

Data Science Advanced
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# Data Cleaning





## Data Cleaning Definition

- Data Cleaning (Data Cleansing) is the process of finding, correcting, and structuring problematic data points within a data set, so that it's generally uniform and prepared for analysis.
- This includes removing corrupt, irrelevant, and incomplete data and formatting it into a language that computers can understand for optimal analysis.





## Data Cleaning Definition

- Goal of Data Cleaning is to Identify and remove corrupt or irrelevant without deleting the necessary data to produce insights to ensure the best balance between data accuracy and minimal loss in sample size.
- It's important to remove these inconsistencies and format it into a language that computers can understand in order to increase the validity of the data set.



## Data Cleaning Purposes

- 1. Removing outliers, duplicate and unnecessary values from the dataset that can potentially skew the results when analysing the data.
- 2. Imputing and changing missing values and standardizing the data format to enhance the quality and consistency.
- 3. Identifying duplicate values and standardizing systems of measurements.
- 4. Detect and fix structural errors and typos, and validate the data to make it easier to handle.



## Data Cleaning Benefits

- 1. Errors Elimination.
- 2. Reduced costs associated with errors.
- 3. Enhances the integrity of data.
- 4. Ensures the highest quality of information for decision making.
- 5. Ensures ease of access to data.
- 6. Faster time to insights.



## Data Wrangling Definition

 Data Wrangling (Data Munging) Is the process of transforming and mapping data from one format into another.

• Purpose: is to prepare the data in a way that makes it accessible for effective use further down the line, to enhance the quality and assure useful data.



## Data Wrangling Purposes

- 1. Renaming columns and indices.
- 2. Detect and remove duplications.
- 3. Detect low variance feature.



- Pandas has a built-in function called rename() to change the column names. It's useful when you want to rename some selected columns.
  - Rename columns using a dictionary mapping.

```
# Rename columns
# PassengerId -> Id
# Pclass -> Classdf.rename(
columns=({ 'PassengerId': 'Id', 'Pclass': 'Class'}),
inplace=True,
df.head()
```



Instead of a dictionary mapping, we can also pass a function to the columns argument. For example, to convert column names into lowercase, we can pass a str.lower function.

df.rename(columns=str.lower).head()



• rename() function can be used to rename index as well. Let's first create a Data

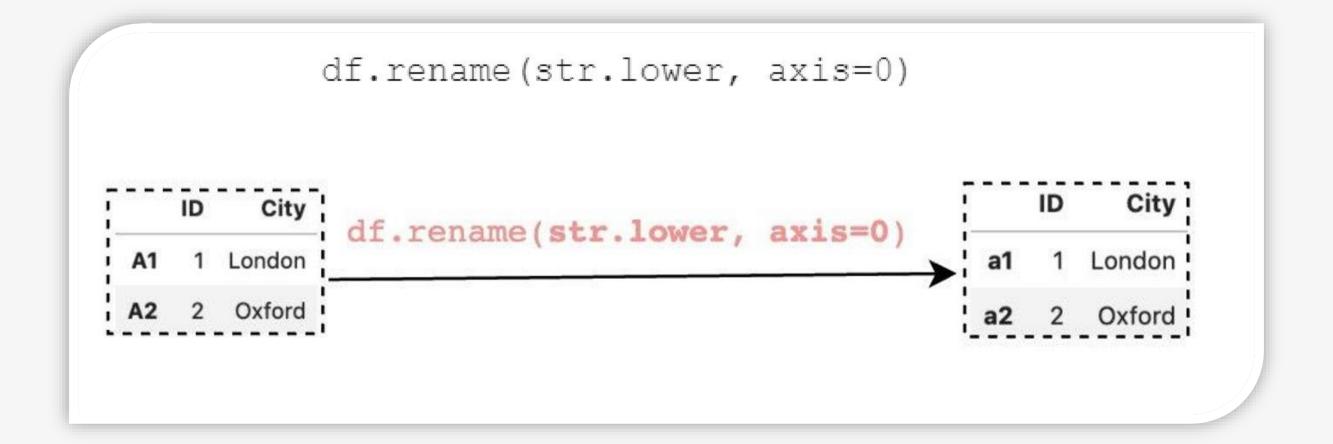
Frame:



• We can use dictionary mapping to rename index:



• We can also pass a function and set the axis argument to 0 or 'index'.





- Rows that have identical data are probably useless, if not dangerously misleading during model evaluation.
- A duplicate row in a row where each value in each column forth at row appears in identically the same order(same column values)in another row.
- Typically, this is not the case and machine learning algorithms will perform better by identifying and removing rows with duplicate data.
- From an algorithm evaluation perspective, duplicate rows will result in misleading performance.



The pandas function duplicated() will indicate whether a given row is duplicated or not.

- Rows are marked as False to indicate that it is not a duplicate.
- Rows are marked as True to indicate that it is a duplicate.

