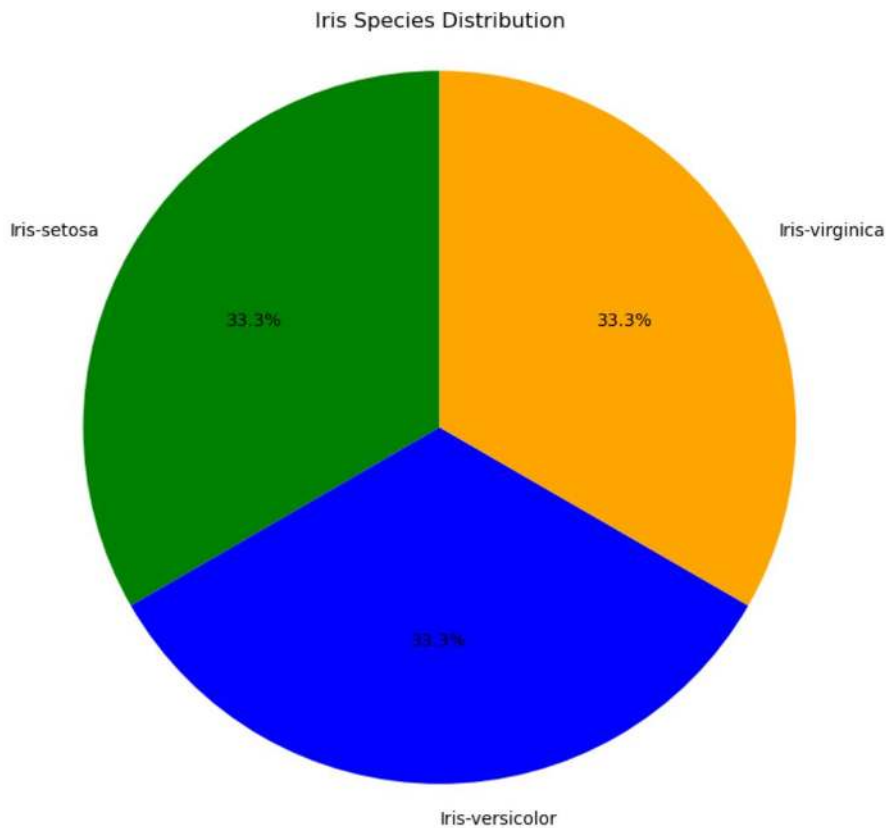


## SLIP 1

Q.2 A) Write a Python program to create a Pie plot to get the frequency of the three species of the Iris data (Use iris.csv)

```
In [8]: 1 from sklearn.datasets import load_iris
2 import pandas as pd
3 import matplotlib.pyplot as plt
4
5 # Load the Iris data from the CSV file
6 iris_data = pd.read_csv('Iris.csv')
7 # Get the frequency of each species
8 species_counts = iris_data['Species'].value_counts()
9
10 ## Create a Pie plot
11 plt.figure(figsize=(8, 8))
12 plt.pie(species_counts, labels=species_counts.index, autopct='%1.1f%%', startangle=90, colors=['green', 'blue', 'orange'])
13 plt.title('Iris Species Distribution')
14 plt.axis('equal') # Equal aspect ratio ensures that pie is drawn as a circle.
15 plt.show()
16
```



B) Write a Python program to view basic statistical details of the data.(Use winequality-red.csv)

```
In [9]: 1 import pandas as pd
2
3 # Load the wine data from the CSV file
4 wine_data = pd.read_csv('winequality-red.csv')
5
6 # Display basic statistical details of the data
7 statistical_details = wine_data.describe()
8 print(statistical_details)
9
```

	fixed acidity	volatile acidity	citric acid	residual sugar	\
count	10.000000	10.0000	10.000000	10.000000	
mean	7.950000	0.631	0.104000	2.330000	
std	1.162612	0.161	0.194548	1.376025	
min	7.300000	0.280	0.000000	1.200000	
25%	7.400000	0.585	0.000000	1.825000	
50%	7.650000	0.655	0.010000	1.900000	
75%	7.800000	0.700	0.055000	2.225000	
max	11.200000	0.880	0.560000	6.100000	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	\
count	10.000000	10.000000	10.000000	10.000000	
mean	0.077000	14.800000	48.900000	0.997080	
std	0.010198	4.467164	25.066356	0.001038	
min	0.065000	9.000000	18.000000	0.994600	
25%	0.071500	11.500000	34.000000	0.996800	
50%	0.075000	15.000000	47.000000	0.997400	
75%	0.076000	16.500000	59.750000	0.997800	
max	0.098000	25.000000	102.000000	0.998000	

	pH	sulphates	alcohol	quality
count	10.000000	10.000000	10.000000	10.000000
mean	3.355000	0.589000	9.700000	5.500000
std	0.127997	0.100161	0.359011	0.849837
min	3.160000	0.460000	9.400000	5.000000
25%	3.270000	0.560000	9.400000	5.000000
50%	3.355000	0.565000	9.650000	5.000000
75%	3.480000	0.632500	9.800000	5.750000
max	3.510000	0.800000	10.500000	7.000000

