Data Analysis with Python

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In this article, we will discuss how to do data analysis with Python. We will discuss all sorts of data analysis i.e. analyzing numerical data with NumPy, Tabular data with Pandas, data visualization Matplotlib, and Exploratory data analysis.

Data Analysis With Python

Data Analysis is the technique of collecting, transforming, and organizing data to make future predictions and informed data-driven decisions. It also helps to find possible solutions for a business problem. There are six steps for Data Analysis. They are:

- Ask or Specify Data Requirements
- Prepare or Collect Data
- Clean and Process
- Analyze
- Share
- Act or Report

Data Analysis with Python

Note: To know more about these steps refer to our Six Steps of Data Analysis Process tutorial.

Analyzing Numerical Data with NumPy

NumPy is an array processing package in Python and provides a highperformance multidimensional array object and tools for working with these arrays. It is the fundamental package for scientific computing with Python.

Arrays in NumPy

NumPy Array is a table of elements (usually numbers), all of the same types, indexed by a tuple of positive integers. In Numpy, the number of dimensions of the array is called the rank of the array. A tuple of integers giving the size of the array along each dimension is known as the shape of the array.

Creating NumPy Array

NumPy arrays can be created in multiple ways, with various ranks. It can also be created with the use of different data types like lists, tuples, etc. The type of the resultant array is deduced from the type of elements in the sequences. NumPy offers several functions to create arrays with initial placeholder content. These minimize the necessity of growing arrays, an expensive operation.

Create Array using numpy.empty/shape, dtype=float, order='C')



import numpy as np

```
a = np.empty([2, 2], dtype = int)
print("\nMatrix a : \n", a)
b = np.empty(2, dtype = int)
print("Matrix b : \n", b)
```

```
Matrix a:
 Π
        94655291709206
                                          0]
 [3543826506195694713
                        34181816989462323]]
Matrix b:
```

Create Array using numpy.zeros(shape, dtype = None, order = 'C')



import numpy as np

```
a = np.zeros([2, 2], dtype = int)
print("\nMatrix a : \n", a)
b = np.zeros(2, dtype = int)
print("Matrix b : \n", b)
c = np.zeros([3, 3])
print("\nMatrix c : \n", c)
```

Output

```
Matrix a:
[[0 0]
[0 0]]

Matrix b:
[0 0]

Matrix c:
[[0. 0. 0.]
[0. 0. 0.]]
```

Operations on Numpy Arrays

Arithmetic Operations

Addition:



import numpy as np

```
a = np.array([5, 72, 13, 100])
b = np.array([2, 5, 10, 30])
add_ans = a+b
```

```
print(add_ans)
add_ans = np.add(a, b)
print(add_ans)

c = np.array([1, 2, 3, 4])
add_ans = a+b+c
print(add_ans)

add_ans = np.add(a, b, c)
print(add_ans)
```

Output

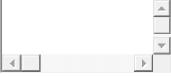
```
      [
      7
      77
      23
      130]

      [
      7
      77
      23
      130]

      [
      8
      79
      26
      134]

      [
      7
      77
      23
      130]
```

• Subtraction:



import numpy as np

```
a = np.array([5, 72, 13, 100])
b = np.array([2, 5, 10, 30])

sub_ans = a-b
print(sub_ans)

sub_ans = np.subtract(a, b)
print(sub_ans)
```

Output

```
[ 3 67 3 70]
[ 3 67 3 70]
```

• Multiplication:

```
△
∀
I
```

import numpy as np

```
a = np.array([5, 72, 13, 100])
b = np.array([2, 5, 10, 30])
```

mul_ans = a*b
print(mul_ans)

mul_ans = np.multiply(a, b)
print(mul_ans)

Output

```
[ 10 360 130 3000]
[ 10 360 130 3000]
```

Division:



import numpy as np

a = np.array([5, 72, 13, 100]) b = np.array([2, 5, 10, 30])

div_ans = a/b
print(div_ans)

div_ans = np.divide(a, b)
print(div_ans)

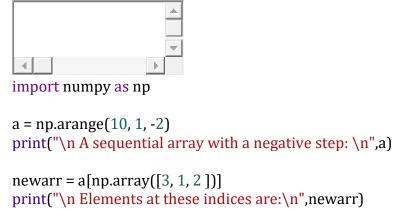
Output

[2.5	14.4	1.3	3.33333333]
[2.5	14.4	1.3	3.33333333]

For more information, refer to our <u>NumPy – Arithmetic Operations Tutorial</u> **NumPy Array Indexing**

Indexing can be done in NumPy by using an array as an index. In the case of the slice, a view or shallow copy of the array is returned but in the index array, a copy of the original array is returned. Numpy arrays can be indexed with other arrays or any other sequence with the exception of tuples. The last element is indexed by -1 second last by -2 and so on.

