

```
In [2]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn import svm
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
```

```
In [4]: df=pd.read_csv('train.csv')
df.head()
```

```
Out[4]:
```

	baseline value	accelerations	fetal_movement	uterine_contractions	light_decelerations	severe_decelerations	prolongued_decelerations	abnormal_short_tern
0	142.0	0.000	0.000	0.007	0.000	0.0	0.0	
1	122.0	0.000	0.000	0.006	0.002	0.0	0.0	
2	129.0	0.005	0.003	0.001	0.000	0.0	0.0	
3	136.0	0.006	0.000	0.008	0.000	0.0	0.0	
4	144.0	0.000	0.000	0.006	0.000	0.0	0.0	

5 rows × 22 columns

```
In [5]: df.info()
```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1700 entries, 0 to 1699
Data columns (total 22 columns):
#   Column                                     Non-Null Count  Dtype
---  -
0   baseline value                           1700 non-null   float64
1   accelerations                           1700 non-null   float64
2   fetal_movement                          1700 non-null   float64
3   uterine_contractions                    1700 non-null   float64
4   light_decelerations                     1700 non-null   float64
5   severe_decelerations                    1700 non-null   float64
6   prolonged_decelerations                  1700 non-null   float64
7   abnormal_short_term_variability          1700 non-null   float64
8   mean_value_of_short_term_variability     1700 non-null   float64
9   percentage_of_time_with_abnormal_long_term_variability 1700 non-null   float64
10  mean_value_of_long_term_variability       1700 non-null   float64
11  histogram_width                           1700 non-null   float64
12  histogram_min                             1700 non-null   float64
13  histogram_max                             1700 non-null   float64
14  histogram_number_of_peaks                 1700 non-null   float64
15  histogram_number_of_zeroes                1700 non-null   float64
16  histogram_mode                            1700 non-null   float64
17  histogram_mean                            1700 non-null   float64
18  histogram_median                          1700 non-null   float64
19  histogram_variance                        1700 non-null   float64
20  histogram_tendency                        1700 non-null   float64
21  fetal_health                             1700 non-null   float64
dtypes: float64(22)
memory usage: 292.3 KB

```

```
In [6]: df.shape
```

```
Out[6]: (1700, 22)
```

```
In [7]: df.isna().sum()
```

```
Out[7]: baseline value      0
accelerations      0
fetal_movement     0
uterine_contractions 0
light_decelerations 0
severe_decelerations 0
prolongued_decelerations 0
abnormal_short_term_variability 0
mean_value_of_short_term_variability 0
percentage_of_time_with_abnormal_long_term_variability 0
mean_value_of_long_term_variability 0
histogram_width    0
histogram_min      0
histogram_max      0
histogram_number_of_peaks 0
histogram_number_of_zeroes 0
histogram_mode      0
histogram_mean      0
histogram_median    0
histogram_variance  0
histogram_tendency  0
fetal_health        0
dtype: int64
```

```
In [8]: df.describe()
```

Out[8]:

	baseline value	accelerations	fetal_movement	uterine_contractions	light_decelerations	severe_decelerations	prolongued_decelerations	abnormal_st
count	1700.000000	1700.000000	1700.000000	1700.000000	1700.000000	1700.000000	1700.000000	
mean	133.213529	0.003212	0.010211	0.004356	0.001899	0.000004	0.000158	
std	9.873344	0.003888	0.050124	0.002943	0.002976	0.000059	0.000587	
min	106.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	
25%	126.000000	0.000000	0.000000	0.002000	0.000000	0.000000	0.000000	
50%	133.000000	0.002000	0.000000	0.004000	0.000000	0.000000	0.000000	
75%	140.000000	0.006000	0.003000	0.006000	0.003000	0.000000	0.000000	
max	159.000000	0.019000	0.481000	0.015000	0.015000	0.001000	0.005000	

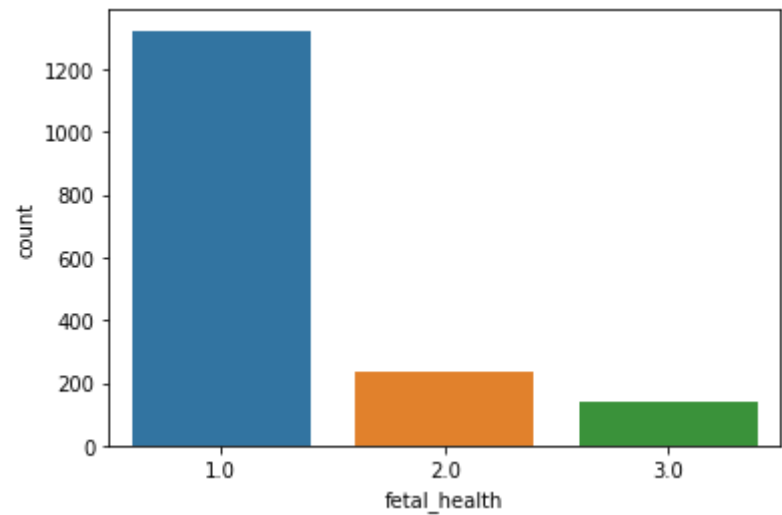
8 rows × 22 columns

In [9]:

```
#To understand the different types in our target variable (fatal_health)
sns.countplot(df.fetal_health)
```

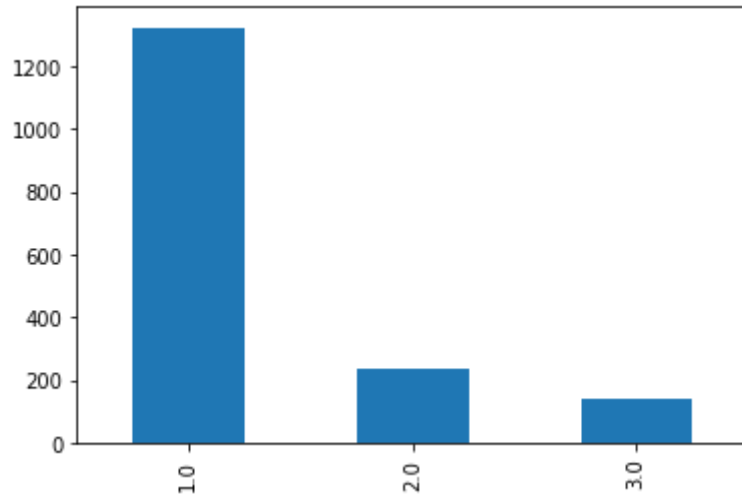
Out[9]:

<AxesSubplot:xlabel='fetal\_health', ylabel='count'>



```
In [10]: df['fetal_health'].value_counts().plot(kind='bar')
```

Out[10]: <AxesSubplot:>

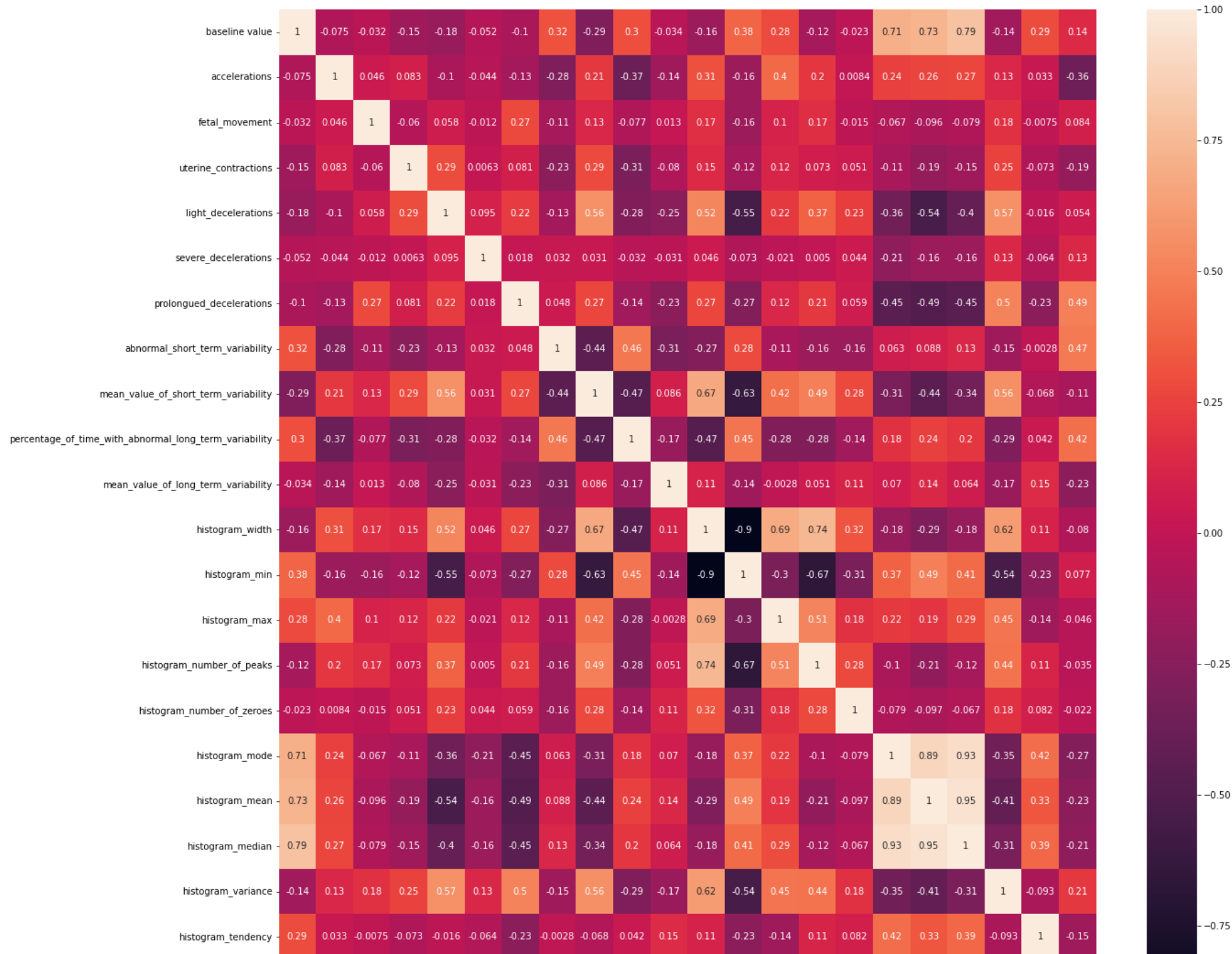