Rows of duplicated content in a dataset should probably be deleted from the dataset prior to modelling.



```
# locate rows of duplicate data
from pandas import read_csv
# define the location of the dataset
 'https://raw.githubusercontent.com/jbrownlee/Datasets/master/iris.csv'
 # load the dataset
df = read_csv(path, header=None)
# calculate duplicates
dups = df.duplicated()
# report if there are any duplicates
print(dups.any())
# list all duplicate rows
print(df[dups])
```



• There are many ways to achieve this, although Pandas provides the drop\_duplicates() function that achieves exactly this.

```
# delete rows of duplicate data from the dataset

from pandas import read_csv

define the location of the dataset

path =
    'https://raw.githubusercontent.com/jbrownlee/Datasets/master/iris.csv'

fload the dataset

df = read_csv(path, header=None)

print(df.shape)

delete duplicate rows

df.drop_duplicates(inplace=True)

print(df.shape)
```



- Running the example first loads the dataset and reports the number of rows and columns.
- Next, the rows of duplicated data are identified and removed from the Data Frame. Then the shape of the Data Frame is reported to confirm the change.

```
1 (150, 5)
2 (147, 5)
```



#### Detect Low Variance Feature

• Variance Threshold is one of the simplest baseline approaches to feature selection. It deletes all features whose variance doesn't meet some threshold. By default, it removes all zero-variance features, i.e. features that have the same value in all samples

$$Var[X]=p(1-p)$$



### Detect Low Variance Feature

```
>>> from sklearn.feature_selection import VarianceThreshold
>>> X = [[0, 0, 1], [0, 1, 0], [1, 0, 0], [0, 1, 1], [0, 1, 0], [0, 1, 1]]
>>> sel = VarianceThreshold(threshold=(.8 * (1 - .8)))
>>> sel.fit_transform(X)
array([[0, 1],
       [1, 0],
       [0, 0],
       [1, 1],
       [1, 0],
       [1, 1]])
```



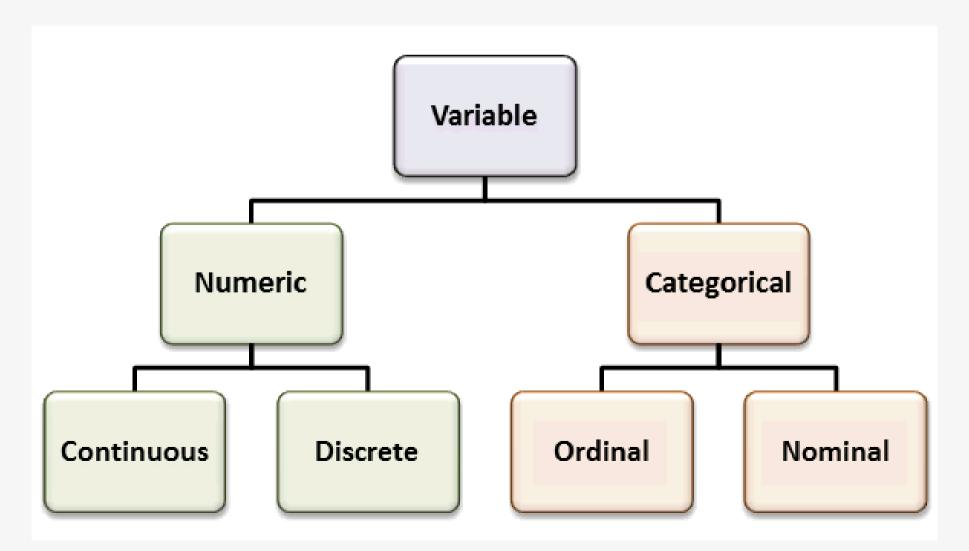
Data Types





## Data Types

 Data can come in many forms, but machine learning models rely on four primary data types. These include numerical data, categorical data, time series data, and text data.





### **Numeric Data**

- Continuous Variables are numeric variables that have an infinite number of values between any two values.
  - Example: The length of a part or the date and time a payment is received.
- Discreate Variables Are numeric variables that have a countable number of values between any two values. A discrete variable is always numeric.
  - Examples:

The number of customer complaints.

The number of flaws or defects.



## Categorical Data

- Categorical Variables is a type of data that consists of a finite number of categories, considered as gathered information that is divided into groups or categories with the aid of names or labels.
- Examples:
  - Gender (Male, Female).
  - Transportation choices (Car, Train, Subway, Bus, Plane).
  - Payment method (Cash, Credit Cards, Mobile Payments).
  - Marriage status (Single, Married).
  - Performance ratings (Poor, Fair, Average, Good, Excellent).



## Categorical Data

- 1. Ordinal Variables the categories have an inherent Order or type of categorical data consisting of a set of orders or scales.
- Example: a list of patients consists of sugar level in the human body which can be classified into high, low, and medium classes.
- 2. Nominal Variables the categories do not have an inherent order or the type of categorical data consists of names or labels without any numerical
- Example: the name of the different departments in any organization such as the human resource department, research, and development department, accounts and billing department, etc.