Python NumPy Array Indexing



Output

```
A sequential array with a negative step:

[10 8 6 4 2]

Elements at these indices are:

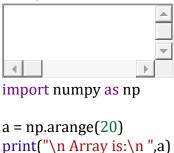
[4 8 6]
```

NumPy Array Slicing

Consider the syntax x[obj] where x is the array and obj is the index. The slice object is the index in the case of <u>basic slicing</u>. Basic slicing occurs when obj is:

- a slice object that is of the form start: stop: step
- an integer
- or a tuple of slice objects and integers

All arrays generated by basic slicing are always the view in the original array.



```
print("\n a[-8:17:1] = ",a[-8:17:1])
print("\n a[10:] = ",a[10:])
```

Output

```
Array is:

[ 0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19]

a[-8:17:1] = [12 13 14 15 16]

a[10:] = [10 11 12 13 14 15 16 17 18 19]

import numpy as np

a = np.arange(20)

print("\n Array is:\n",a)

print("\n a[-8:17:1] = ",a[-8:17:1])

print("\n a[10:] = ",a[10:])
```

Output

```
Array is:

[ 0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19]

a[-8:17:1] = [12 13 14 15 16]

a[10:] = [10 11 12 13 14 15 16 17 18 19]
```

Ellipsis can also be used along with basic slicing. Ellipsis (...) is the number of : objects needed to make a selection tuple of the same length as the dimensions of the array.



import numpy as np

Output

[[2 5] [8 11]]

NumPy Array Broadcasting

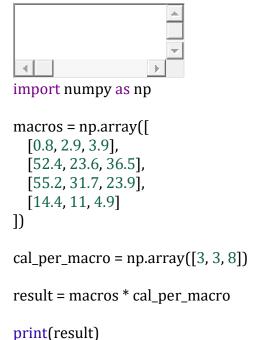
The term <u>broadcasting</u> refers to how numpy treats arrays with different Dimensions during arithmetic operations which lead to certain constraints, the smaller array is broadcast across the larger array so that they have compatible shapes.

Let's assume that we have a large data set, each datum is a list of parameters. In Numpy we have a 2-D array, where each row is a datum and the number of rows is the size of the data set. Suppose we want to apply some sort of scaling to all these data every parameter gets its own scaling factor or say Every parameter is multiplied by some factor. Just to have a clear understanding, let's count calories in foods using a macro-nutrient breakdown. Roughly put, the caloric parts of food are made of fats (9 calories per gram), protein (4 CPG), and carbs (4 CPG). So if we list some foods (our data), and for each food list its macro-nutrient breakdown (parameters), we can then multiply each nutrient by its caloric value (apply scaling) to compute the caloric breakdown of every food item.

_									
	Food	Fats (g)	Protein (g)	Carbs(g)		Food	Fats (g)	Protein (g)	Carbs(g)
	Apple	0.8	2.9	3.9		Apple	2.4	8.7	31.2
	Banana	52.4	23.6	36.5	[3, 3, 8]	Banana	157.2	70.8	292
	Raw Almond	55.2	31.7	23.9		Raw Almond	165.6	95.1	191.2
	Cookies	14.4	11	4.9		Cookies	43.2	33	39.2

With this transformation, we can now compute all kinds of useful information. For example, what is the total number of calories present in some food or, given a breakdown of my dinner know how many calories did I get from protein and so on.

Let's see a naive way of producing this computation with Numpy:



Output

```
[[ 2.4 8.7 31.2]
[157.2 70.8 292.]
[165.6 95.1 191.2]
[ 43.2 33. 39.2]]
```

Broadcasting Rules: Broadcasting two arrays together follow these rules:

- If the arrays don't have the same rank then prepend the shape of the lower rank array with 1s until both shapes have the same length.
- The two arrays are compatible in a dimension if they have the same size in the dimension or if one of the arrays has size 1 in that dimension.
- The arrays can be broadcast together if they are compatible with all dimensions.
- After broadcasting, each array behaves as if it had a shape equal to the element-wise maximum of shapes of the two input arrays.

• In any dimension where one array had a size of 1 and the other array had a size greater than 1, the first array behaves as if it were copied along that dimension.

```
import numpy as np

v = np.array([12, 24, 36])
w = np.array([45, 55])

print(np.reshape(v, (3, 1)) * w)

X = np.array([[12, 22, 33], [45, 55, 66]])

print(X + v)

print((X.T + w).T)

print(X * 2)
```

Output

```
[[ 540 660]
  [1080 1320]
  [1620 1980]]
  [[ 24 46 69]
  [ 57 79 102]]
  [[ 57 67 78]
  [100 110 121]]
  [[ 24 44 66]
  [ 90 110 132]]
```

Note: For more information, refer to our <u>Python NumPy Tutorial</u>.

