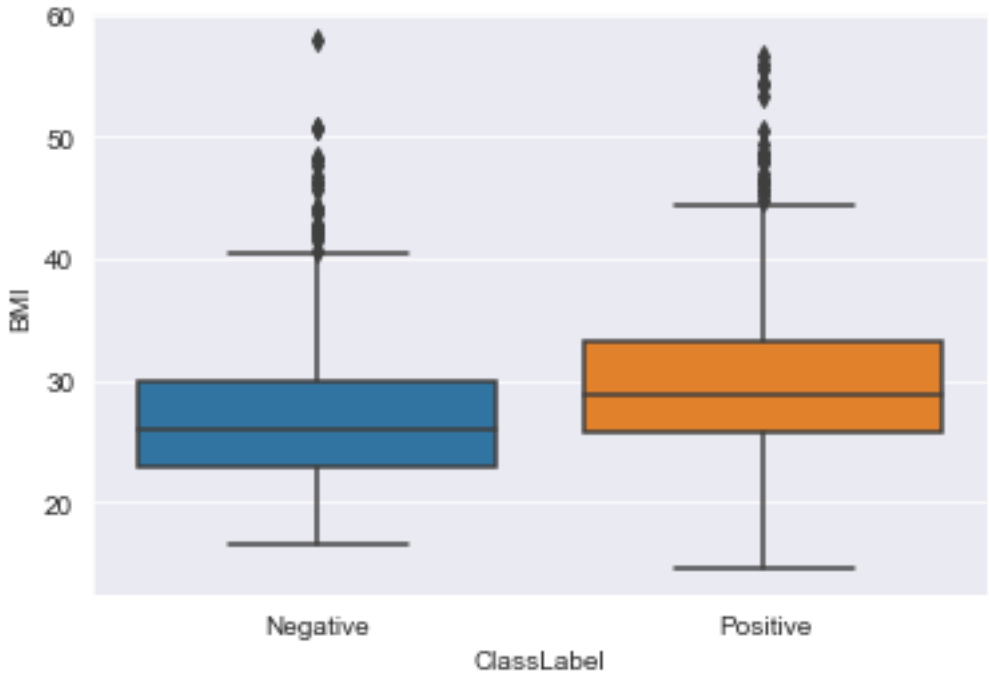
[165]:
```
sns.set_style("darkgrid")
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[0])
```