## Categorical vs Numeric Data

#### **Category**

- 1. Gender
- 2. Neighbourhood
- 3. Scholarship
- 4. Hypertension
- 5. Diabetes

#### Numeric

- 1. PatientID
- 2. AppointmentID
- 3. Age
- 4. Hypertension
- 5. Diabetes



# Missing Values





## Missing Values Definition

- Missing Data is defined as the absent values or the data that is not stored for some feature/s in the given dataset.
- Missing Values types:
- 1. Missing Completely At Random (MCAR).
- 2. Missing At Random (MAR).
- 3. Missing Not At Random (MNAR).



## Missing Values Types

Table 1

Types of Missing Values

Types of missing values	Description	Possible causes
Missing completely at random	Missing data occur completely at random without being influenced by other data.	Consent withdrawal, omission of major exams, death, discontinued follow-up and serious adverse reactions.
Missing at random	Missing data occur at a specific time point in conjunction with participant dissatisfaction with study outcomes and ongoing participation	Refusal to continue measurements.
Not missing at random	Missing data occur when a patient who is not satisfied with study outcomes performs the required measurements on his own, before the scheduled measurement.	If a patient finds the results of self-measurement dissatisfactory in addition to dissatisfaction related to the study, the patient may refuse further measurements.



## Standard Missing Values

- The missing values that Pandas can recognize and detect.
- In the third row, there's an empty cell.
- In the seventh row, there's an "NA" value.
- Pandas will detect both empty cells and "NA" types as missing values.

ST_NUM	ST_NAME	NUM_BEDROOMS	OWN_OCCUPIED
104	PUTNAM	3	Υ
197	LEXINGTON	3	N
	LEXINGTON	n/a	N
201	BERKELEY	1	12
203	BERKELEY	3	Υ
207	BERKELEY	NA	Υ
NA	WASHINGTON	2	
213	TREMONT		Υ
215	TREMONT	na	Υ



## Non-Standard Missing Values

- The missing values have different formats.
- In the NUM\_BEDROOMS column, there are four missing values.
- n/a, NA, --, na It's important to recognize these non-standard types of missing values for purposes of summarizing and transforming missing values.

ST_NUM	ST_NAME	NUM_BEDROOMS	OWN OCCUPIED
104	PUTNAM	3	Υ
197	LEXINGTON	3	N
	LEXINGTON	n/a	N
201	BERKELEY	1	12
203	BERKELEY	3	Υ
207	BERKELEY	NA	Υ
NA	WASHINGTON	2	
213	TREMONT		Υ
215	TREMONT	na	Υ



## Unexpected Missing Values

- For example, if our feature is expected to be a string, but there's a numeric type, then technically this is also a missing value.
- In the fourth row, there's the number 12.

• The response for Owner Occupied should clearly be a string (Y or N), so this numeric

type should be a missing value.

ST_NUM	ST_NAME	NUM_BEDROOMS	OWN_OCCUPIED
104	PUTNAM	3	Υ
197	LEXINGTON	3	N
	LEXINGTON	n/a	N
201	BERKELEY	1	12
203	BERKELEY	3	Υ
207	BERKELEY	NA	Υ
NA	WASHINGTON	2	
213	TREMONT		Υ
215	TREMONT	na	Υ



## Why Handling Missing Values?

- It is important to handle the missing values in the right way.
- Missing values in the dataset cause many machine learning algorithms to fail.
- Lead to building a biased machine learning model which will result in faulty results if the missing values are not handled correctly.
- Missing data can cause a reduction of precision in the statistical analysis.



## Handling Missing Values Methods

#### 1. Complete Case Analysis:

- This method aims to use the data of variables observed after removing all missing values.
- Although it has the advantage of the simplicity of analysis, decreasing the sample size and reducing statistical power are disadvantages because drawing statistical inferences becomes difficult during analysis.



## Handling Missing Values Methods

#### 2. Available Case Analysis:

This method deals with only the data available for each analysis. It allows a larger sample size than that used for complete case analysis. However, this approach causes sample sizes to vary between the variables used in the analysis.



### Handling Missing Values Methods

#### 3. Imputation Analysis:

- This method involves the replacement of values with substituted values computed from statistical analysis to Preserv data set completed without missing values for analysis. Imputations can be created by using either an explicit or an implicit modelling approach.
- It includes different methods of imputation by mean, median, probability, ratio, regression, predictive regression, and assumption of distribution.



```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890
Data columns (total 7 columns):
 # Column Non-Null Count Dtype
0 Survived 891 non-null int64
1 Pclass 891 non-null int64
2 Sex 891 non-null int64
3 Age 714 non-null float64
4 SibSp 891 non-null int64
5 Parch 891 non-null int64
 6 Fare 891 non-null float64
dtypes: float64(2), int64(5)
memory usage: 48.9 KB
```



#### 1. Remove the Column:

- In this case let's remove the Age column from our dataset and check for accuracy.
- This approach can be used when there are many null values in the column.

```
updated_df = df.dropna(axis=1)
updated_df.info()
```



• After removing Age column.

