

# Ways to deal with outliers

```
In [1]: #Importing the dataset
import pandas as p
import numpy as n
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import load_boston
import warnings
warnings.filterwarnings('ignore')
```

```
In [2]: boston=load_boston() #it is stored as dictionary
df= p.DataFrame(boston['data'],columns=boston['feature_names']) #converting dictionary to
```

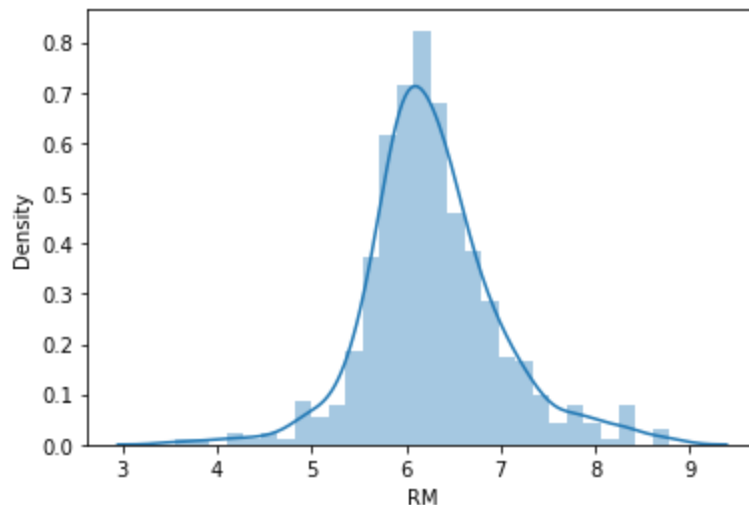
```
In [3]: df.head()
```

```
Out[3]:
```

	CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	B	LSTAT
0	0.00632	18.0	2.31	0.0	0.538	6.575	65.2	4.0900	1.0	296.0	15.3	396.90	4.98
1	0.02731	0.0	7.07	0.0	0.469	6.421	78.9	4.9671	2.0	242.0	17.8	396.90	9.14
2	0.02729	0.0	7.07	0.0	0.469	7.185	61.1	4.9671	2.0	242.0	17.8	392.83	4.03
3	0.03237	0.0	2.18	0.0	0.458	6.998	45.8	6.0622	3.0	222.0	18.7	394.63	2.94
4	0.06905	0.0	2.18	0.0	0.458	7.147	54.2	6.0622	3.0	222.0	18.7	396.90	5.33

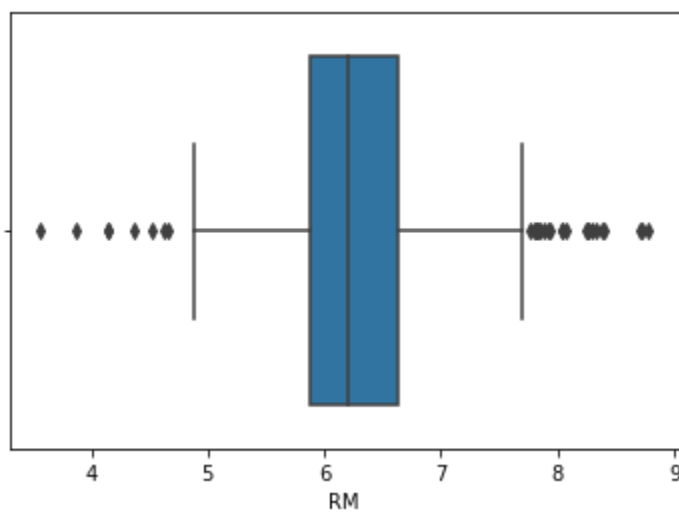
```
In [77]: sns.distplot(df['RM'])
```

```
Out[77]: <AxesSubplot:xlabel='RM', ylabel='Density'>
```



```
In [70]: #As we can see outliers
sns.boxplot(df['RM'])
```

```
Out[70]: <AxesSubplot:xlabel='RM'>
```