```
In [12]: #to find all datatypes in our file
cats = list(df.select_dtypes(include=['object','bool']))
nums = list(df.select_dtypes(include=['int64','float64']))
print(cats)
print(nums)
[]
['baseline_value', 'accelerations', 'fetal_movement',
'uterine_contractions', 'light_decelerations', 'severe_decelerations',
'prolongued_decelerations', 'abnormal_short_term_variability',
'mean_value_of_short_term_variability',
'percentage_of_time_with_abnormal_long_term_variability',
'mean_value_of_long_term_variability', 'histogram_width',
'histogram_min', 'histogram_max', 'histogram_number_of_peaks',
'histogram_number_of_zeroes', 'histogram_mode', 'histogram_mean',
'histogram_median', 'histogram_variance', 'histogram_tendency',
'fetal_health']
#splitting the dataset X,y
X=df.iloc[:, :-1].values
y=df.iloc[:, -1].values
from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.3,random_state=21)
```

```
[]  
['baseline value', 'accelerations', 'fetal_movement', 'uterine_contractions', 'light_decelerations', 'severe_decelerations', 'prolonged_decelerations', 'abnormal_short_term_variability', 'mean_value_of_short_term_variability', 'percentage_of_time_with_abnormal_long_term_variability', 'mean_value_of_long_term_variability', 'histogram_width', 'histogram_min', 'histogram_max', 'histogram_number_of_peaks', 'histogram_number_of_zeroes', 'histogram_mode', 'histogram_mean', 'histogram_median', 'histogram_variance', 'histogram_tendency', 'fetal_health']
```

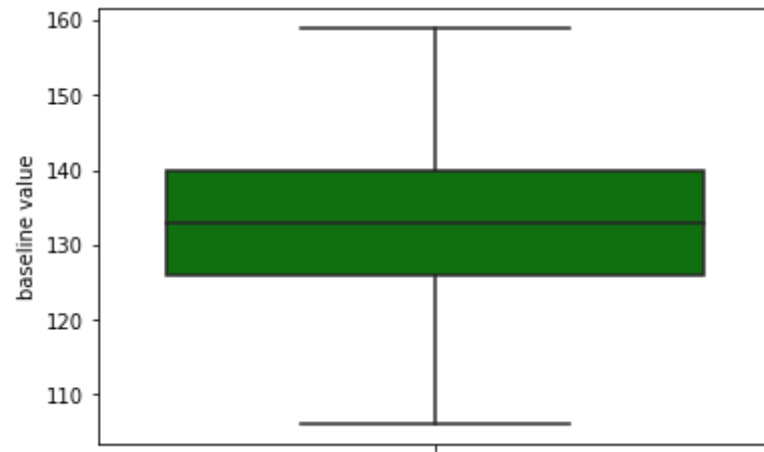
```
In [13]: #Finding the correlation of df  
corr=df.corr()  
plt.figure(figsize=(20,20))  
sns.heatmap(corr,annot=True)
```