[165]:
```
<AxesSubplot:xlabel='ClassLabel', ylabel='Age'>
```



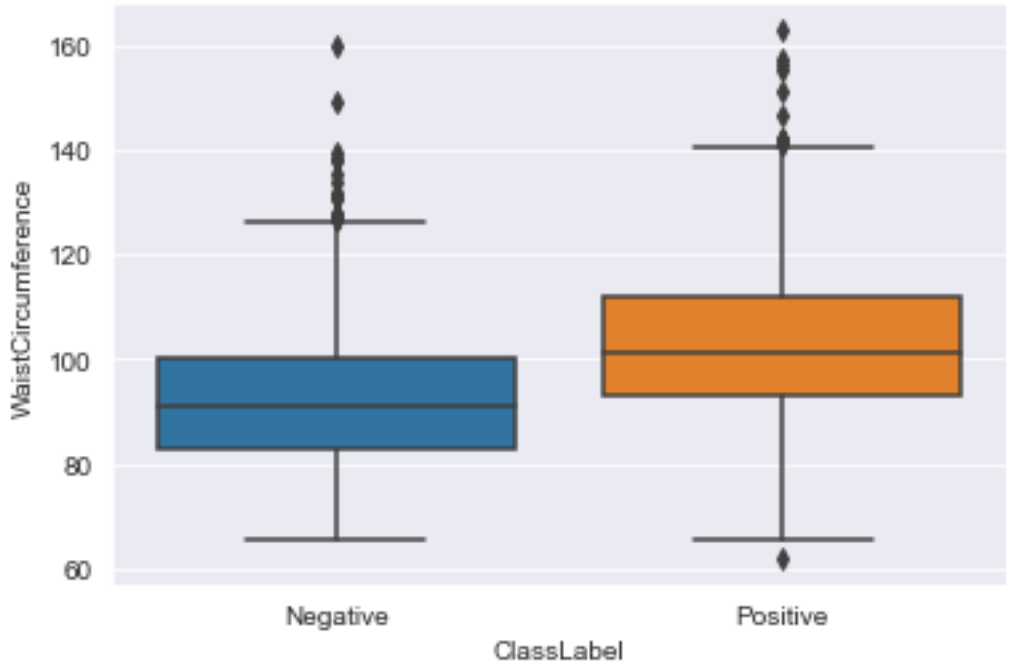
[166]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[1])
```

[166]:
```
<AxesSubplot:xlabel='ClassLabel', ylabel='BMI'>
```



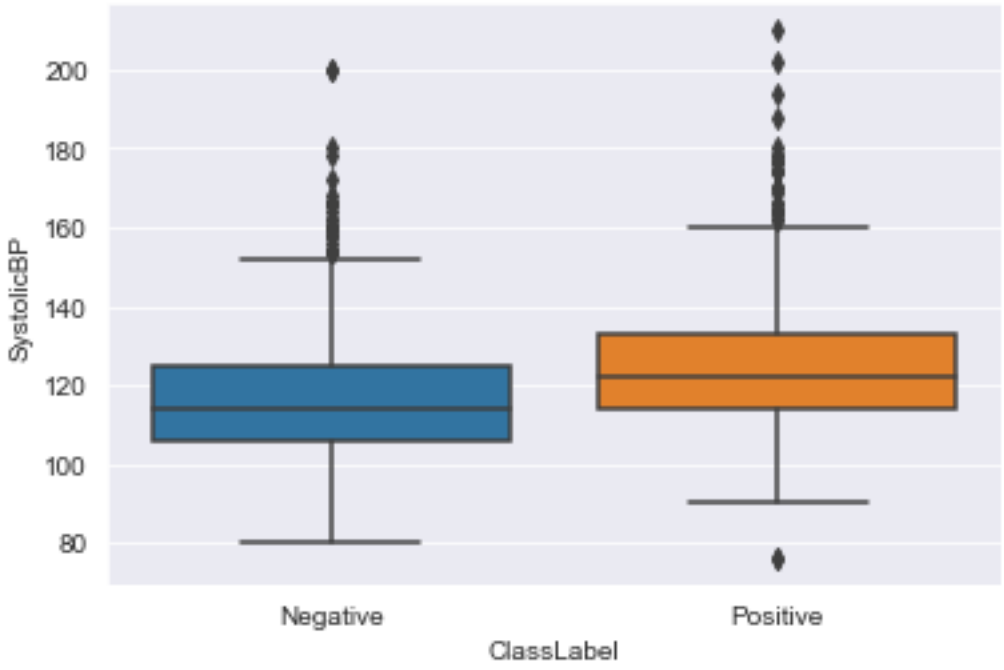
[167]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[2])
```