## SLIP 2

Q.2 A) Write a Python program for Handling Missing Value. Replace missing value of salary, age column with mean of that column.(Use Data.csv file).

```
In [10]: 1 import pandas as pd
2
3 # Load the data from the CSV file
4 data = pd.read_csv('Data.csv')
5
6 # Display the original data
7 print("Original Data:")
8 print(data)
9
10 # Calculate the mean of the 'salary' and 'age' columns
11 mean_salary = data['salary'].mean()
12 mean_age = data['age'].mean()
13
14 # Replace missing values with the mean in the 'salary' and 'age' columns
15 data['salary'].fillna(mean_salary, inplace=True)
16 data['age'].fillna(mean_age, inplace=True)
17
18 # Display the data after handling missing values
19 print("\nData after handling missing values:")
20 print(data)
21
```

Original Data:

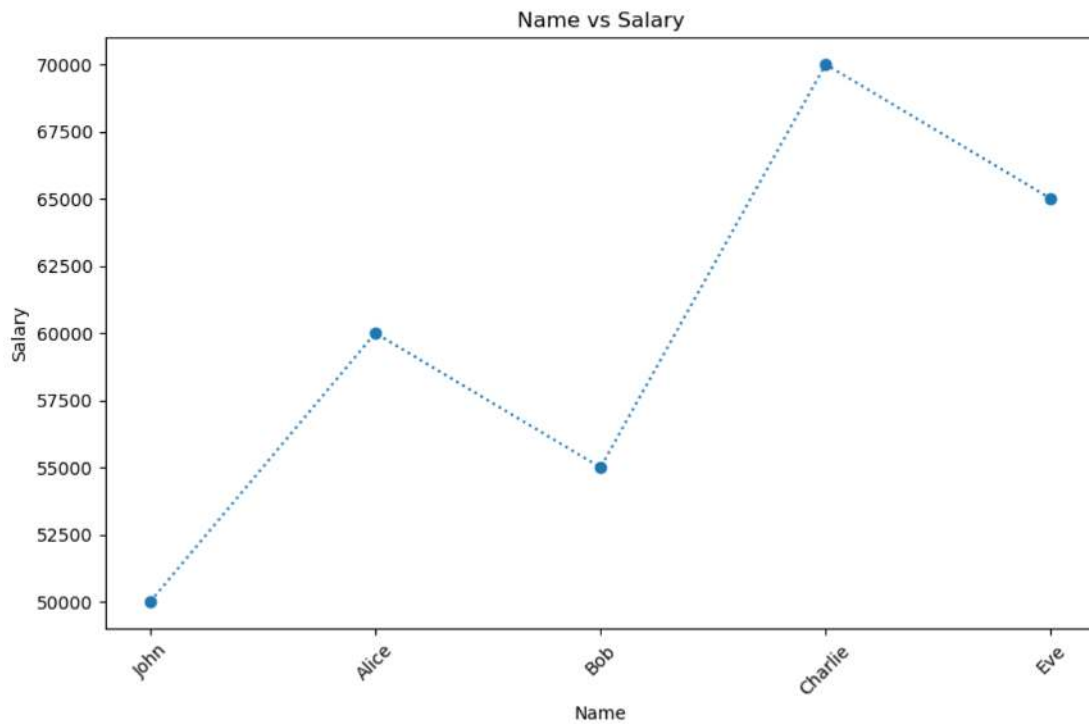
	name	salary	age	gender
0	John	50000.0	NaN	Male
1	Alice	NaN	25.0	Female
2	Bob	60000.0	30.0	NaN
3	Charlie	NaN	NaN	NaN
4	Eve	55000.0	28.0	Female

Data after handling missing values:

	name	salary	age	gender
0	John	50000.0	27.666667	Male
1	Alice	55000.0	25.000000	Female
2	Bob	60000.0	30.000000	NaN
3	Charlie	55000.0	27.666667	NaN
4	Eve	55000.0	28.000000	Female

Q.2 B) Write a Python program to generate a line plot of name Vs salary

```
In [16]: 1 import pandas as pd
2 import matplotlib.pyplot as plt
3
4 # Load the data from the CSV file
5 data = pd.read_csv('Data1.csv')
6
7 # Create a Line plot of name versus salary
8 plt.figure(figsize=(10, 6))
9 plt.plot(data['name'], data['salary'], linestyle='dotted', marker='o')
10 plt.title('Name vs Salary')
11 plt.xlabel('Name')
12 plt.ylabel('Salary')
13 plt.xticks(rotation=45) # Rotate x-axis Labels for better readability
14
15 # Show the plot
16 plt.show()
17
```