## Trimming outliers from the dataset

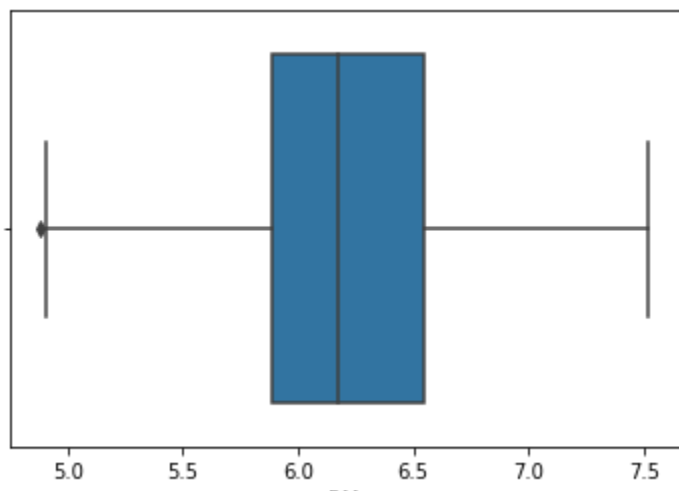
```
In [49]: def outliers(data):
  IQR=data.quantile(0.75)-data.quantile(0.25)
  lr=data.quantile(0.25)-(1.5*IQR) #lower range
  hr=data.quantile(0.70)+(1.5*IQR) #higher range
  return data.loc[~(n.where(data<lr,True,n.where(data>hr,True,False)))] #Without outlier
```

```
In [50]: outliers(df['RM']) #as we csn there is no outliers
```

```
Out[50]: 0      6.575
1      6.421
2      7.185
3      6.998
4      7.147
...
501    6.593
502    6.120
503    6.976
504    6.794
505    6.030
Name: RM, Length: 472, dtype: float64
```

```
In [51]: sns.boxplot(outliers(df['RM']))
```

```
Out[51]: <AxesSubplot:xlabel='RM'>
```



```
In [97]: #We can find outlier with using mean and standard deviation in case of IQR
def outliers(data,k):
    lr=data.mean()-(data.std()*k) #where n is number
    hr=data.mean()+(data.std()*k)
    return data.loc[~(n.where(data<lr,True,n.where(data>hr,True,False)))] #Without outlier
```

```
In [99]: outliers(df['RM'],1.5)
```

```
Out[99]: 0      6.575
1      6.421
2      7.185
3      6.998
4      7.147
...
501     6.593
502     6.120
503     6.976
504     6.794
505     6.030
Name: RM, Length: 449, dtype: float64
```

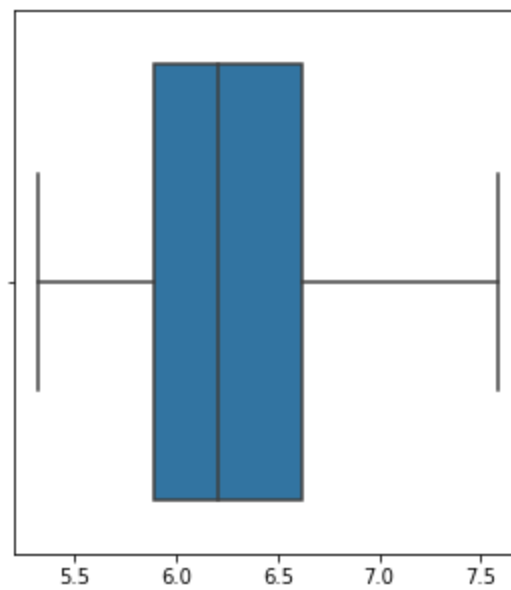
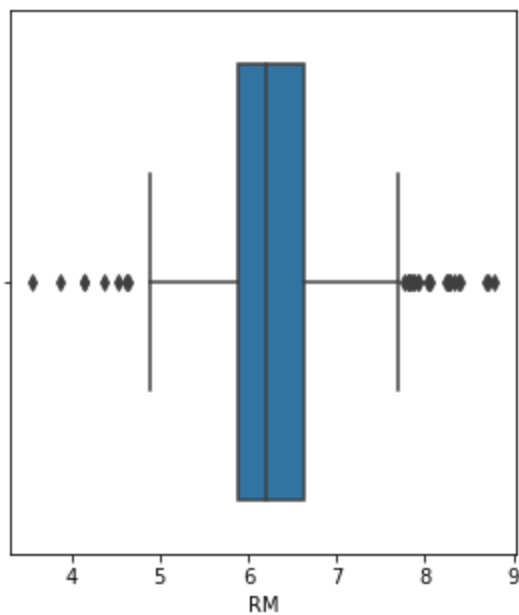
## Performing winsorization