[167]:
```
<AxesSubplot:xlabel='ClassLabel', ylabel='WaistCircumference'>
```



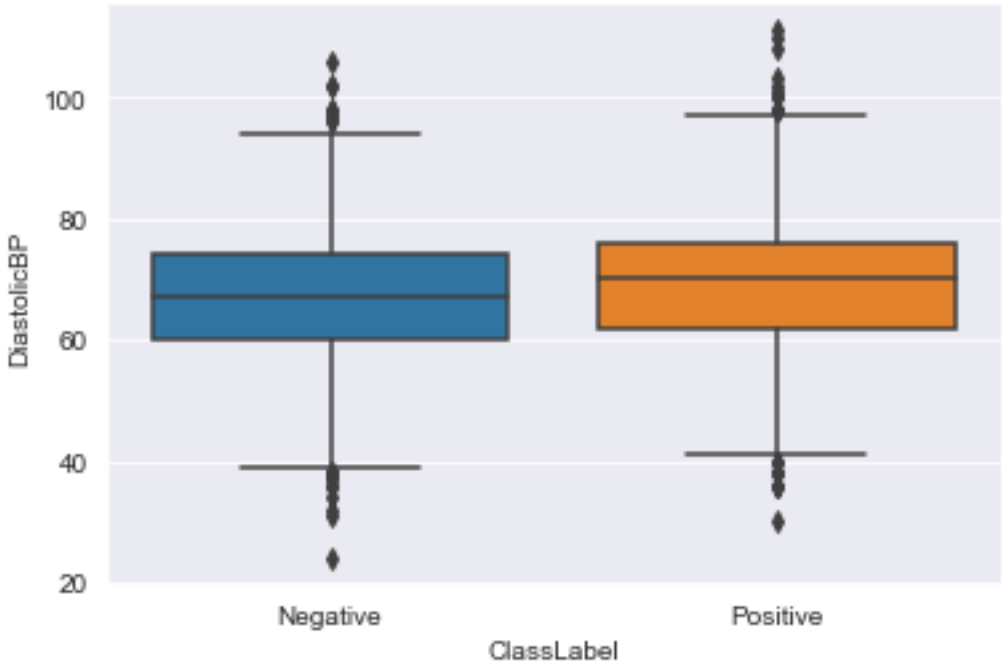
[168]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[3])
```

[168]: <AxesSubplot:xlabel='ClassLabel', ylabel='SystolicBP'>



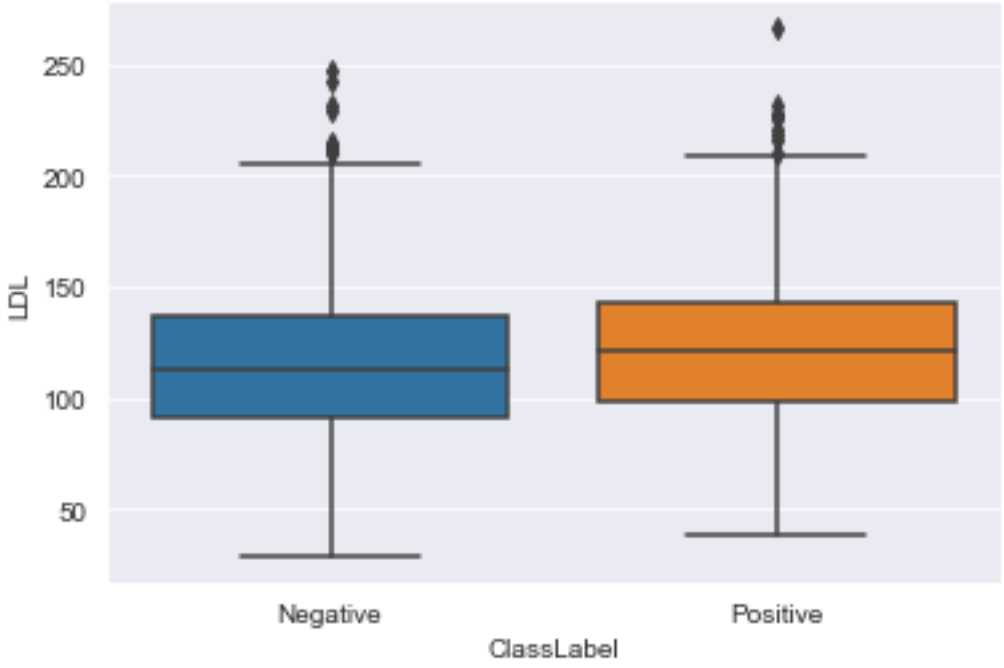
[169]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[4])
```