```
Out[13]: <AxesSubplot:>
```

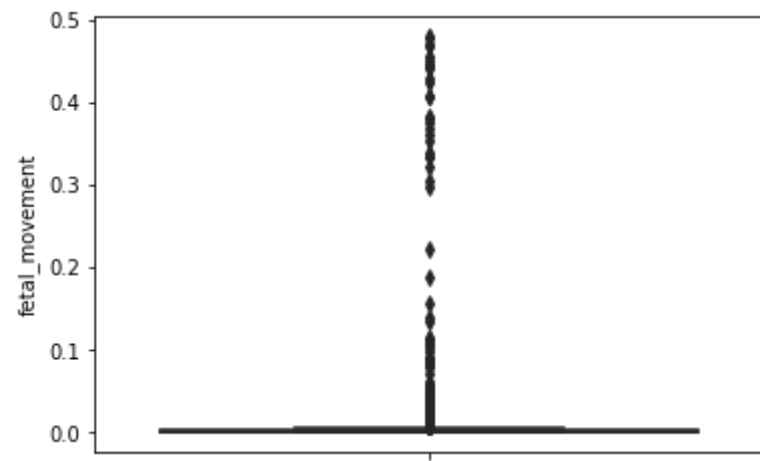
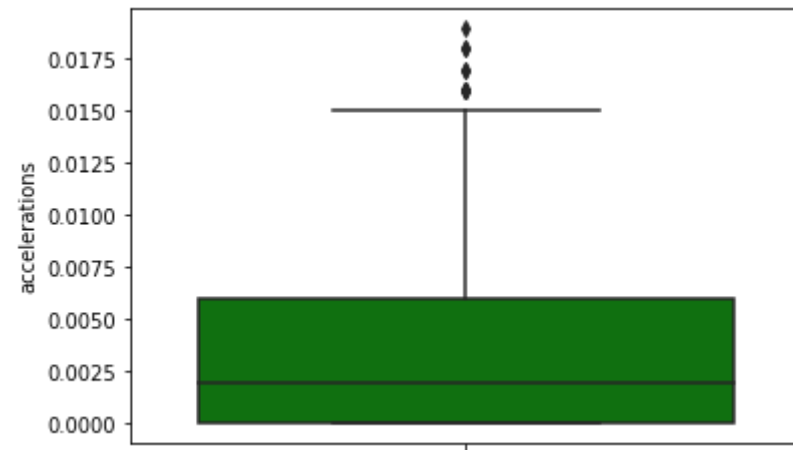


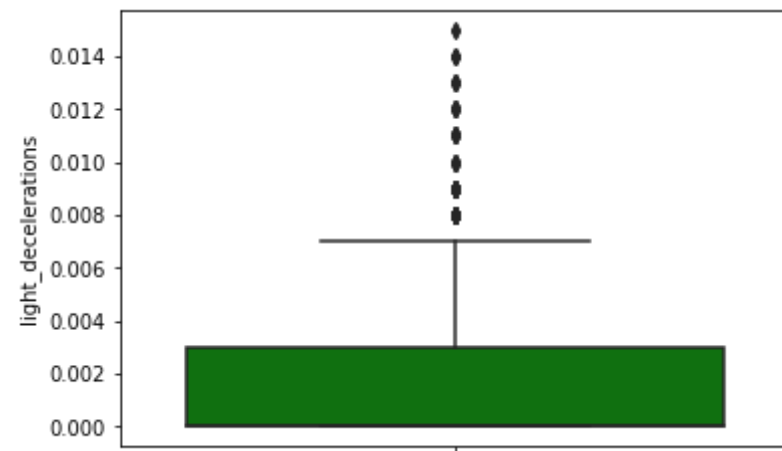
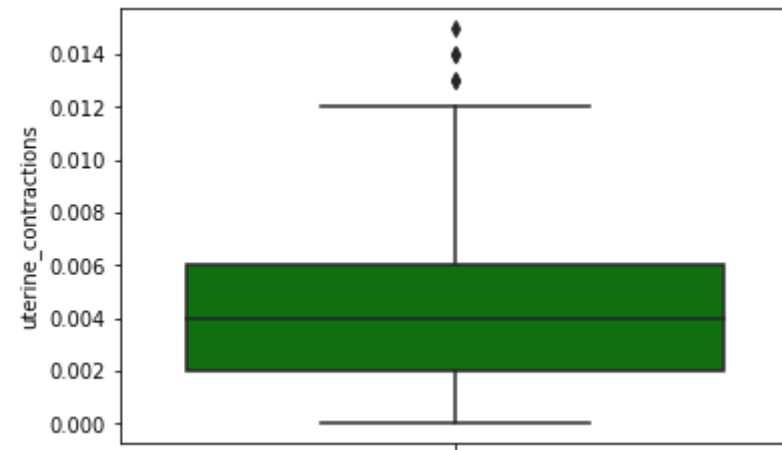
fetal_health	0.14	-0.36	0.084	-0.19	0.054	0.13	0.49	0.47	-0.11	0.42	-0.23	-0.08	0.077	-0.046	-0.035	-0.022	-0.27	-0.23	-0.21	0.21	-0.15	1
	baseline_value	accelerations	fetal_movement	uterine_contractions	light_decelerations	severe_decelerations	prolongued_decelerations	abnormal_short_term_variability	mean_value_of_short_term_variability	percentage_of_time_with_abnormal_long_term_variability	mean_value_of_long_term_variability	histogram_width	histogram_min	histogram_max	histogram_number_of_peaks	histogram_number_of_zeroes	histogram_mode	histogram_mean	histogram_median	histogram_variance	histogram_tendency	fetal_health