Winsorizing is different from trimming because the extreme values are not removed, but are instead replaced by other values. Data greater than quantile 90 percent is replaced by value at 90 quantile similarly less than quantile 5 percent is replaced by value at 5 quantile

```
In [105... def fn(data,lw,h):
    lr=data.quantile(lw)
    hr=data.quantile(h)
    return n.where(data<lr,lr,n.where(data>hr,hr,data))
```

```
In [104... plt.figure(figsize=(10,5))
plt.subplot(1,2,1)
sns.boxplot(df['RM'])
plt.subplot(1,2,2)
sns.boxplot(fn(df['RM'],0.05,0.95))
```

```
Out[104... <AxesSubplot:>
```



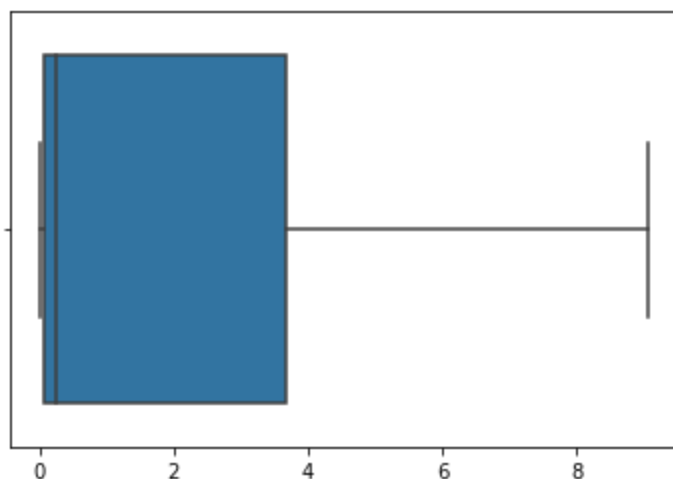
## Capping the variable at arbitrary maximum and minimum values

Similarly to winsorization, we can replace the extreme values by values closer to other values in the variable, by determining the maximum and minimum boundaries with the mean plus or minus the standard deviation, or the inter-quartile range proximity rule.

```
In [21]: def outliers(data):
IQR=data.quantile(0.75)-data.quantile(0.25)
lr=data.quantile(0.25)-(1.5*IQR) #lower range
hr=data.quantile(0.75)+(1.5*IQR) #higher range
return n.where(data<lr,lr,n.where(data>hr,hr,data))
def outliers_mean(data,k):
lr=data.mean()-(data.std()*k) #where k is number
hr=data.mean()+(data.std()*k)
return n.where(data<lr,lr,n.where(data>hr,hr,data))
```

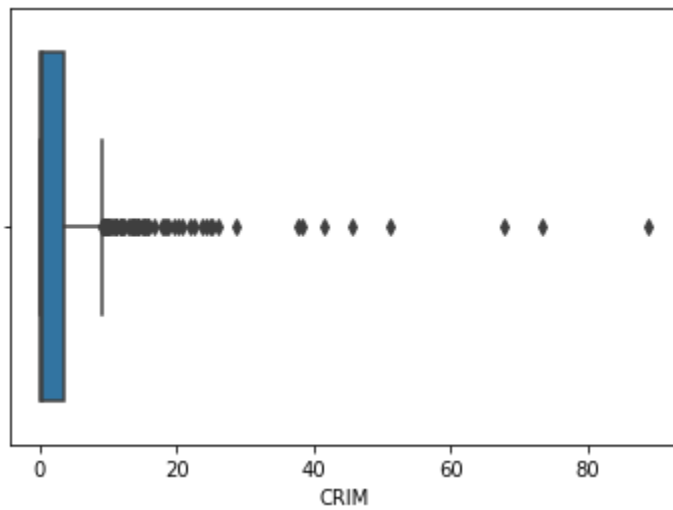
```
In [26]: sns.boxplot(outliers(df['CRIM']))
```

```
Out[26]: <AxesSubplot:>
```