[169]: <AxesSubplot:xlabel='ClassLabel', ylabel='DiastolicBP'>



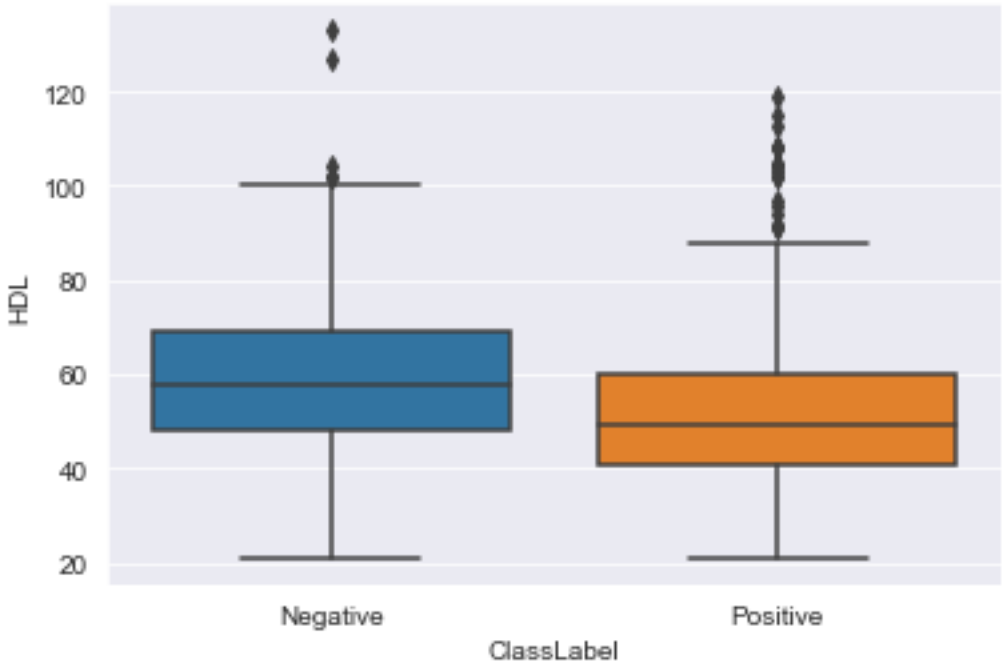
[170]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[5])
```

[170]: <AxesSubplot:xlabel='ClassLabel', ylabel='LDL'>



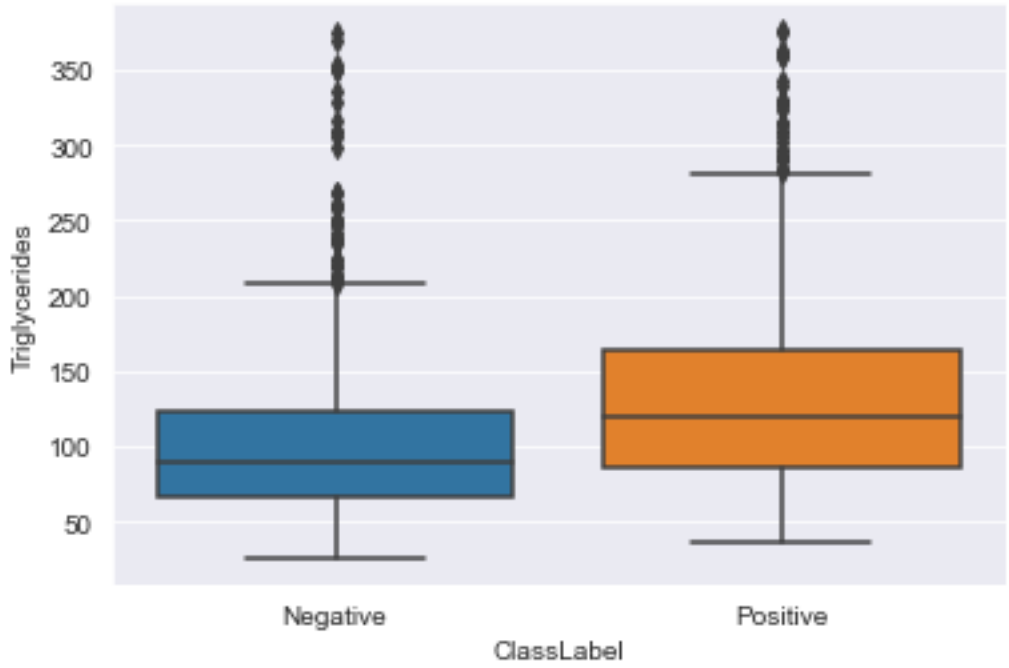
[171]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[6])
```

[171]: <AxesSubplot:xlabel='ClassLabel', ylabel='HDL'>



[172]:
```
sns.boxplot(data=dataset, x='ClassLabel', y=numerical_dataset.columns[7])
```

[172]: <AxesSubplot:xlabel='ClassLabel', ylabel='Triglycerides'>



2.3 Table of Quartiles (Q1,Q2/Median,Q3) for each numerical value

[173]:
```
q1_list = [] ; q2_list = [] ; q3_list = []

for column in numerical_dataset:
    q1 = dataset[column].quantile(0.25)
    q2 = dataset[column].mean()
    q3 = dataset[column].quantile(0.75)
    q1_list.append(q1)
    q2_list.append(q2)
    q3_list.append(q3)

quartiles = {}
quartiles['index']=numerical_dataset.columns.tolist()
quartiles['Q1']=q1_list
quartiles['Q2']=q2_list
quartiles['Q3']=q3_list

quartiles_df = pd.DataFrame(quartiles, index=quartiles['index'],␣
 ↪columns=['Q1','Q2','Q3'])
quartiles_df
```

[173]:

	Q1	Q2	Q3
Age	34.00	49.231537	63.000

```
BMI                  24.16   28.416365   31.715
WaistCircumference   86.50   97.555888  106.350
SystolicBP          108.00  120.359947  130.000
DiastolicBP          61.00   68.152362   76.000
LDL                  94.00  118.815702  140.000
HDL                  44.00   55.480373   65.000
Triglycerides        75.00  117.979375  146.000
```

2.4 Number of Outliers for each continous variable

[174]:
```
print('---Amount of Outliers---')
for index in quartiles_df.index:
    q1 = quartiles_df.loc[index,'Q1']
    q2 = quartiles_df.loc[index,'Q2']
    q3 = quartiles_df.loc[index,'Q3']

    iqr = q3 - q1

    upper_limit = q3 + 1.5*iqr
    lower_limit = q1 - 1.5*iqr

    print (index+': '+str(len(dataset[(dataset[index]> upper_limit) |␣
 ↪(dataset[index] < lower_limit)])))
```