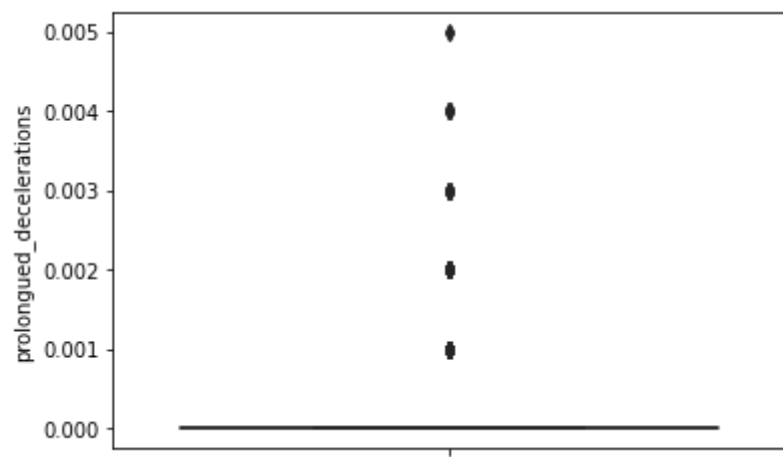
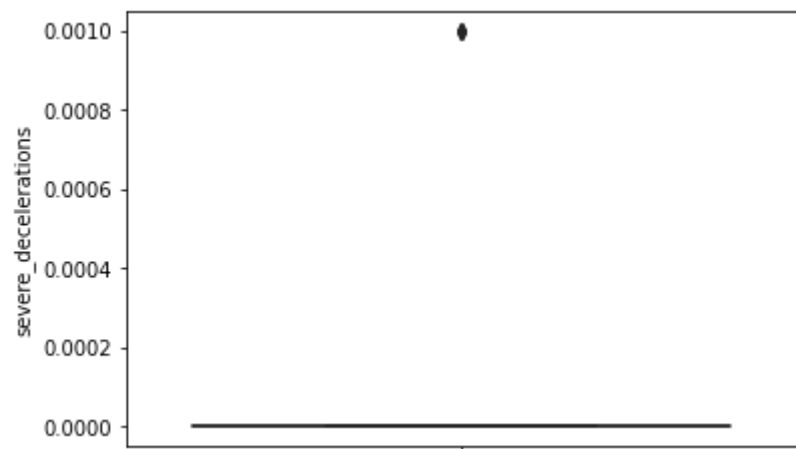
```
In [14]: #Finding the outlier using boxplot
for i in range(0, len(nums )):
    sns.boxplot(y=df[nums[i]],color='green',orient='v')
plt.show()
```

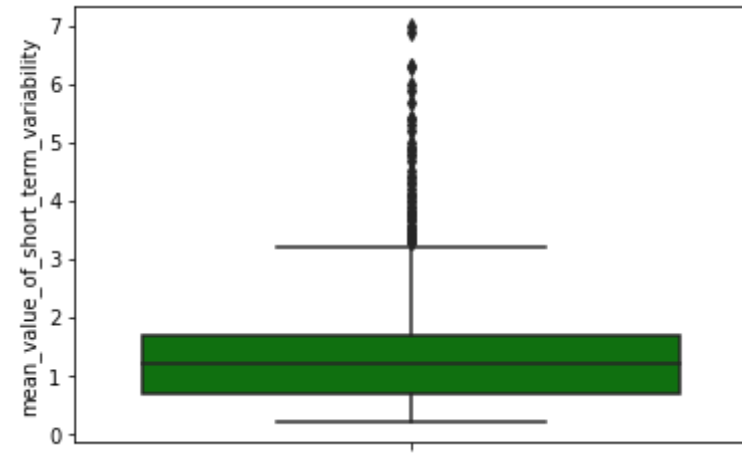
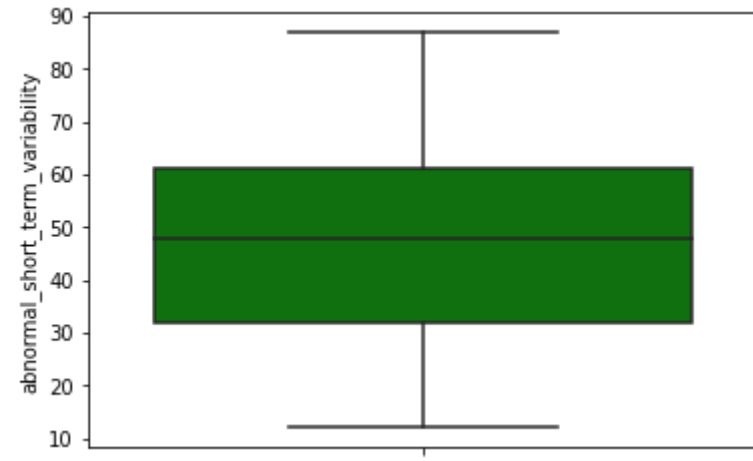




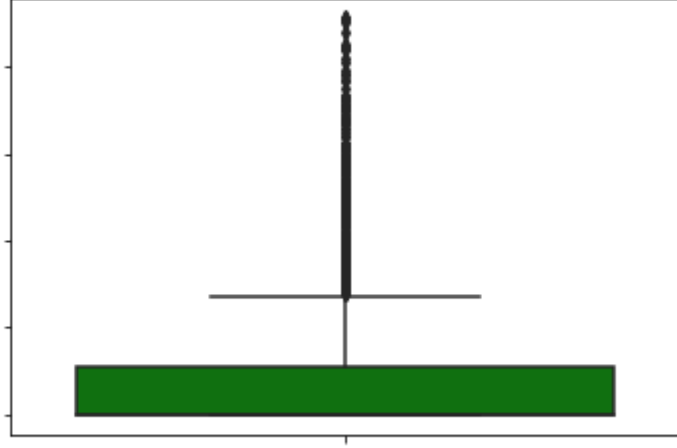




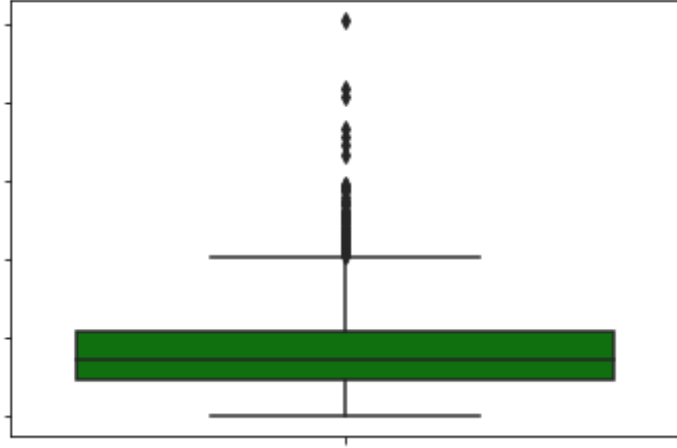


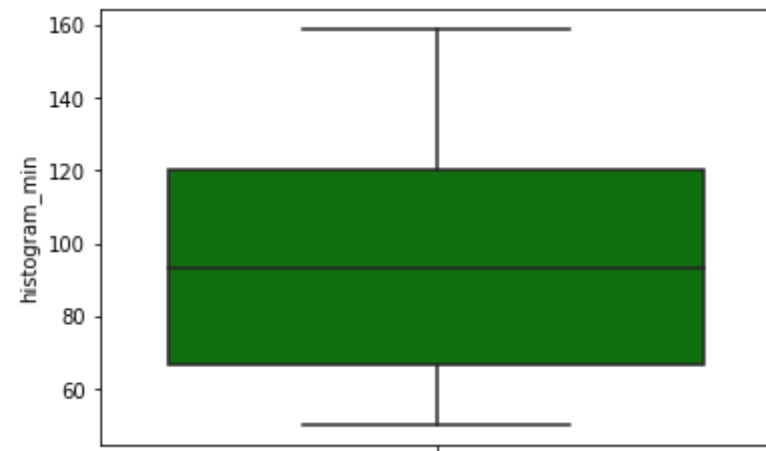
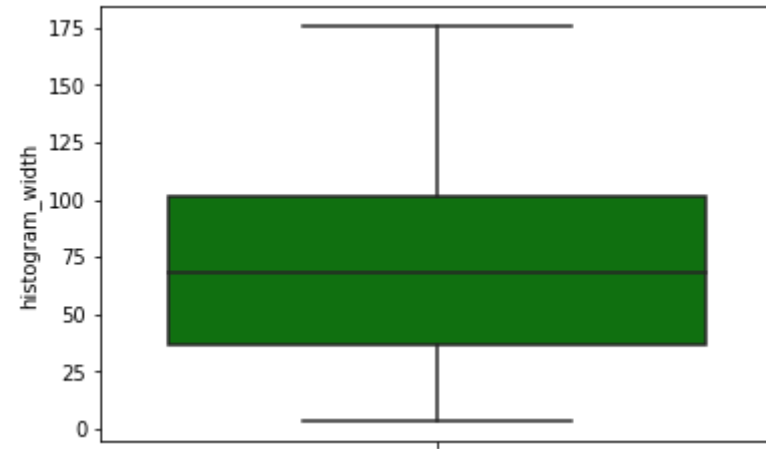


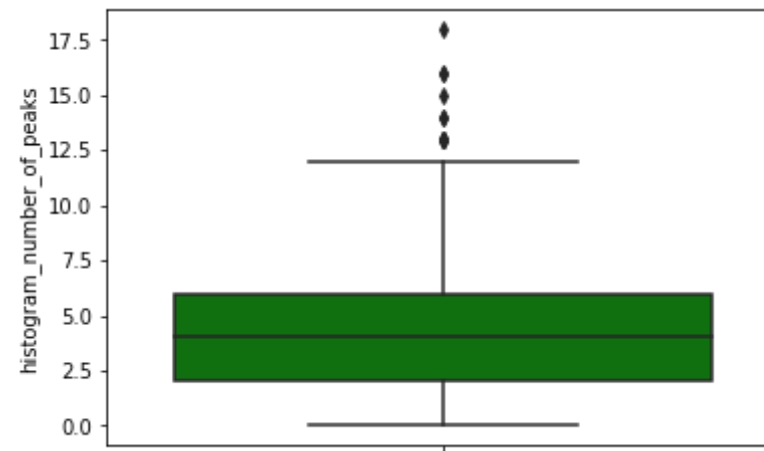
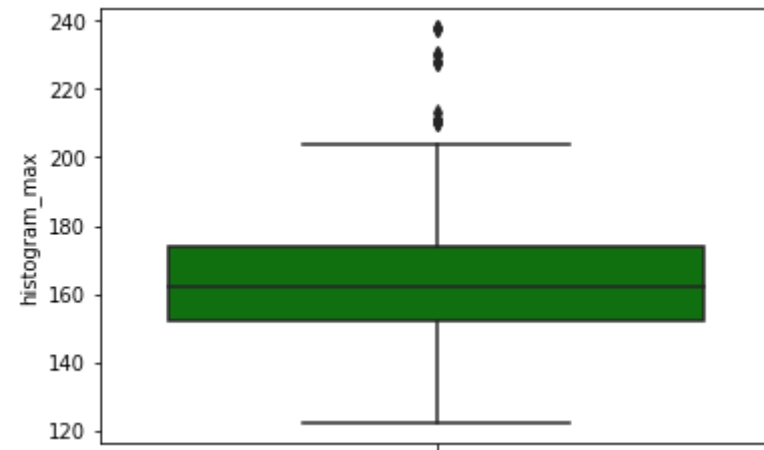
percentage\_of\_time\_with\_abnormal\_long\_term\_variability

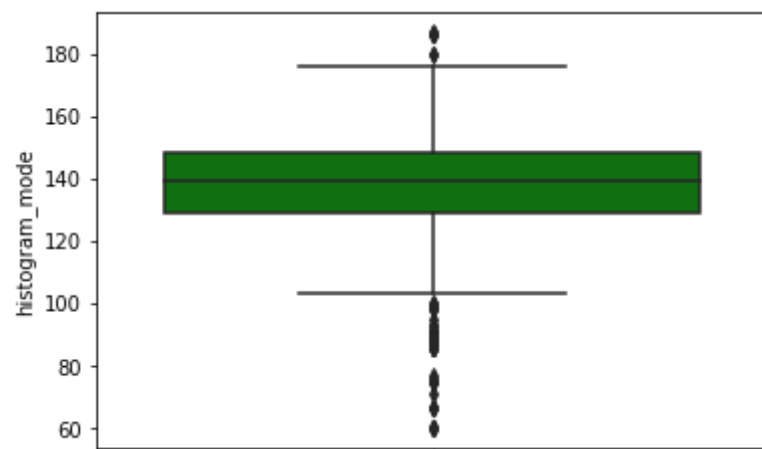