Q.2 C) Download the heights and weights dataset and load the dataset from a given csv file into a dataframe. Print the first, last 10 rows and random 20 rows also display shape of the dataset.

```
In [17]: 1 import pandas as pd
2
3 # Load the dataset from the CSV file
4 df = pd.read_csv('height_weight.csv')
5
6 # Display the first 10 rows
7 print("First 10 rows:")
8 print(df.head(10))
9
10 # Display the last 10 rows
11 print("\nLast 10 rows:")
12 print(df.tail(10))
13
14 # Display random 20 rows
15 print("\nRandom 20 rows:")
16 print(df.sample(20))
17
18 # Display the shape of the dataset
19 print("\nShape of the dataset:")
20 print(df.shape)
21
```

First 10 rows:

	Height	Weight
0	160	55
1	165	60
2	170	65
3	175	70
4	180	75
5	185	80
6	190	85
7	155	50
8	162	57
9	168	62

Last 10 rows:

	Height	Weight
10	172	68
11	178	73
12	182	78
13	188	83
14	195	90
15	150	45
16	158	52
17	164	58
18	169	63
19	174	69

Random 20 rows:

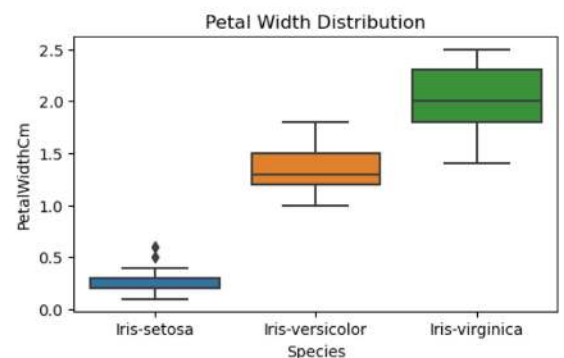
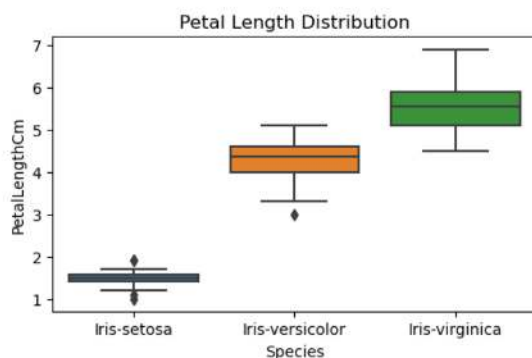
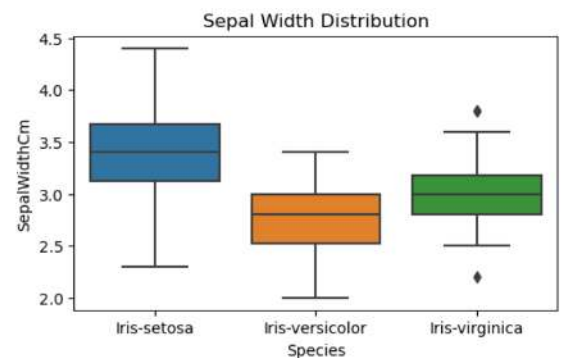
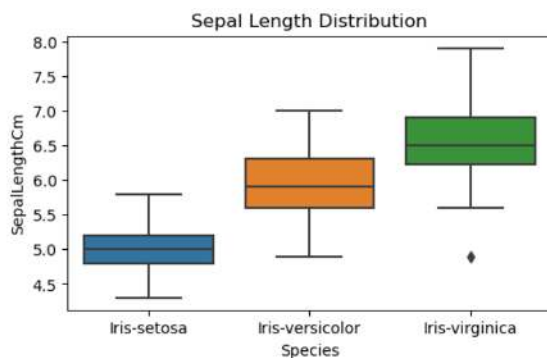
	Height	Weight
15	150	45
0	160	55
5	185	80
4	180	75
14	195	90
3	175	70
8	162	57
12	182	78
7	155	50
17	164	58
10	172	68
11	178	73
2	170	65
16	158	52
18	169	63
13	188	83
9	168	62
6	190	85
19	174	69
1	165	60

Shape of the dataset:

(20, 2)

Q.2 A) Write a Python program to create box plots to see how each feature i.e. Sepal Length, Sepal Width, Petal Length, Petal Width are distributed across the three species. (Use iris.csv dataset)

```
In [36]: 1 import pandas as pd
2 import seaborn as sns
3 import matplotlib.pyplot as plt
4
5 # Load the Iris dataset
6 iris_data = pd.read_csv('iris.csv')
7
8 # Create box plots for each feature across the three species
9 plt.figure(figsize=(14, 8))
10 plt.subplots_adjust(wspace=0.5, hspace=0.5) # Adjust the space between subplots for better readability
11
12 # Box plot for Sepal Length
13 plt.subplot(2, 2, 1)
14 sns.boxplot(x='Species', y='SepalLengthCm', data=iris_data)
15 plt.title('Sepal Length Distribution')
16
17 # Box plot for Sepal Width
18 plt.subplot(2, 2, 2)
19 sns.boxplot(x='Species', y='SepalWidthCm', data=iris_data)
20 plt.title('Sepal Width Distribution')
21
22 # Box plot for Petal Length
23 plt.subplot(2, 2, 3)
24 sns.boxplot(x='Species', y='PetalLengthCm', data=iris_data)
25 plt.title('Petal Length Distribution')
26
27 # Box plot for Petal Width
28 plt.subplot(2, 2, 4)
29 sns.boxplot(x='Species', y='PetalWidthCm', data=iris_data)
30 plt.title('Petal Width Distribution')
31
32 # Show the plots
33 plt.show()
34
35
```