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890
Data columns (total 5 columns):
 # Column Non-Null Count Dtype
  Pclass 891 non-null int64
 1 Sex 891 non-null int64
 2 SibSp 891 non-null int64
 3 Parch 891 non-null int64
   Fare 891 non-null float64
dtypes: float64(1), int64(4)
memory usage: 34.9 KB
```



Accuracy after removing Age column.

```
from sklearn import metrics
from sklearn.model_selection import train_test_split
X_train, X_test,y_train,y_test =
  train_test_split(updated_df,y,test_size=0.3)
  from sklearn.linear_model import LogisticRegression
  lr = LogisticRegression()
  lr.fit(X_train,y_train)
  pred = lr.predict(X_test)
  print(metrics.accuracy_score(pred,y_test))

0.7947761194029851
```



#### 2. Remove the Null rows:

- If there is a certain row with many missing values, then you can remove the entire row with all the features in that row.
- axis=0 is used to drop the row with `NaN` values.

```
updated_df = newdf.dropna(axis=0)

y1 = updated_df['Survived']
updated_df.drop("Survived",axis=1,inplace=True)
updated_df.info()
```



After removing Null rows.

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 714 entries, 0 to 890
Data columns (total 6 columns):
    Column Non-Null Count Dtype
   Pclass 714 non-null int64
   Sex 714 non-null int64
  Age 714 non-null float64
  SibSp 714 non-null int64
  Parch 714 non-null int64
   Fare 714 non-null float64
dtypes: float64(2), int64(4)
memory usage: 39.0 KB
```



Accuracy after removing Null rows.

```
from sklearn import metrics
from sklearn.model_selection import train_test_split
X_train, X_test,y_train,y_test =
  train_test_split(updated_df,y1,test_size=0.3)
from sklearn.linear_model import LogisticRegression
lr = LogisticRegression()
lr.fit(X_train,y_train)
pred = lr.predict(X_test)
print(metrics.accuracy_score(pred,y_test))

0.8232558139534883
```



- 3. Fill missing values:
- The possible ways to fill the Missing Values are:
- 1. Filling the missing data with the mean or median value in case of a numerical variable.
- 2. Filling the missing data with the mode in case of a categorical value.



• fillna() function can be used to fill the null values in the dataset.

```
updated_df = df
updated_df['Age']=updated_df['Age'].fillna(updated_df['Age'].mean())
updated_df.info()
```



• After filling Null values in the Age column with Mean.

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890
Data columns (total 7 columns):
    Column Non-Null Count Dtype
0 Survived 891 non-null int64
1 Pclass 891 non-null int64
2 Sex 891 non-null int64
3 Age 891 non-null float64
4 SibSp 891 non-null int64
5 Parch 891 non-null int64
6 Fare 891 non-null float64
dtypes: float64(2), int64(5)
memory usage: 48.9 KB
```



• Accuracy after filling with Mean.

```
y1 = updated_df['Survived']
updated_df.drop("Survived",axis=1,inplace=True)
from sklearn import metrics
from sklearn.model_selection import train_test_split
X_train, X_test,y_train,y_test =
train_test_split(updated_df,y1,test_size=0.3)
from sklearn.linear_model import LogisticRegression
lr = LogisticRegression()
lr.fit(X_train,y_train)
pred = lr.predict(X_test)
print(metrics.accuracy_score(pred,y_test))
```



# Outliers





#### What is the Outlier?

- Outlier in statistics, an outlier is an observation point that is distant from other observations.
- Let's have a look at some examples. Suppose you have been asked to observe the performance of the Indian cricket team i.e. the Run made by each player and collect the

data.

Players	Scores
Player1	500
Player2	350
Player3	10
Player4	300
Player5	450



#### What is the Outlier?

- Now that we know outliers can either be a mistake or just variance, how would you decide if they are important or not.
- Well, it is pretty simple if they are the result of a mistake, then we can ignore them, but if it is just variance in the data we would need to think a bit further. Before we try to understand whether to ignore the outliers or not, we need to know the ways to identify them.



## Detecting Outliers Ways

- Graphical Ways:
  - 1. Scatter Plots
  - 2. Box Plots
- Numerical Ways:
  - 1. Z-Score
  - 2. IQR



### Example of Outlier

So, Let's get start. We will be using Boston House Pricing Dataset which is included
in the sklearn dataset API. We will load the dataset and separate out the features and
targets.