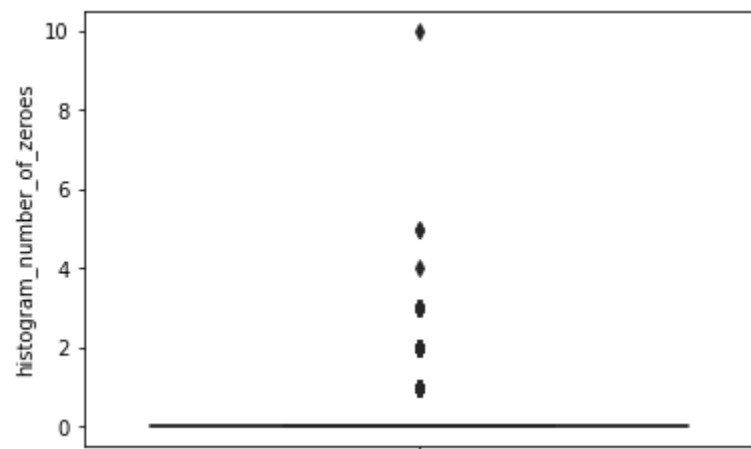


mean\_value\_of\_long\_term\_variability

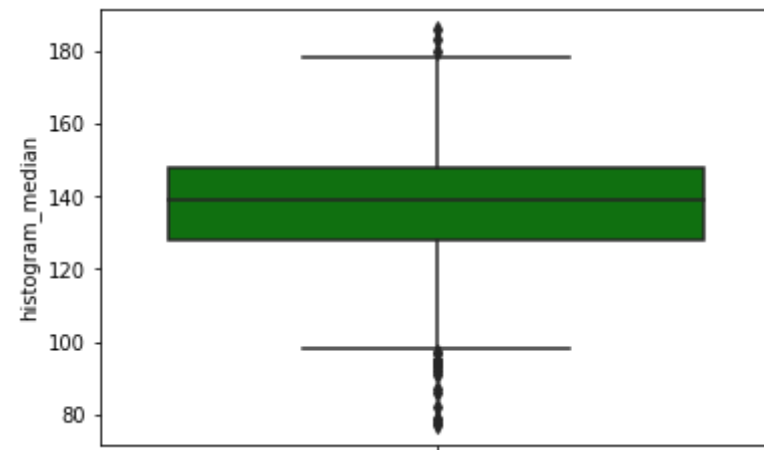
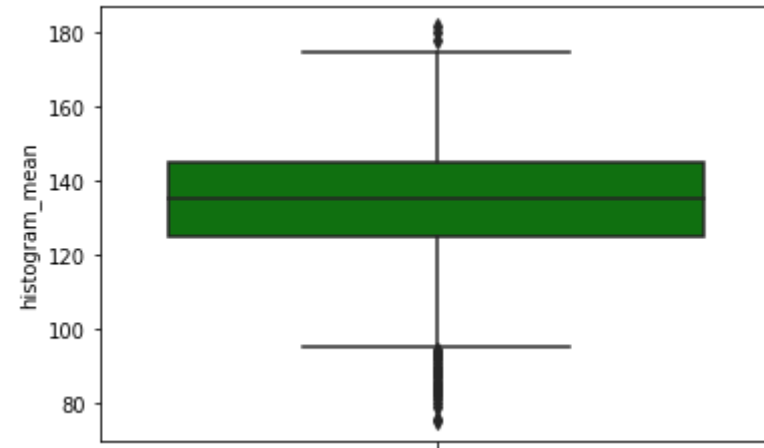


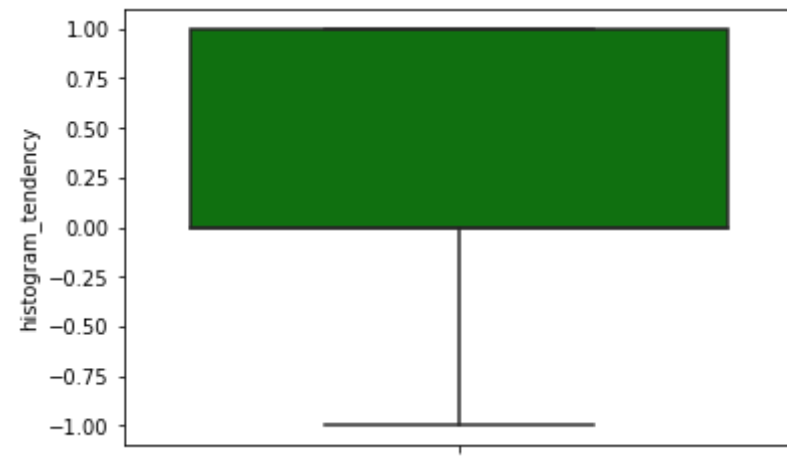
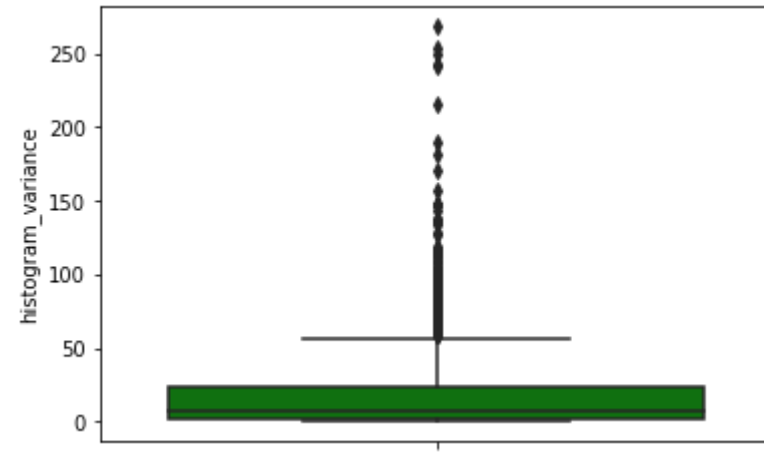


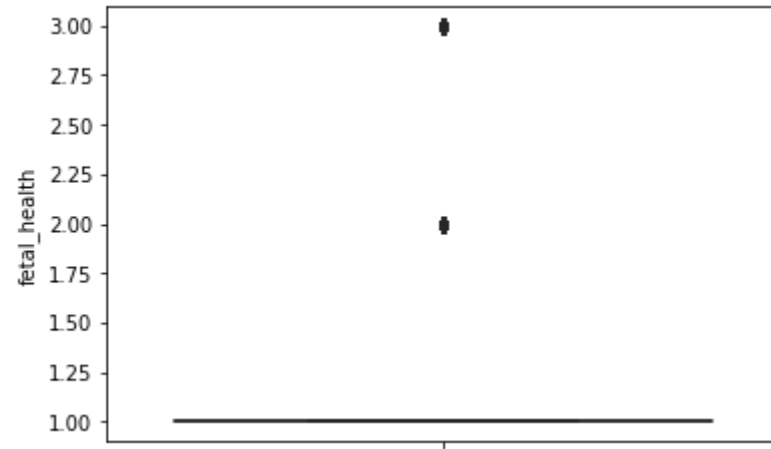




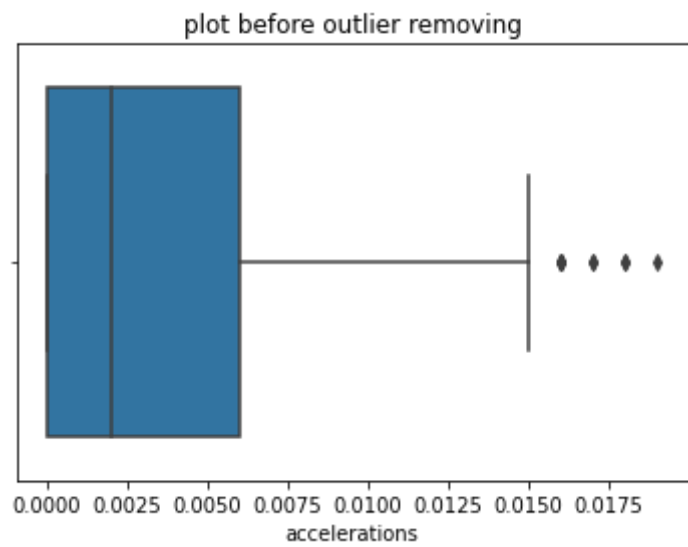
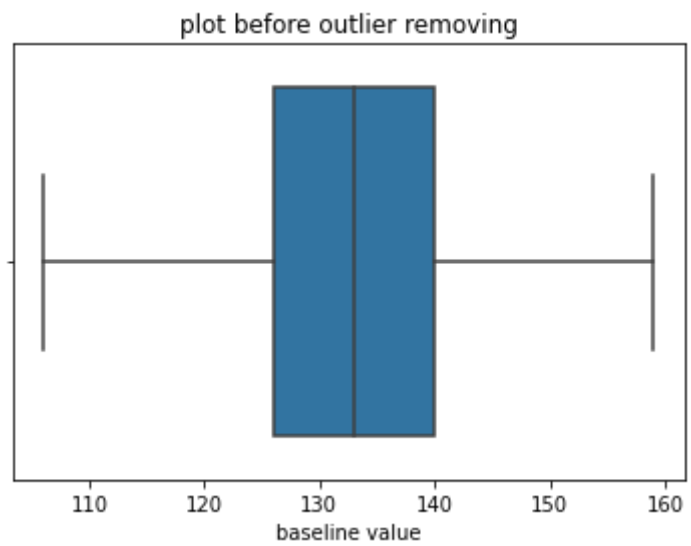


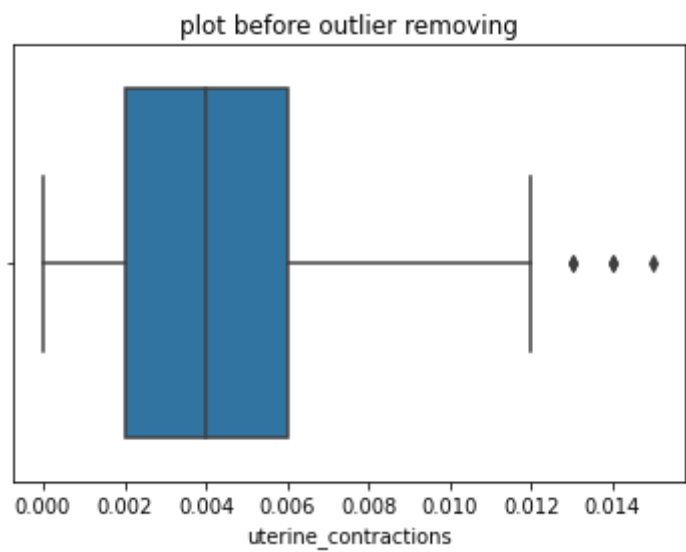
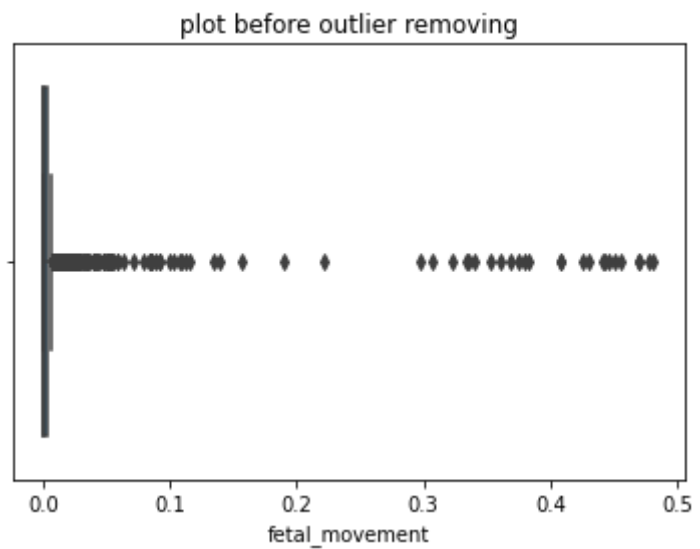


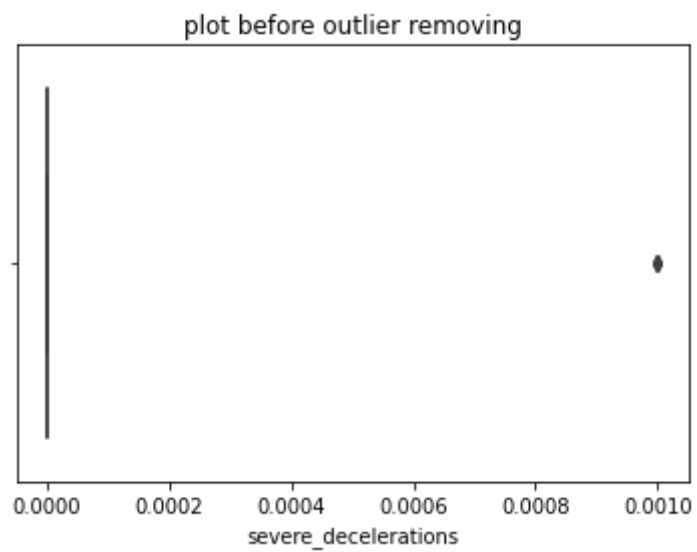
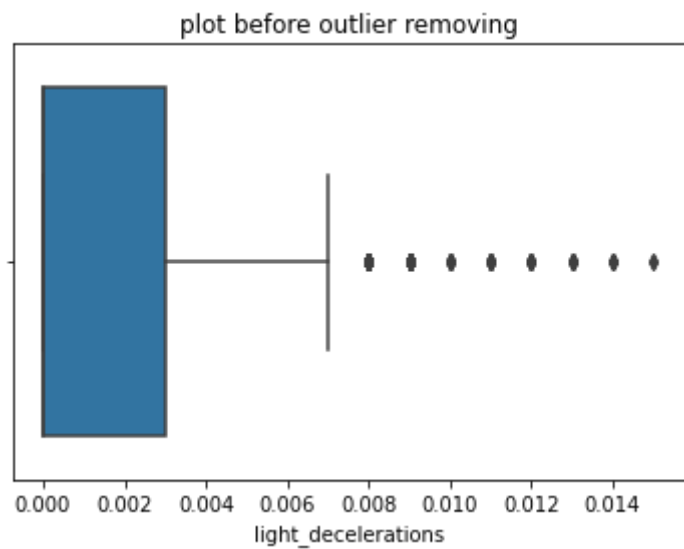


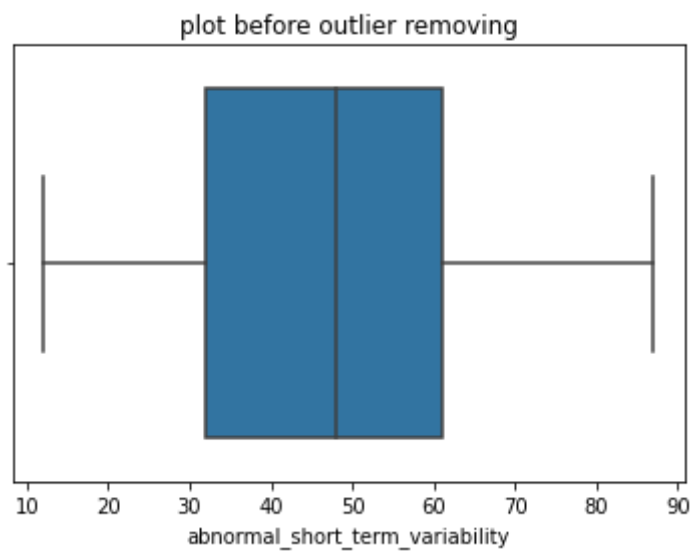
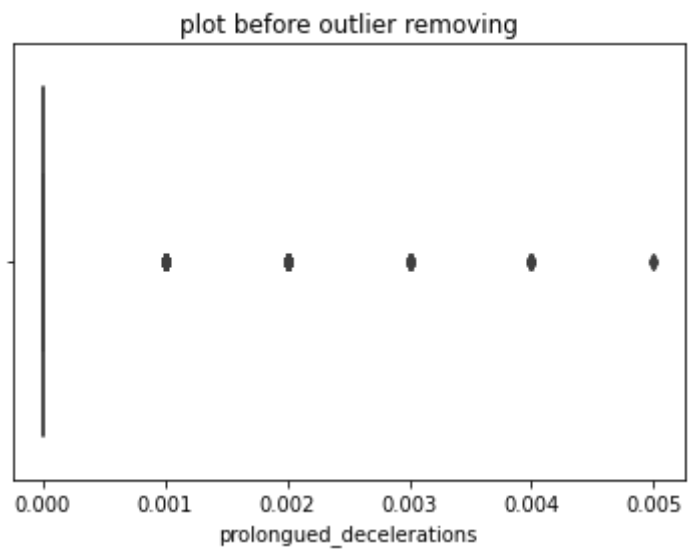


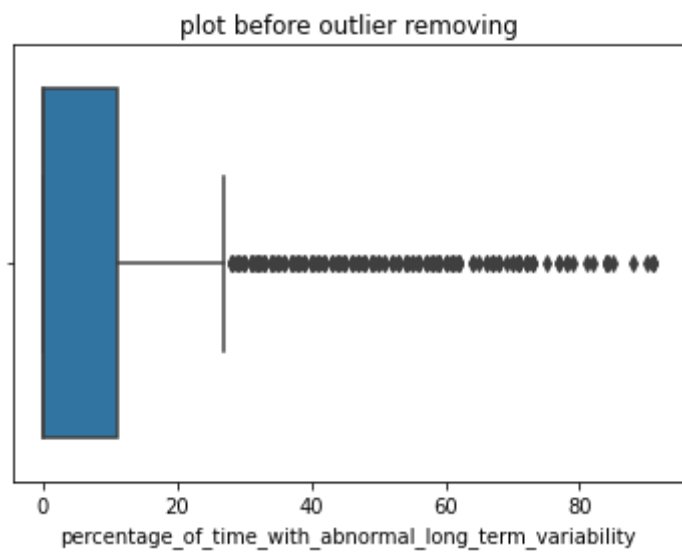
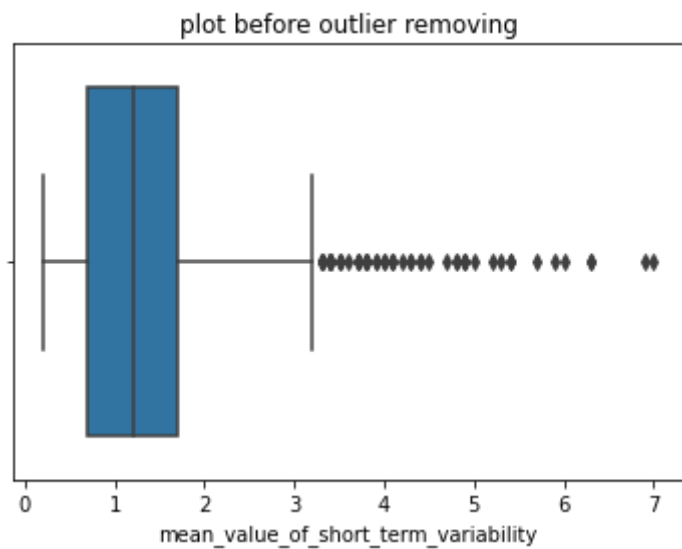
```