Q.2 B) Write a Python program to view basic statistical details of the data (Use Heights and Weights Dataset)

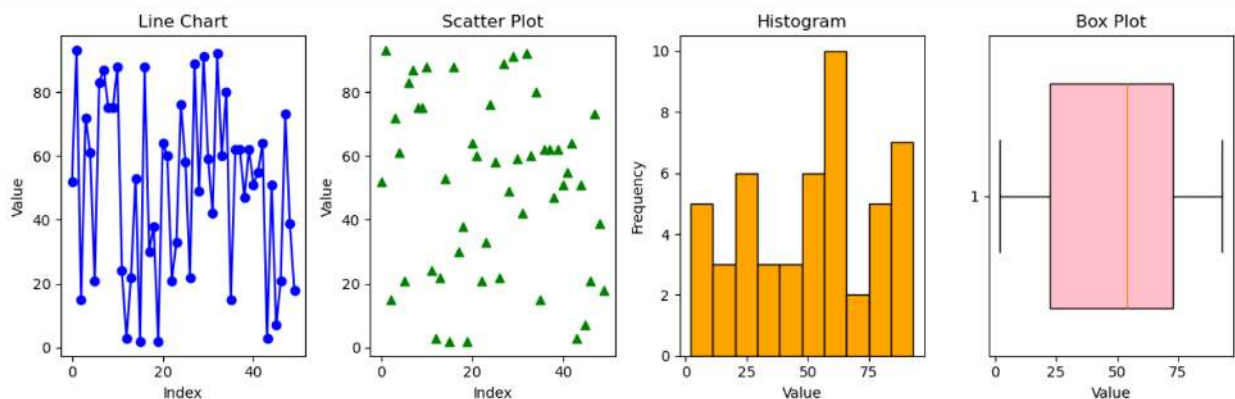
```
In [37]: 1 import pandas as pd
2
3 # Load the dataset from the CSV file
4 df = pd.read_csv('height_weight.csv')
5
6 # Display basic statistical details of the data
7 statistical_details = df.describe()
8 print(statistical_details)
9
```

	Height	Weight
count	20.000000	20.000000
mean	172.000000	66.900000
std	12.238851	12.417729
min	150.000000	45.000000
25%	163.500000	57.750000
50%	171.000000	66.500000
75%	180.500000	75.750000
max	195.000000	90.000000

## SLIP 4

Q.2 A) Generate a random array of 50 integers and display them using a line chart, scatter plot, histogram and box plot. Apply appropriate color, labels and styling options.

```
In [38]: 1 import numpy as np
2 import matplotlib.pyplot as plt
3
4 # Set a seed for reproducibility
5 np.random.seed(42)
6
7 # Generate a random array of 50 integers
8 random_array = np.random.randint(1, 100, 50)
9
10 # Line Chart
11 plt.figure(figsize=(12, 4))
12 plt.subplot(1, 4, 1)
13 plt.plot(random_array, marker='o', color='blue')
14 plt.title('Line Chart')
15 plt.xlabel('Index')
16 plt.ylabel('Value')
17
18 # Scatter Plot
19 plt.subplot(1, 4, 2)
20 plt.scatter(range(len(random_array)), random_array, color='green', marker='^')
21 plt.title('Scatter Plot')
22 plt.xlabel('Index')
23 plt.ylabel('Value')
24
25 # Histogram
26 plt.subplot(1, 4, 3)
27 plt.hist(random_array, bins=10, color='orange', edgecolor='black')
28 plt.title('Histogram')
29 plt.xlabel('Value')
30 plt.ylabel('Frequency')
31
32 # Box Plot
33 plt.subplot(1, 4, 4)
34 plt.boxplot(random_array, vert=False, widths=0.7, patch_artist=True, boxprops=dict(facecolor='pink'))
35 plt.title('Box Plot')
36 plt.xlabel('Value')
37
38 # Adjust Layout for better presentation
39 plt.tight_layout()
40
41 # Show the plots
42 plt.show()
43
```



Q.2 B) Write a Python program to print the shape, number of rows-columns, data types, feature names and the description of the data. (Use User\_Data.csv)

```

In [40]: 1 import pandas as pd
2
3 # Load the dataset from the CSV file
4 df = pd.read_csv('user_data.csv')
5
6 # Print the shape (number of rows and columns)
7 print("Shape of the data:", df.shape)
8
9 # Print the number of rows and columns
10 print("Number of rows:", df.shape[0])
11 print("Number of columns:", df.shape[1])
12
13 # Print data types of each column
14 print("\nData types:")
15 print(df.dtypes)
16
17 # Print feature names
18 print("\nFeature names:")
19 print(df.columns)
20
21 # Print the description of the data
22 print("\nDescription of the data:")
23 print(df.describe())
24