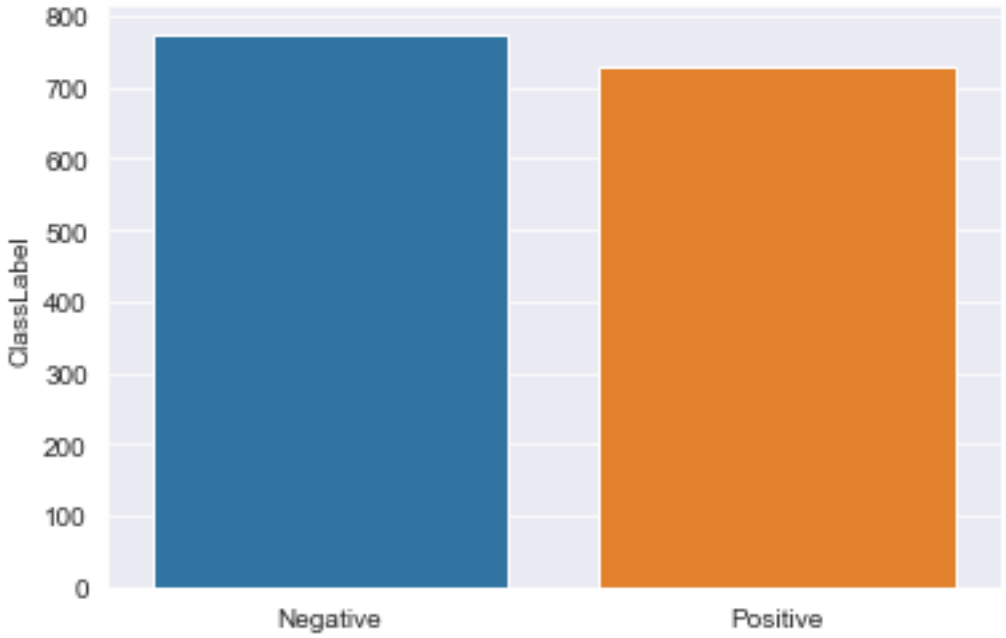
```
---Amount of Outliers---
Age: 0
BMI: 42
WaistCircumference: 22
SystolicBP: 32
DiastolicBP: 30
LDL: 19
HDL: 24
Triglycerides: 59
```

3 Task 3

3.1 BarPlot for each categorical variable

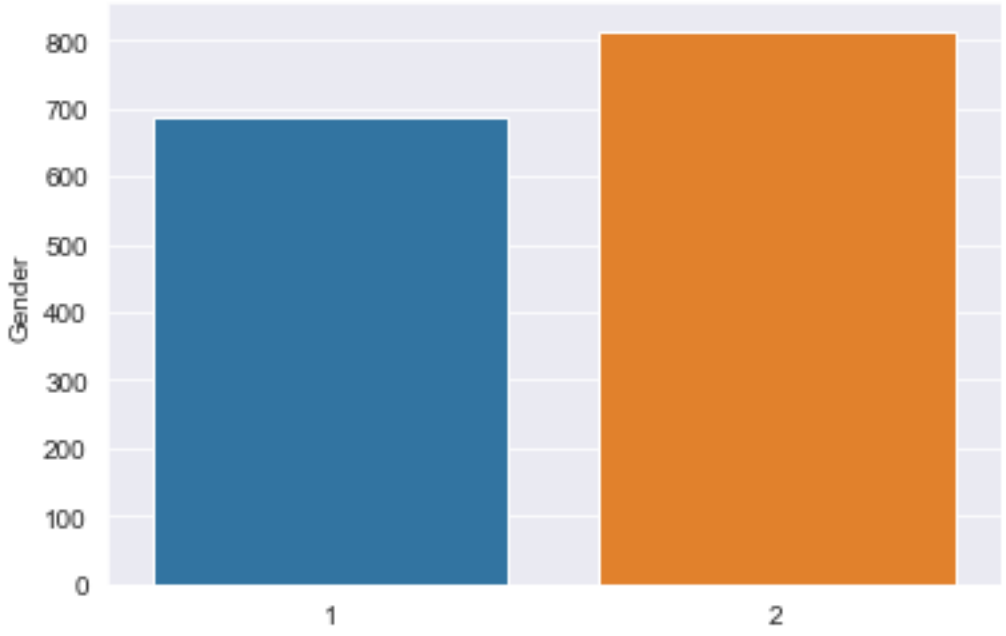
[175]:
```
sns.barplot(x=categorical_dataset.iloc[:, 0].value_counts().index,␣
 ↪y=categorical_dataset.iloc[:, 0].value_counts())
```