In [15]: #Removing the outliers
for i in range(len(nums)):
    sns.boxplot(df[nums[i]])
    plt.title(nums[i])
    plt.title("plot before outlier removing")
    plt.show()
    def drop_outliers(df, field_name):
        iqr = 1.5 * (np.percentile(df[field_name], 75) - np.percentile(df[field_name], 25))
        df.drop(df[df[field_name] > (iqr + np.percentile(df[field_name], 75))].index, inplace=True)
        df.drop(df[df[field_name] < (np.percentile(df[field_name], 25) - iqr)].index, inplace=True)
        iqr = 1.5 * (np.percentile(df[field_name], 75) - np.percentile(df[field_name], 25))
        df.drop(df[df[field_name] > (iqr + np.percentile(df[field_name], 75))].index, inplace=True)
        df.drop(df[df[field_name] < (np.percentile(df[field_name], 25) - iqr)].index, inplace=True)
        drop_outliers(df, nums[i])
    sns.boxplot(df[nums[i]])
    plt.title("plot after outlier removing")
    plt.show()
```





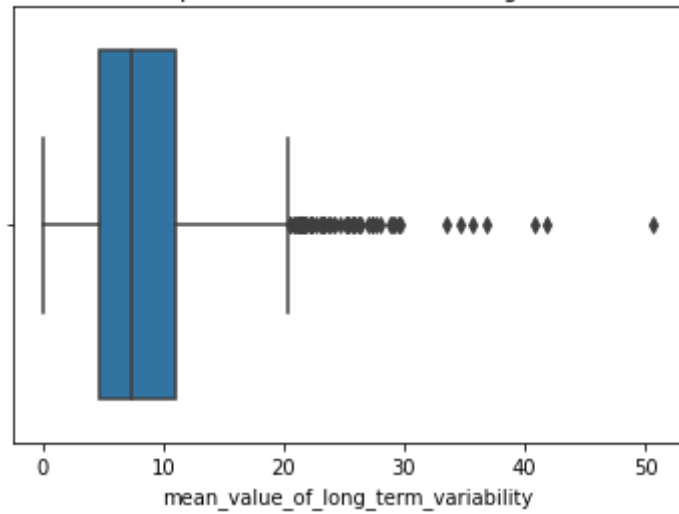




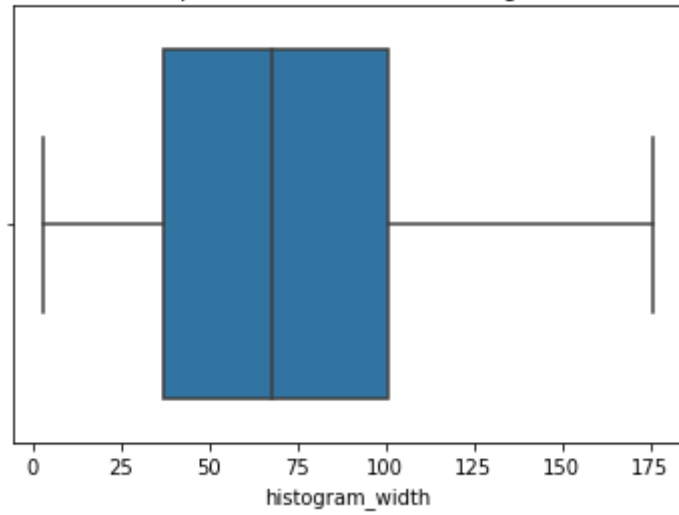


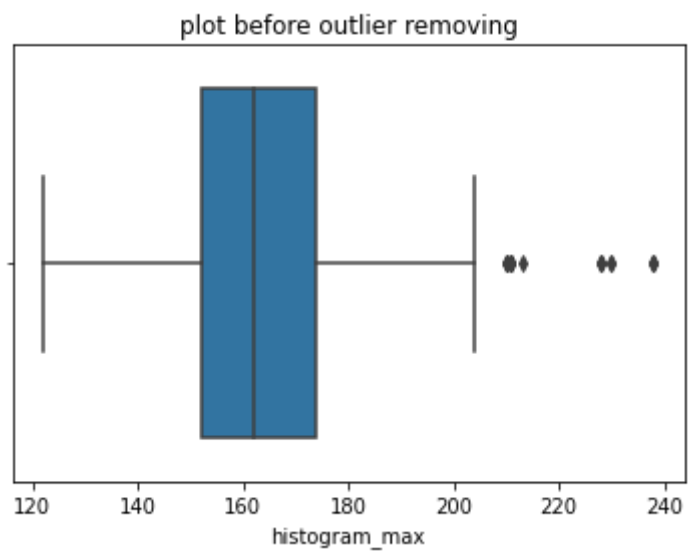
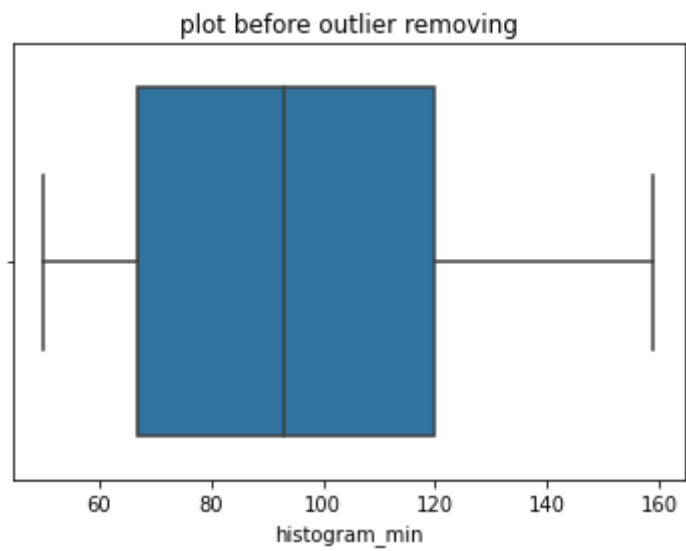


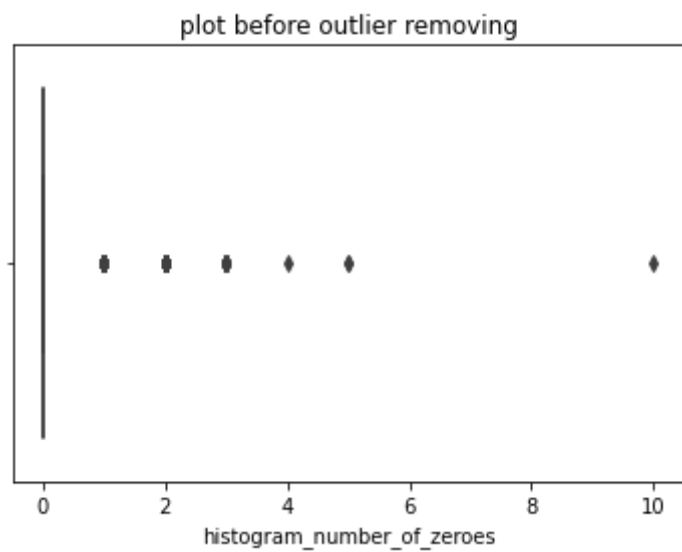
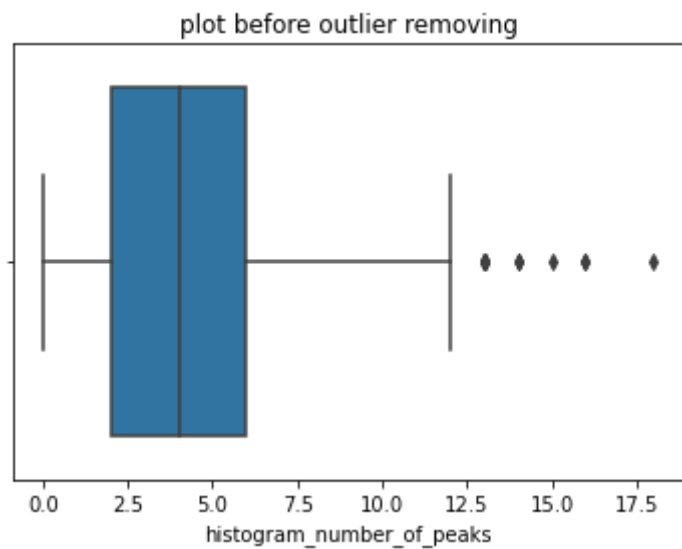
plot before outlier removing

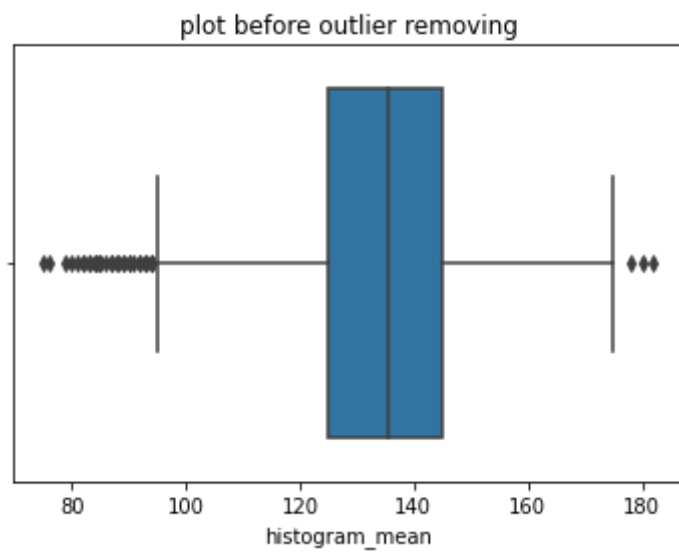
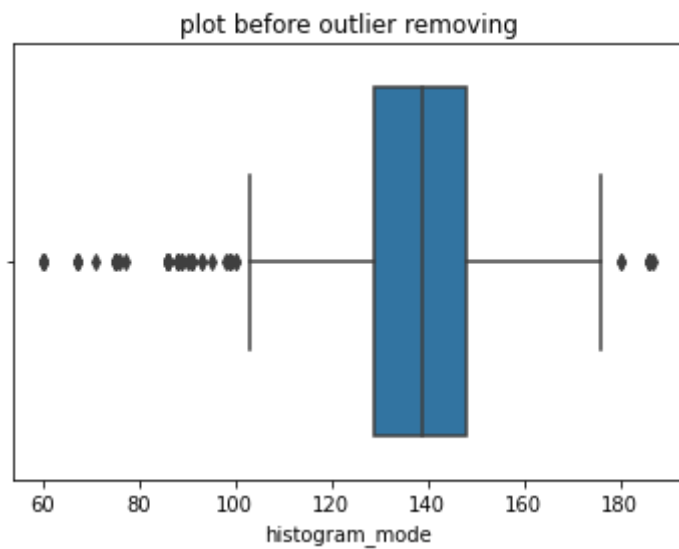


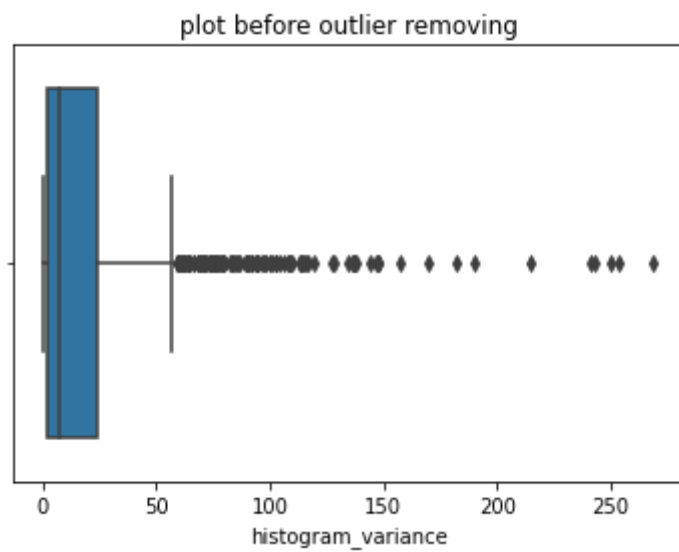
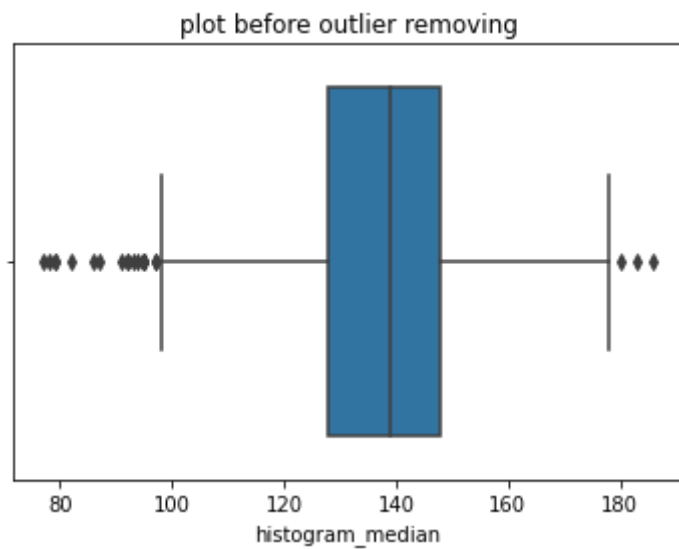
plot before outlier removing

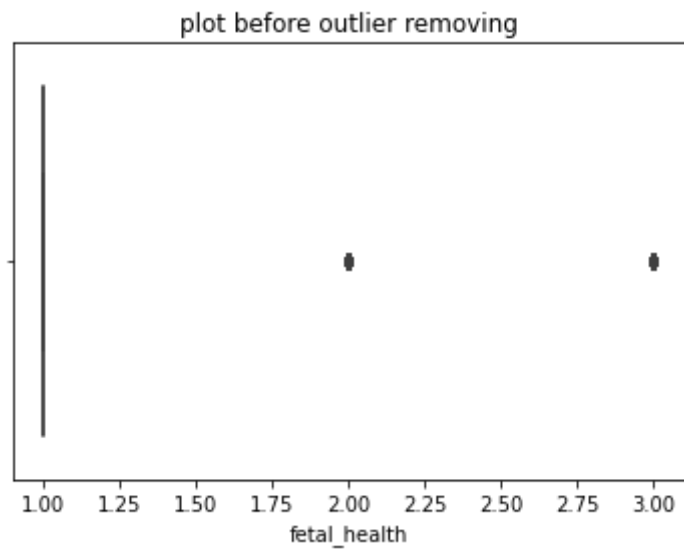
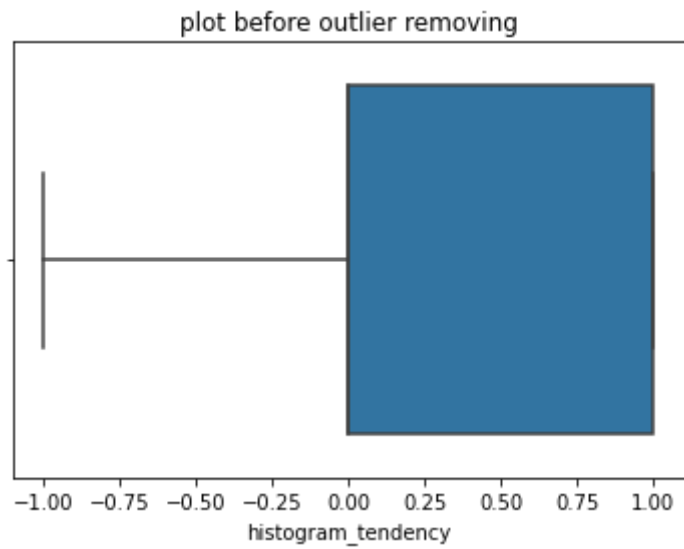