Analyzing Data Using Pandas

Python Pandas Is used for relational or labeled data and provides various data structures for manipulating such data and time series. This library is built on top of the NumPy library. This module is generally imported as:

```
import pandas as pd
```

Here, pd is referred to as an alias to the Pandas. However, it is not necessary to import the library using the alias, it just helps in writing less

amount code every time a method or property is called. Pandas generally provide two data structures for manipulating data, They are:

- Series
- Dataframe

Series:

<u>Pandas Series</u> is a one-dimensional labeled array capable of holding data of any type (integer, string, float, python objects, etc.). The axis labels are collectively called indexes. Pandas Series is nothing but a column in an excel sheet. Labels need not be unique but must be a hashable type. The object supports both integer and label-based indexing and provides a host of methods for performing operations involving the index.

Pandas Series It can be created using the Series() function by loading the dataset from the existing storage like SQL, Database, CSV Files, Excel Files, etc., or from data structures like lists, dictionaries, etc. **Python Pandas Creating Series**

```
import pandas as pd
import numpy as np

ser = pd.Series(dtype="object")

print(ser)

data = np.array(['g', 'e', 'e', 'k', 's'])

ser = pd.Series(data)
print(ser)
```

Output

```
Series([], dtype: object)

0 g

1 e

2 e

3 k

4 s

dtype: object
```

Dataframe:

<u>Pandas DataFrame</u> is a two-dimensional size-mutable, potentially heterogeneous tabular data structure with labeled axes (rows and columns). A Data frame is a two-dimensional data structure, i.e., data is aligned in a tabular fashion in rows and columns. Pandas DataFrame consists of three principal components, the data, rows, and columns.



```
import pandas as pd

df = pd.DataFrame()
print(df)

lst = ['Geeks', 'For', 'Geeks', 'is', 'portal', 'for', 'Geeks']

df = pd.DataFrame(lst, columns=['Words'])
print(df)
```

Output

```
Empty DataFrame
Columns: []
Index: []
    Words
    Geeks
0
1
      For
2
    Geeks
3
       is
4
  portal
      for
5
    Geeks
```

Creating Dataframe from CSV

We can <u>create a dataframe from the CSV</u> files using the <u>read_csv()</u> function.

Python Pandas read CSV

```
import pandas as pd

df = pd.read_csv("Iris.csv")

df.head()
```

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

head

of a dataframe

Filtering DataFrame

Pandas <u>dataframe.filter()</u> function is used to Subset rows or columns of dataframe according to labels in the specified index. Note that this routine does not filter a dataframe on its contents. The filter is applied to the labels of the index.

Python Pandas Filter Dataframe

```
import pandas as pd

df = pd.read_csv("Iris.csv")

df.filter(["Species", "SepalLengthCm", "SepalLengthCm"]).head()
```

Output:

	Species	SepalLengthCm	SepalLengthCm
0	Iris-setosa	5.1	5.1
1	Iris-setosa	4.9	4.9
2	Iris-setosa	4.7	4.7
3	Iris-setosa	4.6	4.6
4	Iris-setosa	5.0	5.0

Applying filter on dataset

Sorting DataFrame

In order to sort the data frame in pandas, the function <u>sort_values()</u> is used. Pandas sort_values() can sort the data frame in Ascending or Descending order.

Python Pandas Sorting Dataframe in Ascending Order

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
13	14	4.3	3.0	1.1	0.1	Iris-setosa
42	43	4.4	3.2	1.3	0.2	Iris-setosa
38	39	4.4	3.0	1.3	0.2	Iris-setosa
8	9	4.4	2.9	1.4	0.2	Iris-setosa
41	42	4.5	2.3	1.3	0.3	Iris-setosa
122	123	7.7	2.8	6.7	2.0	Iris-virginica
118	119	7.7	2.6	6.9	2.3	Iris-virginica
117	118	7.7	3.8	6.7	2.2	Iris-virginica
135	136	7.7	3.0	6.1	2.3	Iris-virginica
131	132	7.9	3.8	6.4	2.0	Iris-virginica

150 rows × 6 columns

Sorted dataset based on a column value

Pandas GroupBy

<u>Groupby</u> is a pretty simple concept. We can create a grouping of categories and apply a function to the categories. In real data science projects, you'll be dealing with large amounts of data and trying things over and over, so for efficiency, we use the Groupby concept. Groupby mainly refers to a process involving one or more of the following steps they are:

- **Splitting:** It is a process in which we split data into group by applying some conditions on datasets.
- **Applying:** It is a process in which we apply a function to each group independently.
- **Combining:** It is a process in which we combine different datasets after applying groupby and results into a data structure.