[175]:
```
<AxesSubplot:ylabel='ClassLabel'>
```



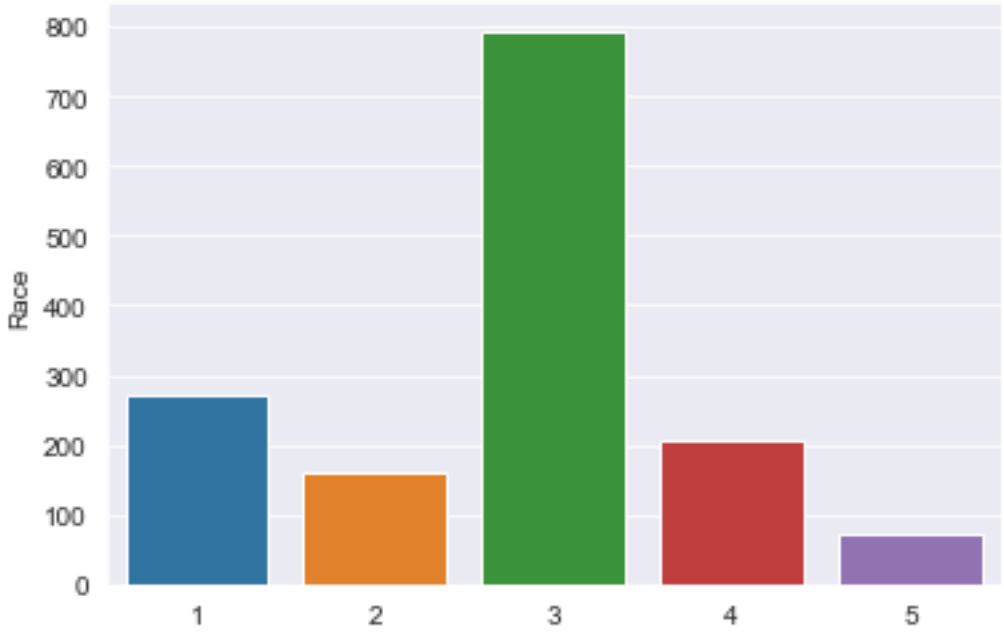
[176]:
```
sns.barplot(x=categorical_dataset.iloc[:, 1].value_counts().index,␣
↪y=categorical_dataset.iloc[:, 1].value_counts())
```

[176]: <AxesSubplot:ylabel='Gender'>



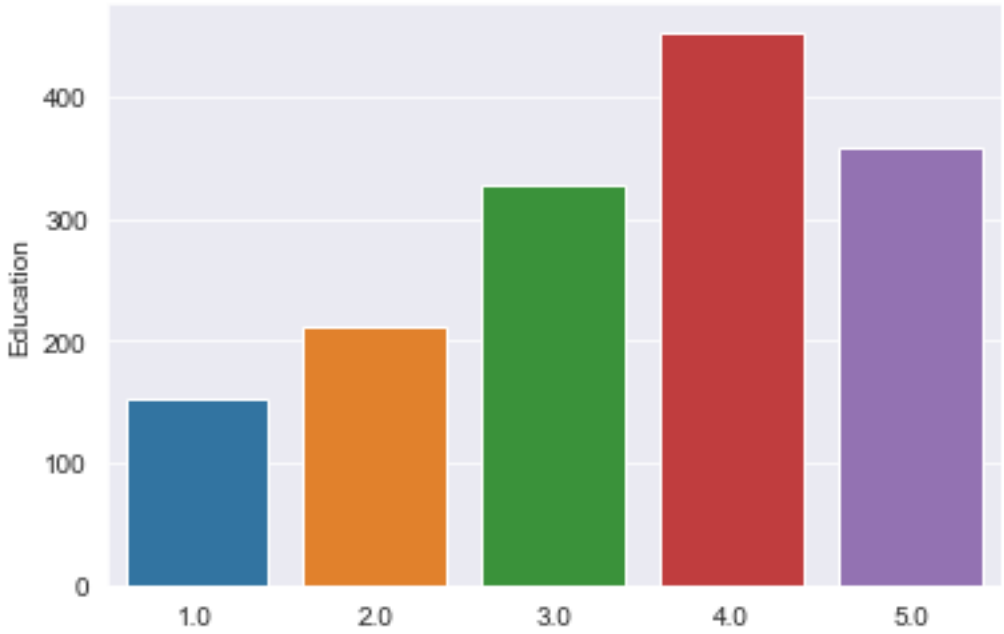
[177]:
```
sns.barplot(x=categorical_dataset.iloc[:, 2].value_counts().index,␣
↪y=categorical_dataset.iloc[:, 2].value_counts())
```