```

Shape of the data: (5, 5)

Number of rows: 5

Number of columns: 5

Data types:

Name object

Age int64

Gender object

Height int64

Weight int64

dtype: object

Feature names:

Index(['Name', 'Age', 'Gender', 'Height', 'Weight'], dtype='object')

Description of the data:

	Age	Height	Weight
count	5.000000	5.000000	5.000000
mean	29.600000	174.000000	71.000000
std	4.159327	9.192388	14.317821
min	24.000000	162.000000	55.000000
25%	28.000000	168.000000	60.000000
50%	29.000000	175.000000	70.000000
75%	32.000000	180.000000	80.000000
max	35.000000	185.000000	90.000000

SLIP 5 same as SLIP 4

SLIP 6 same as SLIP 2

SLIP 7

Write a Python program to perform the following tasks :

a. Apply OneHot coding on Country column.

b. Apply Label encoding on purchased column

(Data.csv have two categorical column the country column, and the purchased column)



```
In [41]: 1 import pandas as pd
2 from sklearn.preprocessing import LabelEncoder, OneHotEncoder
3
4 # Load the dataset from the CSV file
5 df = pd.read_csv('data2.csv')
6
7 # Display the original dataset
8 print("Original Dataset:")
9 print(df)
10
11 # a. Apply OneHot encoding on the 'Country' column
12 df_onehot = pd.get_dummies(df, columns=['Country'], prefix='Country')
13
14 # Display the dataset after OneHot encoding
15 print("\nDataset after OneHot encoding:")
16 print(df_onehot)
17
18 # b. Apply Label encoding on the 'Purchased' column
19 label_encoder = LabelEncoder()
20 df['Purchased'] = label_encoder.fit_transform(df['Purchased'])
21
22 # Display the dataset after Label encoding
23 print("\nDataset after Label encoding:")
24 print(df)
25
```

Original Dataset:

	Country	Age	Salary	Purchased
0	France	25	50000	No
1	Spain	30	60000	Yes
2	Germany	35	70000	No
3	Spain	28	62000	No
4	Germany	40	80000	Yes
5	France	32	75000	Yes
6	Spain	45	90000	No
7	France	50	95000	Yes
8	Germany	22	55000	No
9	France	48	98000	Yes

Dataset after OneHot encoding:

	Age	Salary	Purchased	Country_France	Country_Germany	Country_Spain
0	25	50000	No	True	False	False
1	30	60000	Yes	False	False	True
2	35	70000	No	False	True	False
3	28	62000	No	False	False	True
4	40	80000	Yes	False	True	False
5	32	75000	Yes	True	False	False
6	45	90000	No	False	False	True
7	50	95000	Yes	True	False	False
8	22	55000	No	False	True	False
9	48	98000	Yes	True	False	False

Dataset after Label encoding:

	Country	Age	Salary	Purchased
0	France	25	50000	0
1	Spain	30	60000	1
2	Germany	35	70000	0
3	Spain	28	62000	0
4	Germany	40	80000	1
5	France	32	75000	1
6	Spain	45	90000	0
7	France	50	95000	1
8	Germany	22	55000	0
9	France	48	98000	1

## SLIP 8

Q.2) Write a program in python to perform following task Standardizing Data (transform them into a standard Gaussian distribution with a mean of 0 and a standard deviation of 1) (Use winequality-red.csv)

```

In [42]: 1 import pandas as pd
2 from sklearn.preprocessing import StandardScaler
3
4 # Load the dataset from the CSV file
5 df = pd.read_csv('winequality-red.csv')
6
7 # Display the original dataset
8 print("Original Dataset:")
9 print(df.head())
10
11 # Extract the features (excluding the target variable)
12 features = df.drop('quality', axis=1)
13
14 # Initialize the StandardScaler
15 scaler = StandardScaler()
16
17 # Fit and transform the features using StandardScaler
18 features_standardized = scaler.fit_transform(features)
19
20 # Create a new DataFrame with standardized features
21 df_standardized = pd.DataFrame(features_standardized, columns=features.columns)
22
23 # Add the 'quality' column back to the DataFrame
24 df_standardized['quality'] = df['quality']
25
26 # Display the dataset after standardization
27 print("\nDataset after Standardization:")
28 print(df_standardized.head())
29

```

Original Dataset:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
0	7.4	0.70	0.00	1.9	0.076	
1	7.8	0.88	0.00	2.6	0.098	
2	7.8	0.76	0.04	2.3	0.092	
3	11.2	0.28	0.56	1.9	0.075	
4	7.4	0.70	0.00	1.9	0.076	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
0	11.0	34.0	0.9978	3.51	0.56	
1	25.0	67.0	0.9968	3.20	0.68	
2	15.0	54.0	0.9970	3.26	0.65	
3	17.0	60.0	0.9980	3.16	0.58	
4	11.0	34.0	0.9978	3.51	0.56	