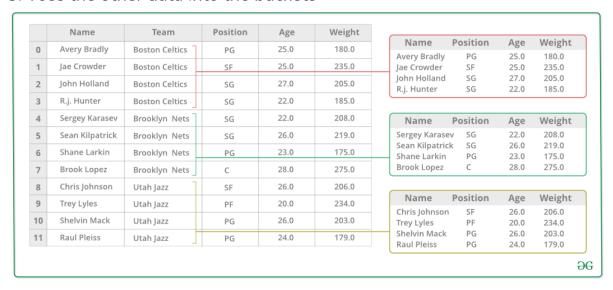
The following image will help in understanding the process involve in the Groupby concept.

1. Group the unique values from the Team column

Pandas Groupby Method

2. Now there's a bucket for each group

3. Toss the other data into the buckets



4. Apply a function on the weight column of each bucket.

Applying Function on the weight column of each column

Python Pandas GroupBy:

Orginal Dataframe

	Name	Age	Address	Qualification
0	Jai	27	Nagpur	Msc
1	Anuj	24	Kanpur	MA
2	Jai	22	Allahabad	MCA
3	Princi	32	Kannuaj	Phd
4	Gaurav	33	Jaunpur	B.Tech
5	Anuj	36	Kanpur	B.com
6	Princi	27	Allahabad	Msc
7	Abhi	32	Aligarh	MA

After Creating Groups

	Age	Address	Qualification
Name			
Abhi	32	Aligarh	MA
Anuj	24	Kanpur	MA
Gaurav	33	Jaunpur	B.Tech
Jai	27	Nagpur	Msc
Princi	32	Kannuaj	Phd

pandas groupby

Applying function to group:

After splitting a data into a group, we apply a function to each group in order to do that we perform some operations they are:

- Aggregation: It is a process in which we compute a summary statistic (or statistics) about each group. For Example, Compute group sums or means
- Transformation: It is a process in which we perform some groupspecific computations and return a like-indexed. For Example, Filling NAs within groups with a value derived from each group
- Filtration: It is a process in which we discard some groups, according to a group-wise computation that evaluates True or False. For Example, Filtering out data based on the group sum or mean

Pandas Aggregation

<u>Aggregation</u> is a process in which we compute a summary statistic about each group. The aggregated function returns a single aggregated value for each group. After splitting data into groups using groupby function, several aggregation operations can be performed on the grouped data.

Python Pandas Aggregation

Output:

-Use of sum aggregate function on dataset

Concatenating DataFrame

In order to concat the dataframe, we use <u>concat()</u> function which helps in concatenating the dataframe. This function does all the heavy lifting of performing concatenation operations along with an axis of Pandas objects while performing optional set logic (union or intersection) of the indexes (if any) on the other axes.

Python Pandas Concatenate Dataframe

Output:

	key	Name	Age
0	K0	Jai	27
1	K1	Princi	24
2	K2	Gaurav	22
3	К3	Anuj	32

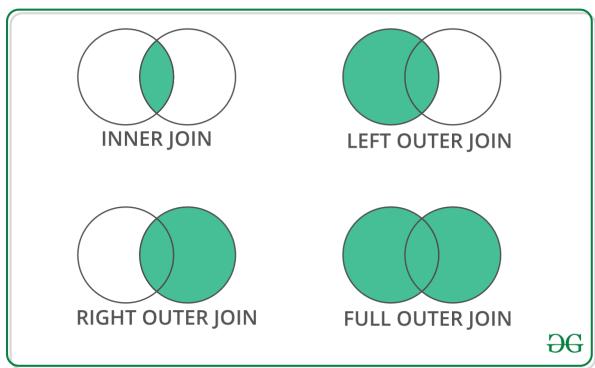
	key	Address	Qualification
0	K0	Nagpur	Btech
1	K1	Kanpur	B.A
2	K2	Allahabad	Bcom
3	КЗ	Kannuaj	B.hons

	key	Name	Age	key	Address	Qualification
() КО	Jai	27	К0	Nagpur	Btech
1	L K1	Princi	24	K1	Kanpur	B.A
2	2 K2	Gaurav	22	K2	Allahabad	Bcom
3	3 K3	Anuj	32	К3	Kannuaj	B.hons

Merging DataFrame

When we need to combine very large DataFrames, joins serve as a powerful way to perform these operations swiftly. Joins can only be done on two DataFrames at a time, denoted as left and right tables. The key is the common column that the two DataFrames will be joined on. It's a good practice to use keys that have unique values throughout the column to avoid unintended duplication of row values. Pandas provide a single function, merge(), as the entry point for all standard database join operations between DataFrame objects.

There are four basic ways to handle the join (inner, left, right, and outer), depending on which rows must retain their data.