```
boston = load_boston()
x = boston.data
y = boston.target
columns = boston.feature_names

#create the dataframe
boston_df = pd.DataFrame(boston.data)
boston_df.columns = columns
boston_df.head()
```



## Example of Outlier

• Features/independent variable will be used to look for any outlier. Looking at the data below, it s seems, we only have numeric values.

CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	В	LSTAT
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
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
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
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
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
					Bost	on Hou	ısing Dat	a				



## Example of Outlier

• There are two types of analysis we will follow to find the outliers: Uni-variate(one variable outlier analysis) and Multi-variate(two or more variable outlier analysis). When you will start coding and plotting the data, you will see yourself that how easy it was to detect the outlier. To keep things simple, we will start with the basic method of detecting outliers and slowly move on to the advance methods.



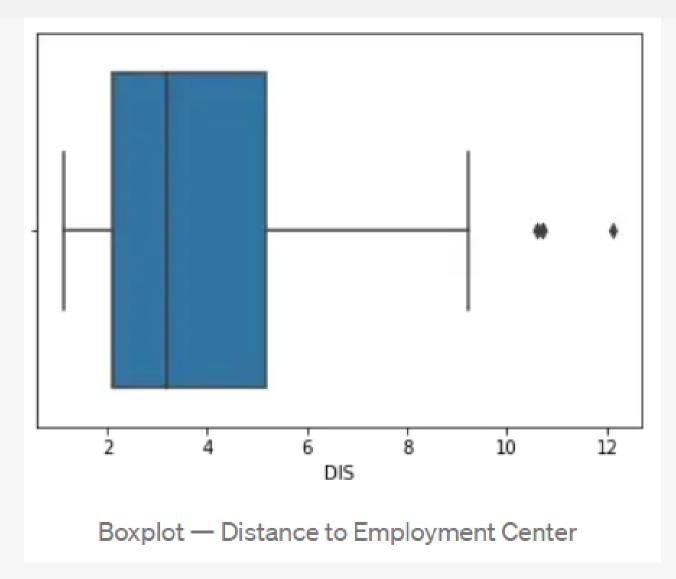
#### Discover outliers with Box Plot

- Box Plot is a method for graphically depicting groups of numerical data through their quartiles. Box plots may also have lines extending vertically from the boxes (whiskers) indicating variability outside the upper and lower quartiles, hence the terms box-and-whisker plot and box-and-whisker diagram. Outliers may be plotted as individual points.
- Above definition suggests, that if there is an outlier it will plotted as point in boxplot but other population will be grouped together and display as boxes. Let's try and see it ourselves.



#### Discover outliers with Box Plot

import seaborn as sns
sns.boxplot(x=boston\_df['DIS'])





#### Discover outliers with Box Plot

- Previous plot shows three points between 10 to 12, these are outliers as there are not included in the box of other observation i.e. no where near the quartiles.
- Here we analysed Uni-variate outlier i.e. we used DIS column only to check the outlier. But we can do multivariate outlier analysis too. Can we do the multivariate analysis with Box plot? Well it depends, if you have a categorical values then you can use that with any continuous variable and do multivariate outlier analysis. As we do not have categorical value in our Boston Housing dataset, we might need to forget about using box plot for multivariate outlier analysis.



#### Discover outliers with Scatter Plot

- Scatter Plot is a type of plot or mathematical diagram using Cartesian coordinates to display values for typically two variables for a set of data. The data are displayed as a collection of points, each having the value of one variable determining the position on the horizontal axis and the value of the other variable determining the position on the vertical axis.
- As the definition suggests, the scatter plot is the collection of points that shows values for two variables. We can try and draw scatter plot for two variables from our housing dataset.



#### Discover outliers with Scatter Plot

• Looking at the plot below, we can most of data points are lying bottom left side but there are points which are far from the population like top right corner.

```
fig, ax = plt.subplots(figsize=(16,8))
ax.scatter(boston_df['INDUS'], boston_df['TAX'])
ax.set_xlabel('Proportion of non-retail business acres per town')
ax.set_ylabel('Full-value property-tax rate per $10,000')
plt.show()

700

600

900

Froportion of non-retail business acres per town

Scatter plot — Proportion of non-retail business acres per town v/s Full value property tax
```



• **Z-Score** is the signed number of standard deviations by which the value of an observation or data point is above the mean value of what is being observed or measured.

• The intuition behind Z-score is to describe any data point by finding their relationship with the Standard Deviation and Mean of the group of data points. Z-score is finding the distribution of data where mean is 0 and standard deviation is 1 i.e. normal distribution.



• You must be wondering that, how does this help in identifying the outliers? Well, while calculating the Z-score we re-scale and centre the data and look for data points which are too far from zero. These data points which are way too far from zero will be treated as the outliers. In most of the cases a threshold of 3 or -3 is used i.e. if the Z-score value is greater than or less than 3 or -3 respectively, that data point will be identified as outliers.



• We will use **Z-score function** defined in **SciPy library** to detect the outliers.

```
from scipy import stats
 import numpy as np
 z = np.abs(stats.zscore(boston_df))
 print(z)
[[0.41771335 0.28482986 1.2879095 ... 1.45900038 0.44105193 1.0755623 ]
[0.41526932 0.48772236 0.59338101 ... 0.30309415 0.44105193 0.49243937]
[0.41527165 0.48772236 0.59338101 ... 0.30309415 0.39642699 1.2087274 ]
 [0.41137448 0.48772236 0.11573841 ... 1.17646583 0.44105193 0.98304761]
[0.40568883 0.48772236 0.11573841 ... 1.17646583 0.4032249 0.86530163]
[0.41292893 0.48772236 0.11573841 ... 1.17646583 0.44105193 0.66905833]]
                          Z-score of Boston Housing Data
```



• Looking the code and the output above, it is difficult to say which data point is an outlier. Let's try and define a threshold to identify an outlier.