[177]: <AxesSubplot:ylabel='Race'>



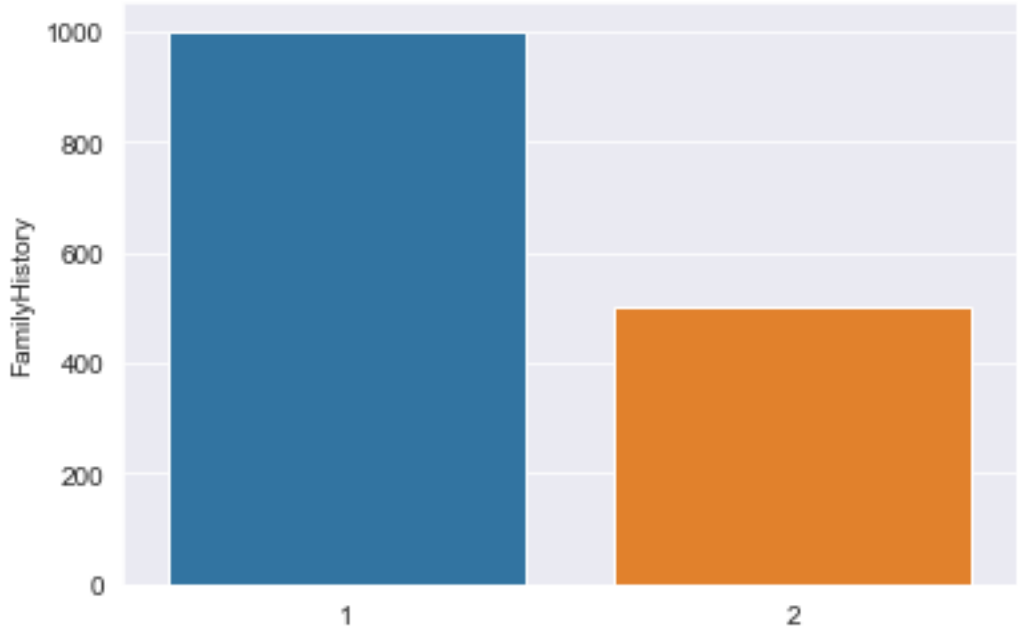
[178]:
```
sns.barplot(x=categorical_dataset.iloc[:, 3].value_counts().index,␣
↪y=categorical_dataset.iloc[:, 3].value_counts())
```

[178]: <AxesSubplot:ylabel='Education'>



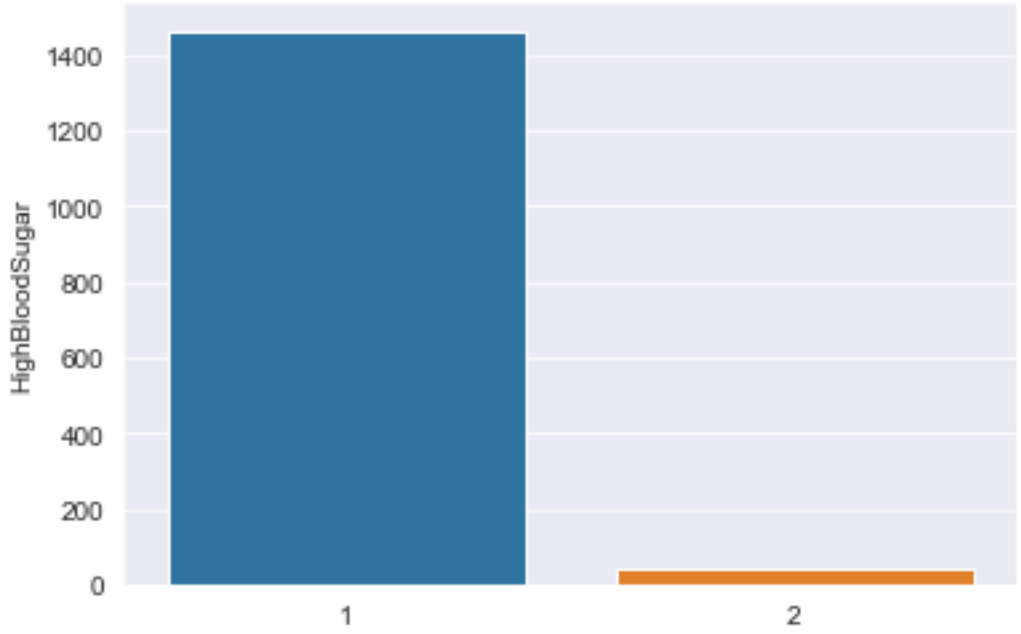
[179]:
```
sns.barplot(x=categorical_dataset.iloc[:, 4].value_counts().index,␣
↪y=categorical_dataset.iloc[:, 4].value_counts())
```