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5
3	9.8	6
4	9.4	5

Dataset after Standardization:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
0	-0.498662	0.451753	-0.563489	-0.329398	-0.103362	
1	-0.135999	1.630239	-0.563489	0.206831	2.170608	
2	-0.135999	0.844582	-0.346763	-0.022981	1.550434	
3	2.946642	-2.298048	2.470683	-0.329398	-0.206725	
4	-0.498662	0.451753	-0.563489	-0.329398	-0.103362	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
0	-0.896665	-0.626576	0.731200	1.276466	-0.305196	
1	2.406839	0.761143	-0.284356	-1.276466	0.957683	
2	0.047193	0.214466	-0.081244	-0.782350	0.641963	
3	0.519122	0.466778	0.934311	-1.605877	-0.094716	
4	-0.896665	-0.626576	0.731200	1.276466	-0.305196	

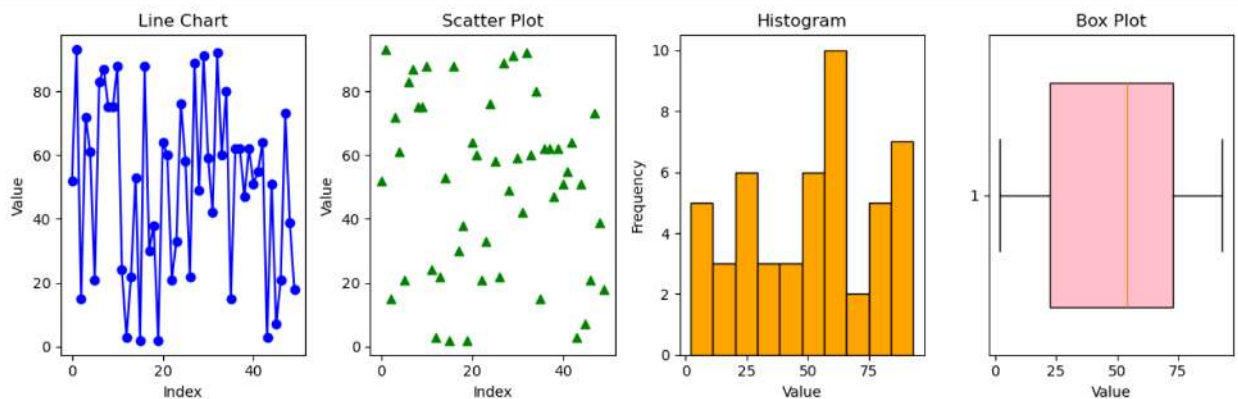
	alcohol	quality
0	-0.88083	5
1	0.29361	5
2	0.29361	5
3	0.29361	6
4	-0.88083	5

Q.2 A) Generate a random array of 50 integers and display them using a line chart, scatter plot. Apply appropriate color, labels and styling options.

```

In [38]: 1 import numpy as np
2 import matplotlib.pyplot as plt
3
4 # Set a seed for reproducibility
5 np.random.seed(42)
6
7 # Generate a random array of 50 integers
8 random_array = np.random.randint(1, 100, 50)
9
10 # Line Chart
11 plt.figure(figsize=(12, 4))
12 plt.subplot(1, 4, 1)
13 plt.plot(random_array, marker='o', color='blue')
14 plt.title('Line Chart')
15 plt.xlabel('Index')
16 plt.ylabel('Value')
17
18 # Scatter Plot
19 plt.subplot(1, 4, 2)
20 plt.scatter(range(len(random_array)), random_array, color='green', marker='^')
21 plt.title('Scatter Plot')
22 plt.xlabel('Index')
23 plt.ylabel('Value')
24
25 # Histogram
26 plt.subplot(1, 4, 3)
27 plt.hist(random_array, bins=10, color='orange', edgecolor='black')
28 plt.title('Histogram')
29 plt.xlabel('Value')
30 plt.ylabel('Frequency')
31
32 # Box Plot
33 plt.subplot(1, 4, 4)
34 plt.boxplot(random_array, vert=False, widths=0.7, patch_artist=True, boxprops=dict(facecolor='pink'))
35 plt.title('Box Plot')
36 plt.xlabel('Value')
37
38 # Adjust layout for better presentation
39 plt.tight_layout()
40
41 # Show the plots
42 plt.show()
43