```
threshold = 3
  print(np.where(z > 3))
This will give a result as below -
 (array([ 55, 56, 57, 102, 141, 142, 152, 154, 155, 160, 162, 163, 199,
       200, 201, 202, 203, 204, 208, 209, 210, 211, 212, 216, 218, 219,
       220, 221, 222, 225, 234, 236, 256, 257, 262, 269, 273, 274, 276,
       277, 282, 283, 283, 284, 347, 351, 352, 353, 353, 354, 355, 356,
       357, 358, 363, 364, 364, 365, 367, 369, 370, 372, 373, 374, 374,
       380, 398, 404, 405, 406, 410, 410, 411, 412, 412, 414, 414, 415,
       416, 418, 418, 419, 423, 424, 425, 426, 427, 427, 429, 431, 436,
       437, 438, 445, 450, 454, 455, 456, 457, 466], dtype=int64), array([ 1, 1, 1, 11, 12, 3, 3,
 3, 3, 3, 3, 3, 1, 1, 1, 1, 1,
       1, 3, 3, 3, 3, 3, 3, 3, 3, 3, 3, 5, 3, 3, 5,
        5, 3, 3, 3, 3, 3, 1, 3, 1, 1, 7, 7, 1, 7, 7, 7,
       3, 3, 3, 3, 5, 5, 5, 3, 3, 12, 5, 12, 0, 0, 0,
       0, 5, 0, 11, 11, 11, 12, 0, 12, 11, 11, 0, 11, 11, 11, 11, 11,
       dtype=int64))
                          Data points where Z-scores is greater than 3
```



 Don't be confused by the results. The first array contains the list of row numbers and second array respective column numbers, which mean z[55][1] have a Z-score higher than 3.

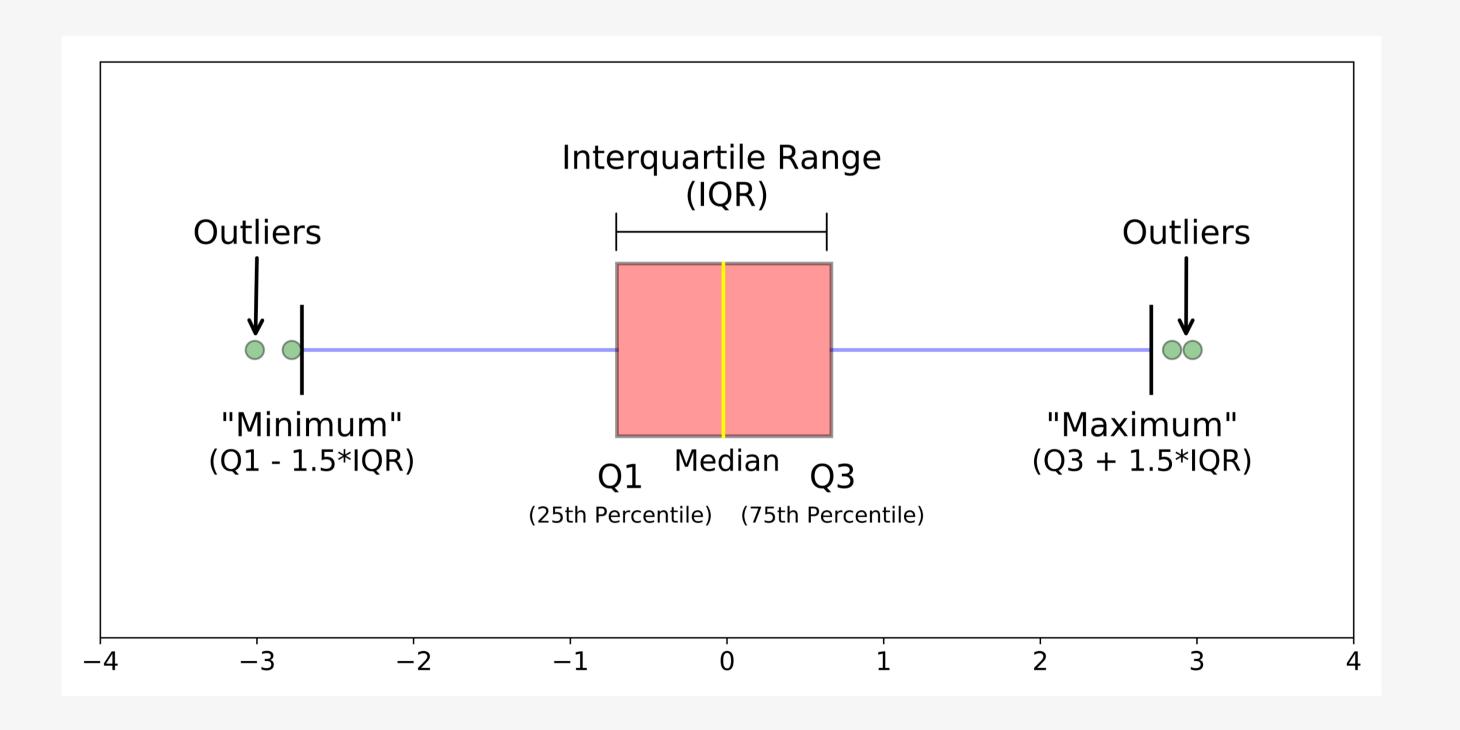
```
print(z[55][1])
3.375038763517309
```

So, the data point — 55th record on column ZN is an outlier.



- The interquartile range (IQR) is a measure of statistical dispersion, being equal to the difference between 75th and 25th percentiles, or between upper and lower quartiles, IQR = Q3 Q1.
- In other words, the IQR is the first quartile subtracted from the third quartile; these quartiles can be clearly seen on a box plot on the data.
- IQR is somewhat similar to Z-score in terms of finding the distribution of data and then keeping some threshold to identify the outlier.







• Let's find out we can box plot uses IQR and how we can use it to find the list of outliers as we did using Z-score calculation. First we will calculate IQR.