Python Pandas Merge Dataframe

	key	Name	Age
0	K0	Jai	27
1	K1	Princi	24
2	K2	Gaurav	22
3	КЗ	Anuj	32

	key	Address	Qualification
0	K0	Nagpur	Btech
1	K1	Kanpur	B.A
2	K2	Allahabad	Bcom
3	КЗ	Kannuaj	B.hons

	key	Name	Age	Address	Qualification
0	K0	Jai	27	Nagpur	Btech
1	K1	Princi	24	Kanpur	B.A
2	K2	Gaurav	22	Allahabad	Bcom
3	К3	Anuj	32	Kannuaj	B.hons

Concatinating Two datasets

Joining DataFrame

In order to join the dataframe, we use <u>.join()</u> function this function is used for combining the columns of two potentially differently indexed DataFrames into a single result DataFrame.

Python Pandas Join Dataframe

	Name	Age
K0	Jai	27
K1	Princi	24
K2	Gaurav	22
К3	Anuj	32

	Address	Qualification
K0	Allahabad	MCA
K2	Kannuaj	Phd
К3	Allahabad	Bcom
K4	Kannuaj	B.hons

	Name	Age	Address	Qualification
K0	Jai	27	Allahabad	MCA
K1	Princi	24	NaN	NaN
K2	Gaurav	22	Kannuaj	Phd
К3	Anuj	32	Allahabad	Bcom

Joining two datasets

For more information, refer to our <u>Pandas Merging</u>, <u>Joining</u>, <u>and</u> <u>Concatenating</u> tutorial

For a complete guide on Pandas refer to our <u>Pandas Tutorial</u>.

Visualization with Matplotlib

Matplotlib is easy to use and an amazing visualizing library in Python. It is built on NumPy arrays and designed to work with the broader SciPy stack and consists of several plots like line, bar, scatter, histogram, etc.

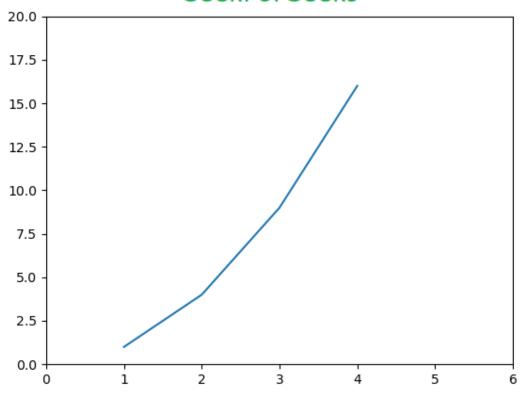
Pyplot

<u>Pyplot</u> is a Matplotlib module that provides a MATLAB-like interface. Pyplot provides functions that interact with the figure i.e. creates a figure, decorates the plot with labels, and creates a plotting area in a figure.

```
import matplotlib.pyplot as plt

plt.plot([1, 2, 3, 4], [1, 4, 9, 16])
plt.axis([0, 6, 0, 20])
plt.show()
```





Bar chart

A <u>bar plot</u> or bar chart is a graph that represents the category of data with rectangular bars with lengths and heights that is proportional to the values which they represent. The bar plots can be plotted horizontally or vertically. A bar chart describes the comparisons between the discrete categories. It can be created using the bar() method.

Python Matplotlib Bar Chart Here we will use the iris dataset only

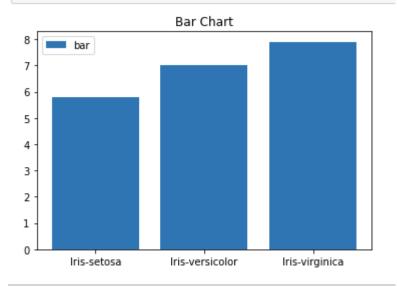
```
import matplotlib.pyplot as plt
import pandas as pd

df = pd.read_csv("Iris.csv")

plt.bar(df['Species'], df['SepalLengthCm'])

plt.title("Iris Dataset")

plt.legend(["bar"])
plt.show()
```



Bar chart using matplotlib

library

Histograms

A <u>histogram</u> is basically used to represent data in the form of some groups. It is a type of bar plot where the X-axis represents the bin ranges while the Y-axis gives information about frequency. To create a histogram the first step is to create a bin of the ranges, then distribute the whole range of the values into a series of intervals, and count the values which fall into each of the intervals. Bins are clearly identified as consecutive, non-overlapping intervals of variables. The <u>hist()</u> function is used to compute and create a histogram of x.

Python Matplotlib Histogram

```
import matplotlib.pyplot as plt
import pandas as pd

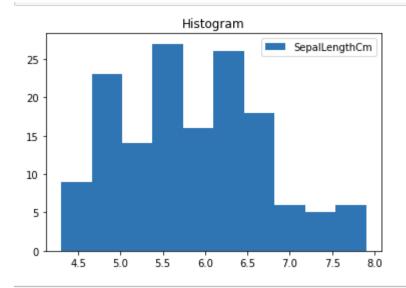
df = pd.read_csv("Iris.csv")

plt.hist(df["SepalLengthCm"])

plt.title("Histogram")

plt.legend(["SepalLengthCm"])

plt.show()
```



Histplot using

matplotlib library

Scatter Plot

Scatter plots are used to observe relationship between variables and uses dots to represent the relationship between them. The <u>scatter()</u> method in the matplotlib library is used to draw a scatter plot.