[179]: <AxesSubplot:ylabel='FamilyHistory'>



[180]:
```
sns.barplot(x=categorical_dataset.iloc[:, 5].value_counts().index,␣
↪y=categorical_dataset.iloc[:, 5].value_counts())
```

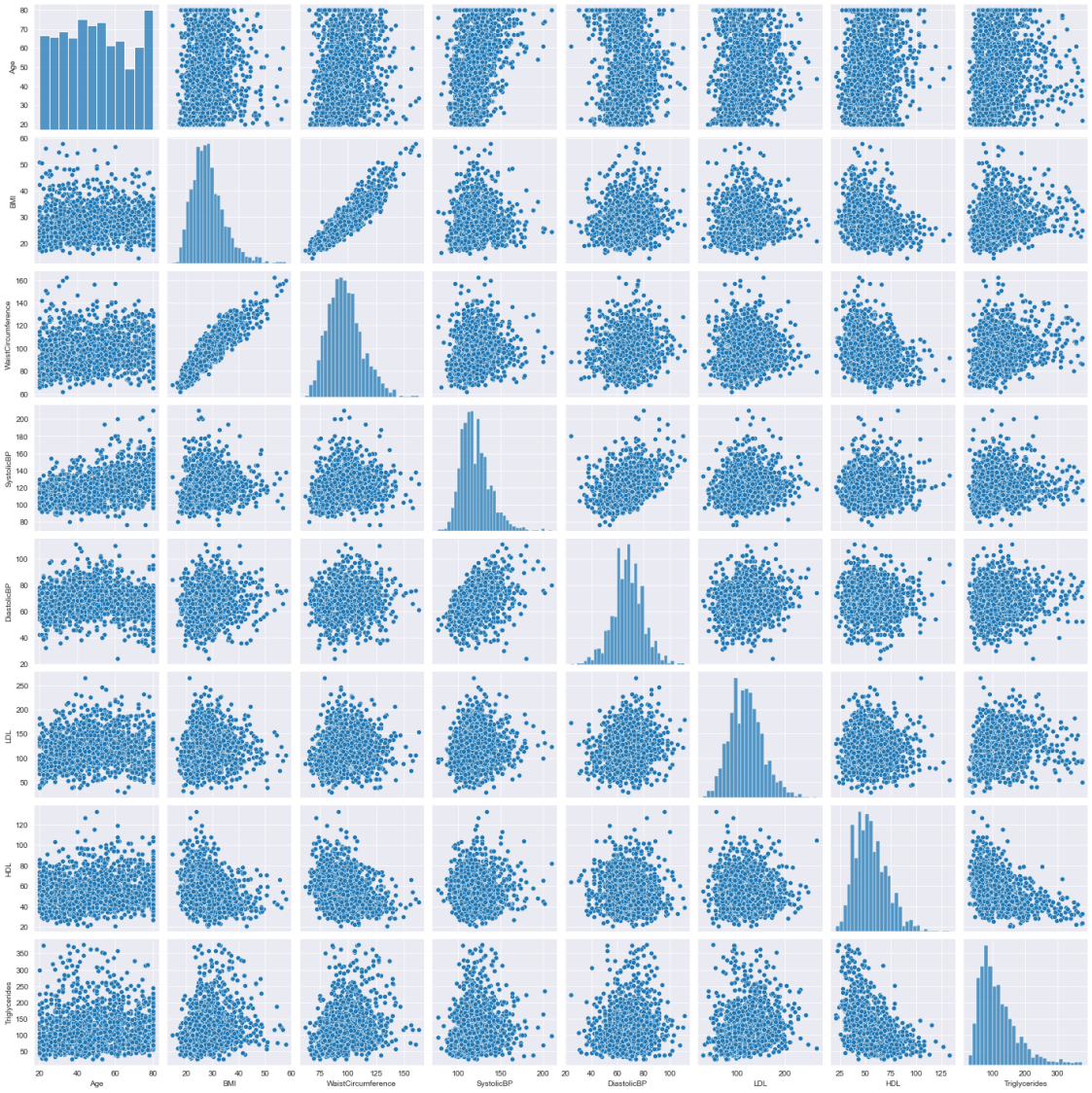
[180]: <AxesSubplot:ylabel='HighBloodSugar'>



4 Task 4

[181]:
```
sns.pairplot(numerical_dataset)
```

[181]: <seaborn.axisgrid.PairGrid at 0x16891c8bbb0>


Age
BMI
WaistCircumference
SystolicBP
DiastolicBP
LDL
HDL
Triglycerides