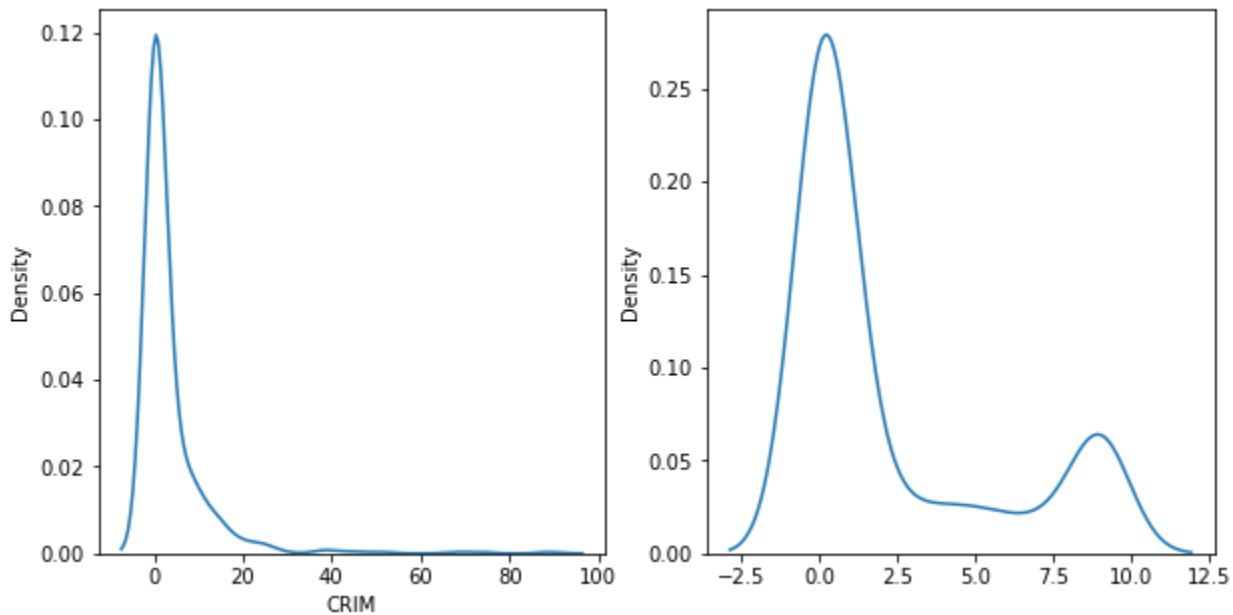
```
In [24]: sns.boxplot(df['CRIM'])
```

Out[24]: <AxesSubplot: xlabel='CRIM'>



```
In [42]: plt.figure(figsize=(10,5))
plt.subplot(1,2,1)
sns.kdeplot(df['CRIM'])
plt.subplot(1,2,2)
sns.kdeplot(outliers(df['CRIM']))
```

Out[42]: <AxesSubplot: ylabel='Density'>



here is one problem in this and winsorization when we have outliers in one tail so it will create the bump that is shown in this graph .