Python Matplotlib Scatter Plot

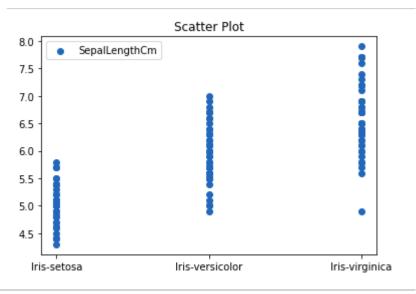
```
import matplotlib.pyplot as plt
import pandas as pd

df = pd.read_csv("Iris.csv")

plt.scatter(df["Species"], df["SepalLengthCm"])

plt.title("Scatter Plot")

plt.legend(["SepalLengthCm"])
plt.show()
```



Scatter plot using

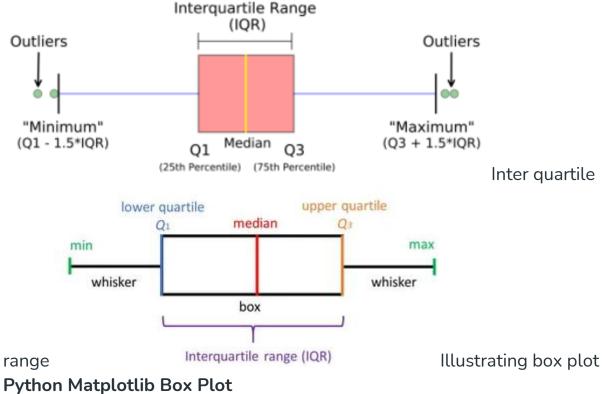
matplotlib library

Box Plot

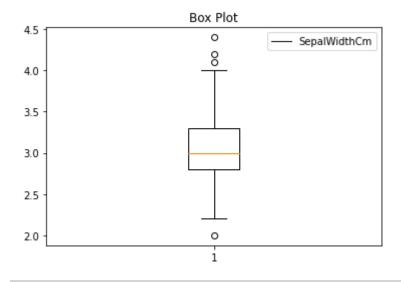
A <u>boxplot</u>, Correlation also known as a box and whisker plot. It is a very good visual representation when it comes to measuring the data distribution. Clearly plots the median values, outliers and the quartiles. Understanding data distribution is another important factor which leads to better model building. If data has outliers, box plot is a recommended way to identify them and take necessary actions. The box and whiskers chart shows how data is spread out. Five pieces of information are generally included in the chart

- The minimum is shown at the far left of the chart, at the end of the left 'whisker'
- First quartile, Q1, is the far left of the box (left whisker)
- The median is shown as a line in the center of the box
- Third quartile, Q3, shown at the far right of the box (right whisker)
- The maximum is at the far right of the box

Representation of box plot



```
import matplotlib.pyplot as plt
import pandas as pd
df = pd.read_csv("Iris.csv")
plt.boxplot(df["SepalWidthCm"])
plt.title("Box Plot")
plt.legend(["SepalWidthCm"])
plt.show()
```



Boxplot using

matplotlib library

Correlation Heatmaps

A 2-D Heatmap is a data visualization tool that helps to represent the magnitude of the phenomenon in form of colors. A correlation heatmap is a heatmap that shows a 2D correlation matrix between two discrete dimensions, using colored cells to represent data from usually a monochromatic scale. The values of the first dimension appear as the rows of the table while the second dimension is a column. The color of the cell is proportional to the number of measurements that match the dimensional value. This makes correlation heatmaps ideal for data analysis since it makes patterns easily readable and highlights the differences and variation in the same data. A correlation heatmap, like a regular heatmap, is assisted by a colorbar making data easily readable and comprehensible.

Note: The data here has to be passed with corr() method to generate a correlation heatmap. Also, corr() itself eliminates columns that will be of no use while generating a correlation heatmap and selects those which can be used.