```

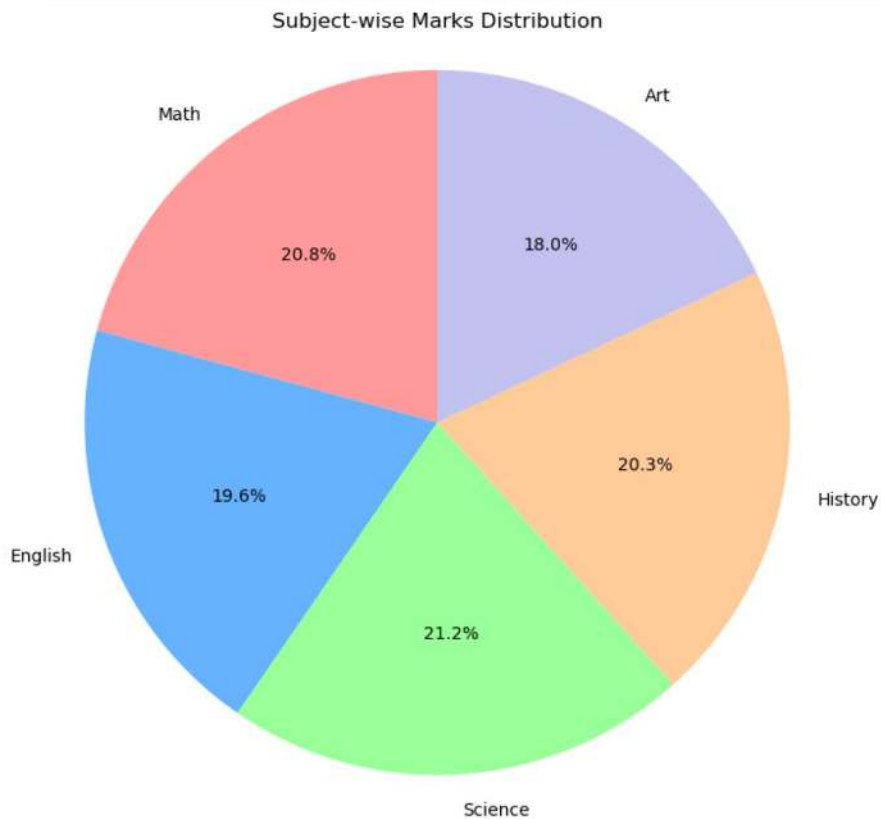


Q.2 B) Create two lists, one representing subject names and the other representing marks obtained in those subjects. Display the data in a pie chart.

```

In [1]: 1 import matplotlib.pyplot as plt
2
3 # Sample data - replace with your own data
4 subjects = ['Math', 'English', 'Science', 'History', 'Art']
5 marks = [90, 85, 92, 88, 78]
6
7 # Create a pie chart
8 plt.figure(figsize=(8, 8))
9 plt.pie(marks, labels=subjects, autopct='%1.1f%%', startangle=90, colors=['#FF9999', '#66B2FF', '#99FF99', '#FFCC99', '#c2c2ff'])
10 plt.title('Subject-wise Marks Distribution')
11 plt.axis('equal') # Equal aspect ratio ensures that pie is drawn as a circle.
12
13 # Show the plot
14 plt.show()
15

```



Q.2 C) Write a program in python to perform following task (Use winequality-red.csv ) Import Dataset and do the followings:

- Describing the dataset
- Shape of the dataset
- Display first 3 rows from dataset

```

In [2]: 1 import pandas as pd
2
3 # a) Describing the dataset
4 # b) Shape of the dataset
5 # Load the dataset from the CSV file
6 df = pd.read_csv('winequality-red.csv')
7
8 # Display basic statistical details of the data
9 print("a) Describing the dataset:")
10 print(df.describe())
11
12 # Display the shape of the dataset
13 print("\nb) Shape of the dataset:")
14 print(df.shape)
15
16 # c) Display first 3 rows from the dataset
17 print("\nc) Display first 3 rows from the dataset:")
18 print(df.head(3))
19

```

a) Describing the dataset:

	fixed acidity	volatile acidity	citric acid	residual sugar	\
count	10.000000	10.000	10.000000	10.000000	
mean	7.950000	0.631	0.104000	2.330000	
std	1.162612	0.161	0.194548	1.376025	
min	7.300000	0.280	0.000000	1.200000	
25%	7.400000	0.585	0.000000	1.825000	
50%	7.650000	0.655	0.010000	1.900000	
75%	7.800000	0.700	0.055000	2.225000	
max	11.200000	0.880	0.560000	6.100000	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	\
count	10.000000	10.000000	10.000000	10.000000	
mean	0.077000	14.800000	48.900000	0.997080	
std	0.010198	4.467164	25.066356	0.001038	
min	0.065000	9.000000	18.000000	0.994600	
25%	0.071500	11.500000	34.000000	0.996800	
50%	0.075000	15.000000	47.000000	0.997400	
75%	0.076000	16.500000	59.750000	0.997800	
max	0.098000	25.000000	102.000000	0.998000	

	pH	sulphates	alcohol	quality
count	10.000000	10.000000	10.000000	10.000000
mean	3.355000	0.589000	9.700000	5.500000
std	0.127997	0.100161	0.359011	0.849837
min	3.160000	0.460000	9.400000	5.000000
25%	3.270000	0.560000	9.400000	5.000000
50%	3.355000	0.565000	9.650000	5.000000
75%	3.480000	0.632500	9.800000	5.750000
max	3.510000	0.800000	10.500000	7.000000