```
In [16]: #Feature scaling
from sklearn.preprocessing import StandardScaler
SC=StandardScaler()
X_train=SC.fit_transform(X_train)
X_test=SC.fit_transform(X_test)
print(X_train)
```

```

[[-1.46513509 -0.83485626 -0.20820521 ... -1.18281773 -0.60623421
  -2.14821249]
 [ 0.16393359 -0.83485626 -0.0746156 ... -0.08025974 -0.53911612
  1.13129086]
 [-1.2615015  2.2251405 -0.00782079 ... -0.42480911 -0.37132092
  -2.14821249]
 ...
 [ 1.28391831  1.71514104 -0.20820521 ...  1.160118  -0.50555708
  -0.50846081]
 [ 1.08028472 -0.83485626  0.6823922 ...  0.74665876 -0.06928955
  1.13129086]
 [ 0.97846793 -0.3248568  0.94957142 ...  0.74665876  1.00459976
  -0.50846081]]

```

In [17]: `print(X_test)`

```

[[ 0.92243937 -0.80741306  0.01293697 ...  0.39981121 -0.48257105
  1.06361291]
 [-1.28386413 -0.80741306 -0.20203067 ... -2.47496427  1.27487407
  -0.59523291]
 [-1.08329109 -0.80741306 -0.20203067 ... -0.83223542 -0.37919193
  1.06361291]
 ...
 [-1.18357761  0.24506147 -0.20203067 ... -0.76378839  0.34446195
  1.06361291]
 [-0.78243152  1.297536  -0.08627887 ... -0.010871  1.58501145
  1.06361291]
 [ 0.01986066 -0.80741306 -0.1854947 ... -0.010871  -0.65486959
  -0.59523291]]

```

In [18]: *#Building the Model*

```

from sklearn.neighbors import KNeighborsClassifier
classifier=KNeighborsClassifier(n_neighbors=5,metric='minkowski',p=2)
classifier.fit(X_train,y_train)
KNeighborsClassifier()
y_pred=classifier.predict(X_test)
print(np.concatenate((y_pred.reshape(len(y_pred),1),y_test.reshape(len
(y_test),1)),1))

```

```
[[1. 1.]
 [3. 3.]
 [1. 1.]
 ...
 [1. 1.]
 [1. 1.]
 [2. 1.]]
```

```
In [19]: #Cross validation
parameters ={'n_neighbors' : [5,7,9,11,13,15],
             'weights' : ['uniform','distance'],
             'metric' : ['minkowski','euclidean','manhattan']}
from sklearn.model_selection import RandomizedSearchCV
cv = RandomizedSearchCV(classifier,parameters ,cv=5)
cv.fit(X_train,y_train)
RandomizedSearchCV(cv=5, estimator=KNeighborsClassifier(),param_distributions={'metric': ['minkowski','euclidean','manhattan'],'n_neighbors': [5,7,9,11,13,15]}, scoring='accuracy')
y_pred = cv.predict(X_test)
```

```
In [21]: from sklearn.metrics import accuracy_score
print('\n Hyperparametric tuned knn accuracy:',accuracy_score(y_pred,y_test))
```

Hyperparametric tuned knn accuracy: 0.8960784313725491

```
In [22]: test_set=pd.read_csv('test.csv')
y_pre =classifier.predict(test_set)
print(y_pre )
```



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```

```
In [25]: #Saving the file in csv format
y = pd.DataFrame(y_pred).astype(int)
y.to_csv('Result.csv')
```

```
In [ ]:
```