Python Matplotlib Correlation Heatmap

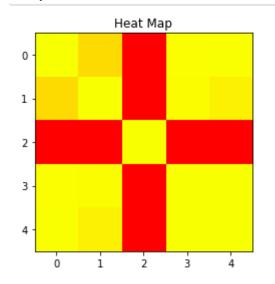
```
import matplotlib.pyplot as plt
import pandas as pd

df = pd.read_csv("Iris.csv")

plt.imshow(df.corr() , cmap = 'autumn' , interpolation = 'nearest' )

plt.title("Heat Map")
plt.show()
```

Output:



Heatmap using matplotlib library

For more information on data visualization refer to our below tutorials

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- Data Visualization using Matplotlib
- Data Visualization with Python Seaborn
- Data Visualisation in Python using Matplotlib and Seaborn
- Using Plotly for Interactive Data Visualization in Python
- Interactive Data Visualization with Bokeh

Exploratory Data Analysis

Exploratory Data Analysis (EDA) is a technique to analyze data using some visual Techniques. With this technique, we can get detailed information about the statistical summary of the data. We will also be able to deal with the duplicates values, outliers, and also see some trends or patterns present in the dataset.

Note: We will be using Iris Dataset.

Getting Information about the Dataset

We will use the shape parameter to get the shape of the dataset.

Shape of Dataframe

df.shape

Output:

(150, 6)

We can see that the dataframe contains 6 columns and 150 rows. Now, let's also the columns and their data types. For this, we will use the info() method.

Information about Dataset

df.info()

Output:

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 6 columns):
    Column
                  Non-Null Count Dtype
    -----
                   -----
                  150 non-null
                                  int64
 0
    Ιd
 1
    SepalLengthCm 150 non-null
                                 float64
                   150 non-null
 2
    SepalWidthCm
                                  float64
 3
    PetalLengthCm 150 non-null
                                  float64
    PetalWidthCm
                                 float64
 4
                  150 non-null
 5
    Species
                   150 non-null
                                object
dtypes: float64(4), int64(1), object(1)
memory usage: 7.2+ KB
```

information about the

dataset

We can see that only one column has categorical data and all the other columns are of the numeric type with non-Null entries.

Let's get a quick statistical summary of the dataset using the <u>describe()</u> method. The describe() function applies basic statistical computations on the dataset like extreme values, count of data points standard deviation, etc. Any missing value or NaN value is automatically skipped. describe() function gives a good picture of the distribution of data.

Description of dataset

df.describe()

Output:

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
count	150.000000	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.054000	3.758667	1.198667
std	43.445368	0.828066	0.433594	1.764420	0.763161
min	1.000000	4.300000	2.000000	1.000000	0.100000
25%	38.250000	5.100000	2.800000	1.600000	0.300000
50%	75.500000	5.800000	3.000000	4.350000	1.300000
75%	112.750000	6.400000	3.300000	5.100000	1.800000
max	150.000000	7.900000	4.400000	6.900000	2.500000

Description about the dataset

We can see the count of each column along with their mean value, standard deviation, minimum and maximum values.

Checking Missing Values

We will check if our data contains any missing values or not. Missing values can occur when no information is provided for one or more items or for a whole unit. We will use the isnull() method.

python code for missing value

df.isnull().sum()

Output:

Id	0
SepalLengthCm	0
SepalWidthCm	0
PetalLengthCm	0
PetalWidthCm	0
Species	0
dtype: int64	

Missing values in the dataset

We can see that no column has any missing value.

Checking Duplicates

Let's see if our dataset contains any duplicates or not. Pandas <u>drop_duplicates()</u> method helps in removing duplicates from the data frame.

Pandas function for missing values

```
data = df.drop_duplicates(subset ="Species",)
data
```

Output:

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
50	51	7.0	3.2	4.7	1.4	Iris-versicolor
100	101	6.3	3.3	6.0	2.5	Iris-virginica

Dropping duplicate value in the dataset

We can see that there are only three unique species. Let's see if the dataset is balanced or not i.e. all the species contain equal amounts of rows or not. We will use the Series.value_counts() function. This function returns a Series containing counts of unique values.

Python code for value counts in the column

df.value counts("Species")

Output:

Species
Iris-setosa 50
Iris-versicolor 50
Iris-virginica 50
dtype: int64

_____value count in the dataset

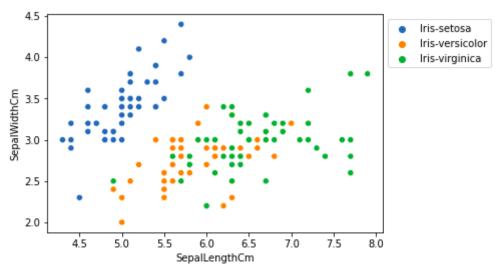
We can see that all the species contain an equal amount of rows, so we should not delete any entries.

Relation between variables

We will see the relationship between the sepal length and sepal width and also between petal length and petal width.

Comparing Sepal Length and Sepal Width

Output:



Scatter plot

using matplotlib library

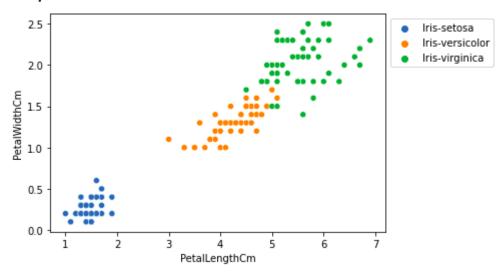
From the above plot, we can infer that -

- Species Setosa has smaller sepal lengths but larger sepal widths.
- Versicolor Species lies in the middle of the other two species in terms of sepal length and width
- Species Virginica has larger sepal lengths but smaller sepal widths.

Comparing Petal Length and Petal Width