```
Q1 = boston_df_o1.quantile(0.25)
  Q3 = boston_df_o1.quantile(0.75)
  IQR = Q3 - Q1
  print(IQR)
Here we will get IQR for each column.
                          CRIM
                                        3.565378
                          ΖN
                                       12.500000
                          INDUS
                                       12.910000
                          CHAS
                                        0.000000
                          NOX
                                        0.175000
                           RM
                                        0.738000
                           AGE
                                       49.050000
                          DIS
                                        3.088250
                           RAD
                                       20.000000
                          TAX
                                      387.000000
                          PTRATIO
                                        2.800000
                                       20.847500
                          LSTAT
                                       10.005000
                          dtype: float64
                                IQR for each column
```



• The data point where we have False that means these values are valid whereas True indicates presence of an outlier.

QR)	,												
4	False	Fals											
5	False	Fals											
6	False	Fals											
7	False	Fals											
8	False	Fals											
9	False	Fals											
10	False	Fals											
11	False	Fals											
12	False	Fals											
13	False	Fals											
14	False	Fals											
15	False	Fals											
16	False	Fals											
17	False	Fals											
18	False	True	Fals										
19	False	Fals											
20	False	Fals											
21	False	Fals											



### Removing outliers with Z-Score

• We saw how one can detect the outlier using Z-score but now we want to remove or filter the outliers and get the clean data. This can be done with just one line code as we have already calculated the Z-score.

```
boston_df_o = boston_df_o[(z < 3).all(axis=1)]

boston_df.shape

(506, 13)

boston_df_o.shape

(415, 13)

With and without outlier size of the dataset</pre>
```



#### Removing outliers with IQR-Score

• Just like Z-score we can use previously calculated IQR score to filter out the outliers by keeping only valid values.