## Performing zero-coding – capping the variable at zero

Zero-coding is a variant of bottom-coding and refers to the process of capping, usually the lower value of the variable, at zero. It is commonly used for variables that cannot take negative values, such as age or income.

```
In [43]: #creating a dummy dataset
n.random.seed(29)
x=n.random.randn(200)
y=n.random.randn(200)*2
z=n.random.randn(200)*6+5
```

```
In [45]: df.head()
```

```
Out[45]:
```

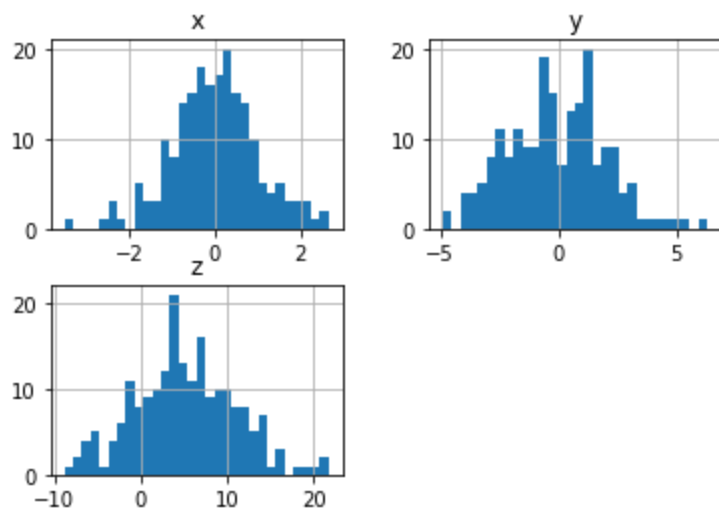
	x	y	z
0	-0.417482	2.903260	4.634786
1	0.706032	0.930279	10.236479
2	1.915985	0.688840	11.964026
3	-2.141755	-0.511348	13.884419
4	0.719057	-1.611499	18.030882

```
In [51]: df.min() #minmum values are negative
```

```
Out[51]: x    -3.505401  
         y    -4.901451  
         z    -8.863583  
         dtype: float64
```

```
In [55]: plt.figure(figsize=(15,5))  
df.hist(bins= 30)
```

```
Out[55]: array([[<AxesSubplot:title={'center':'x'}>,  
                <AxesSubplot:title={'center':'y'}>],  
          [<AxesSubplot:title={'center':'z'}>, <AxesSubplot:>]], dtype=object)  
<Figure size 1080x360 with 0 Axes>
```



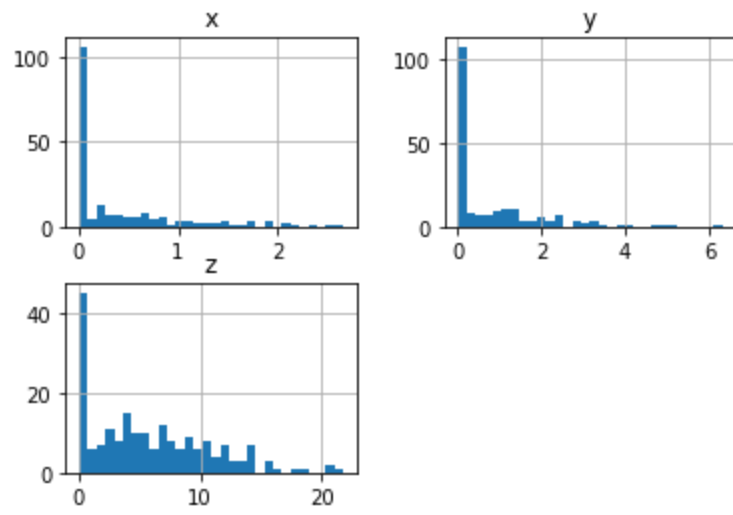
```
In [56]: #AS we can see the negative values
```

```
In [59]: #Replacing the negative values with zeros  
df['x'].loc[df['x']<0]=0  
df['y'].loc[df['y']<0]=0  
df['z'].loc[df['z']<0]=0
```

```
In [60]: plt.figure(figsize=(15,5))  
df.hist(bins= 30)
```

```
Out[60]: array([[<AxesSubplot:title={'center':'x'}>,  
                <AxesSubplot:title={'center':'y'}>],  
          [<AxesSubplot:title={'center':'z'}>, <AxesSubplot:>]], dtype=object)
```

<Figure size 1080x360 with 0 Axes>



```
In [61]: #now we can see increase in the nubor of zero values present
```

```
In [ ]:
```