```
plt.legend(bbox_to_anchor=(1, 1), loc=2)
plt.show()
```

Output:



sactter plot

petal length

From the above plot, we can infer that -

- The species Setosa has smaller petal lengths and widths.
- Versicolor Species lies in the middle of the other two species in terms of petal length and width
- Species Virginica has the largest petal lengths and widths.

Let's plot all the column's relationships using a pairplot. It can be used for multivariate analysis.

Python code for pairplot

Pairplot for the dataset

We can see many types of relationships from this plot such as the species Seotsa has the smallest of petals widths and lengths. It also has the smallest sepal length but larger sepal widths. Such information can be gathered about any other species.

Handling Correlation

Pandas <u>dataframe.corr()</u> is used to find the pairwise correlation of all columns in the dataframe. Any NA values are automatically excluded. Any non-numeric data type columns in the dataframe are ignored.

Example:

```
data.corr(method='pearson')
```

Output:

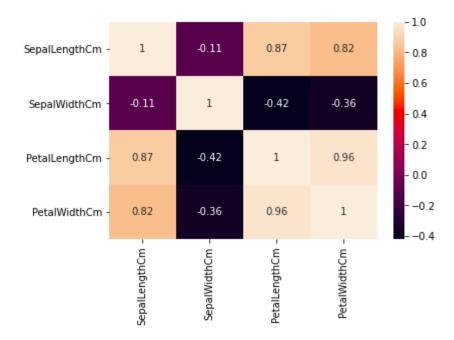
	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
ld	1.000000	0.624413	-0.654654	0.969909	0.999685
SepalLengthCm	0.624413	1.000000	-0.999226	0.795795	0.643817
SepalWidthCm	-0.654654	-0.999226	1.000000	-0.818999	-0.673417
PetalLengthCm	0.969909	0.795795	-0.818999	1.000000	0.975713
PetalWidthCm	0.999685	0.643817	-0.673417	0.975713	1.000000

correlation between columns in the dataset

Heatmaps

The heatmap is a data visualization technique that is used to analyze the dataset as colors in two dimensions. Basically, it shows a correlation between all numerical variables in the dataset. In simpler terms, we can plot the above-found correlation using the heatmaps.

python code for heatmap



Heatmap for

correlation in the dataset

From the above graph, we can see that -

- Petal width and petal length have high correlations.
- Petal length and sepal width have good correlations.
- Petal Width and Sepal length have good correlations.

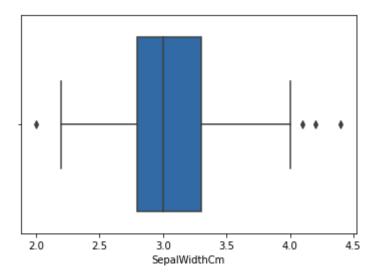
Handling Outliers

An Outlier is a data item/object that deviates significantly from the rest of the (so-called normal)objects. They can be caused by measurement or execution errors. The analysis for outlier detection is referred to as outlier mining. There are many ways to detect outliers, and the removal process is the data frame same as removing a data item from the panda's dataframe. Let's consider the iris dataset and let's plot the boxplot for the SepalWidthCm column.

python code for Boxplot

```
import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('Iris.csv')
sns.boxplot(x='SepalWidthCm', data=df)
```



Boxplot for sepalwidth

column

In the above graph, the values above 4 and below 2 are acting as outliers.

Removing Outliers

For removing the outlier, one must follow the same process of removing an entry from the dataset using its exact position in the dataset because in all the above methods of detecting the outliers end result is the list of all those data items that satisfy the outlier definition according to the method used.

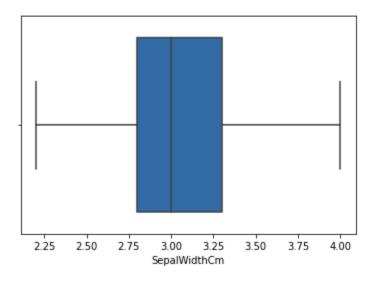
We will detect the outliers using <u>IQR</u> and then we will remove them. We will also draw the boxplot to see if the outliers are removed or not.

```
print("New Shape: ", df.shape)
sns.boxplot(x='SepalWidthCm', data=df)
```

Output:

```
Old Shape: (150, 6)
New Shape: (146, 6)
```

<AxesSubplot:xlabel='SepalWidthCm'>



boxplot using seaborn

library