```
boston_df_out = boston_df_o1[~((boston_df_o1 < (Q1 - 1.5 * IQR)) |
  (boston_df_o1 > (Q3 + 1.5 * IQR))).any(axis=1)]
boston_df_out.shape
```









#### **Encoding Categorical Data Types**

- 1. Label Encoding (Ordinal Encoding) this type of encoding can be used when the data consists of ordinal variables, ordinal encoding converts each label into integer values and the encoded data represents the sequence of labels.
- 2. One Hot Encoding for categorical variables where there is no ordinal relationship, integer coding may be insufficient, at best, or misleading to the model at worst.
  - The one-hot encoding transform is available in the scikit-learn Python machine learning library via the OneHotEncoder class.



#### **Encoding Categorical Data Types**

In one hot encoding, for each level of a categorical feature, we create a new variable.

Each category is mapped with binary numbers (0 or 1). Here, 0 represents the

absence, and 1 represents the presence of that category.



#### One Hot Encoding Example





### Label/Ordinal Encoding Example





## Feature Scaling





#### Feature Scaling Techniques

- Normalization is a scaling technique in which values are shifted and rescaled so that they end up ranging between 0 and 1. It is also known as Min-Max scaling.
- Here's the formula for normalization:

$$X' = \frac{X - X_{min}}{X_{max} - X_{min}}$$



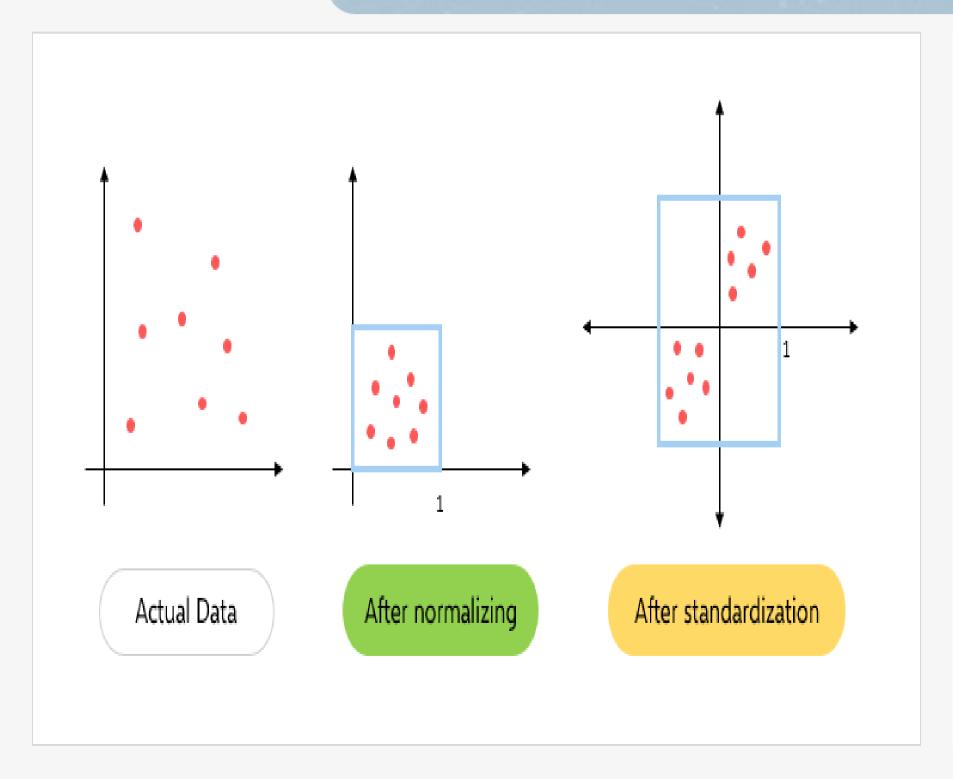
#### Feature Scaling Techniques

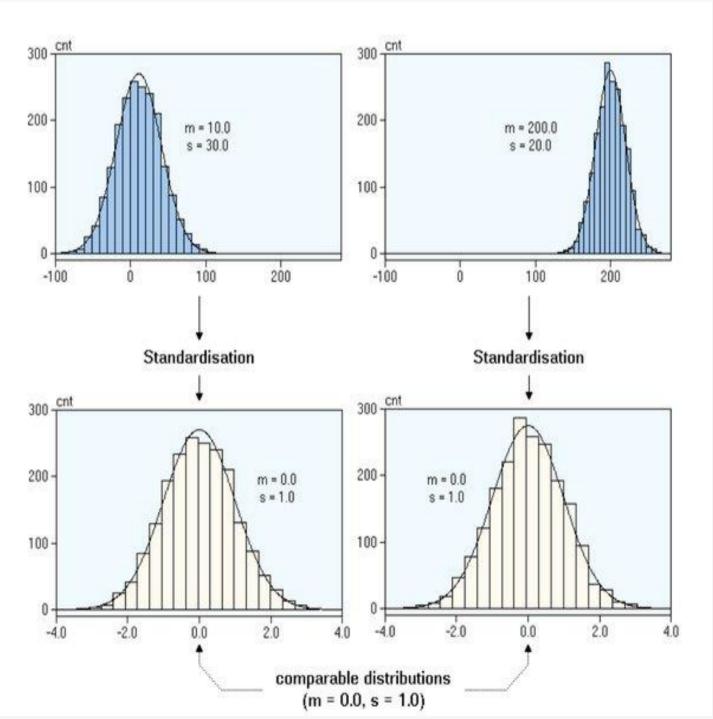
- 2. Standardization is another scaling technique where the values are cantered around the mean with a unit standard deviation. This means that the mean of the attribute becomes zero and the resultant distribution has a unit standard deviation.
- Here's the formula for standardization:

$$X^{'} = \frac{X - \mu}{\sigma}$$



#### Feature Scaling Techniques







#### Normalization using Sklearn

• To normalize your data, you need to import the MinMaxScalar from the sklearn

library and apply it to our dataset.

```
# data normalization with sklearn
from sklearn.preprocessing import
MinMaxScaler

# fit scaler on training data
norm = MinMaxScaler().fit(X_train)

# transform training data

X_train_norm =
norm.transform(X_train)

# transform testing dataabs

X_test_norm =
norm.transform(X_test)
```



#### Standardization using Sklearn

• To standardize your data, you need to import the **StandardScaler** from the sklearn library and apply it to our dataset.

```
# data standardization with sklearn
from sklearn.preprocessing import StandardScaler

# copy of datasets

X_train_stand = X_train.copy()

X_test_stand = X_test.copy()

# numerical features
num_cols =
['Item_Weight','Item_Visibility','Item_MRP','Outlet_Establish
ment_Year']
```



#### Standardization using Sklearn

```
# apply standardization on numerical features
for i in num_cols:
   # fit on training data column
   scale = StandardScaler().fit(X_train_stand[[i]])
   # transform the training data column
  X train stand[i] = scale.transform(X train stand[[i]])
   # transform the testing data column
  X_test_stand[i] = scale.transform(X_test_stand[[i]])
```



#### References

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# THANKYOU