b) Shape of the dataset:  
(10, 12)

c) Display first 3 rows from the dataset:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
0	7.4	0.70	0.00	1.9	0.076	
1	7.8	0.88	0.00	2.6	0.098	
2	7.8	0.76	0.04	2.3	0.092	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
0	11.0	34.0	0.9978	3.51	0.56	
1	25.0	67.0	0.9968	3.20	0.68	
2	15.0	54.0	0.9970	3.26	0.65	

	alcohol	quality
0	9.4	5
1	9.8	5
2	9.8	5

## SLIP 10

Q.2 A) Write a python program to Display column-wise mean, and median for SOCR- HeightWeight dataset.

```
In [3]: 1 import pandas as pd
2
3 # Assuming 'SOCR-HeightWeight.csv' is the dataset file
4 # Make sure the file is in the same directory as your Python script or provide the correct path
5 df = pd.read_csv('height_weight.csv')
6
7 # Display column-wise mean
8 print("Column-wise Mean:")
9 mean_values = df.mean()
10 print(mean_values)
11
12 # Display column-wise median
13 print("\nColumn-wise Median:")
14 median_values = df.median()
15 print(median_values)
16
```

Column-wise Mean:  
Height 172.0  
Weight 66.9  
dtype: float64

Column-wise Median:  
Height 171.0  
Weight 66.5  
dtype: float64

Q.2 B) Write a python program to compute sum of Manhattan distance between all pairs of points.

```
In [4]: 1 import itertools
2
3 def manhattan_distance(point1, point2):
4     return abs(point1[0] - point2[0]) + abs(point1[1] - point2[1])
5
6 # Sample dataset (replace with your own dataset)
7 points = [(1, 2), (3, 4), (5, 6), (7, 8)]
8
9 # Compute the sum of Manhattan distance between all pairs of points
10 total_distance = sum(manhattan_distance(point1, point2) for point1, point2 in itertools.combinations(points, 2))
11
12 # Display the result
13 print("Sum of Manhattan distance between all pairs of points:", total_distance)
14
```

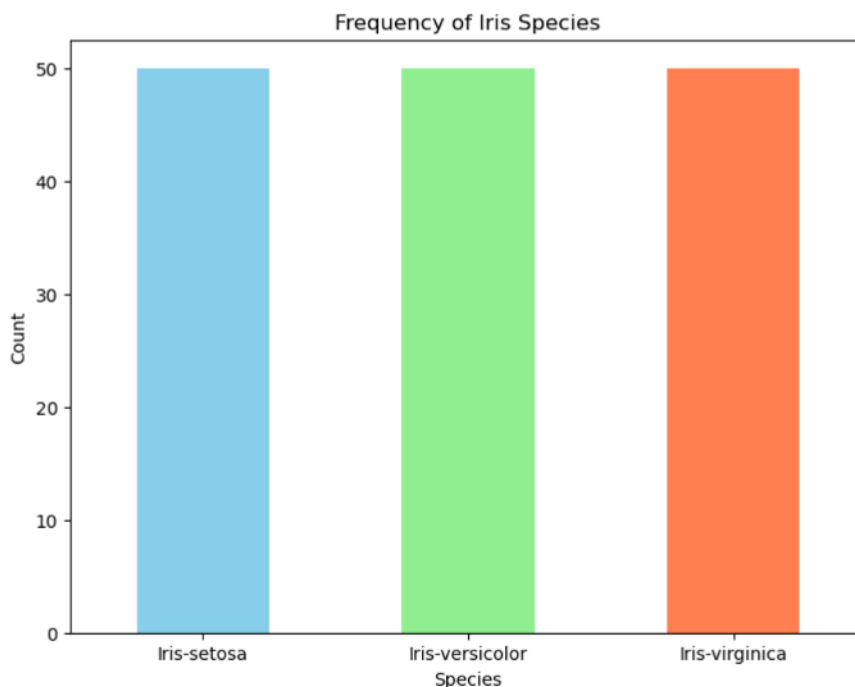
Sum of Manhattan distance between all pairs of points: 40

SLIP 11 same as SLIP 1

SLIP 18 same as SLIP 3

Q.2 A) Import dataset “iris.csv”. Write a Python program to create a Bar plot to get the frequency of the three species of the Iris data.

```
In [2]: 1 import pandas as pd
2 import matplotlib.pyplot as plt
3
4 # Load the Iris dataset
5 iris_data = pd.read_csv('iris.csv')
6
7 # Count the frequency of each species
8 species_counts = iris_data['Species'].value_counts()
9
10 # Create a bar plot
11 plt.figure(figsize=(8, 6))
12 species_counts.plot(kind='bar', color=['skyblue', 'lightgreen', 'coral'])
13 plt.title('Frequency of Iris Species')
14 plt.xlabel('Species')
15 plt.ylabel('Count')
16 plt.xticks(rotation=0) # Rotate x-axis labels for better readability
17
18 # Show the plot
19 plt.show()
20
```



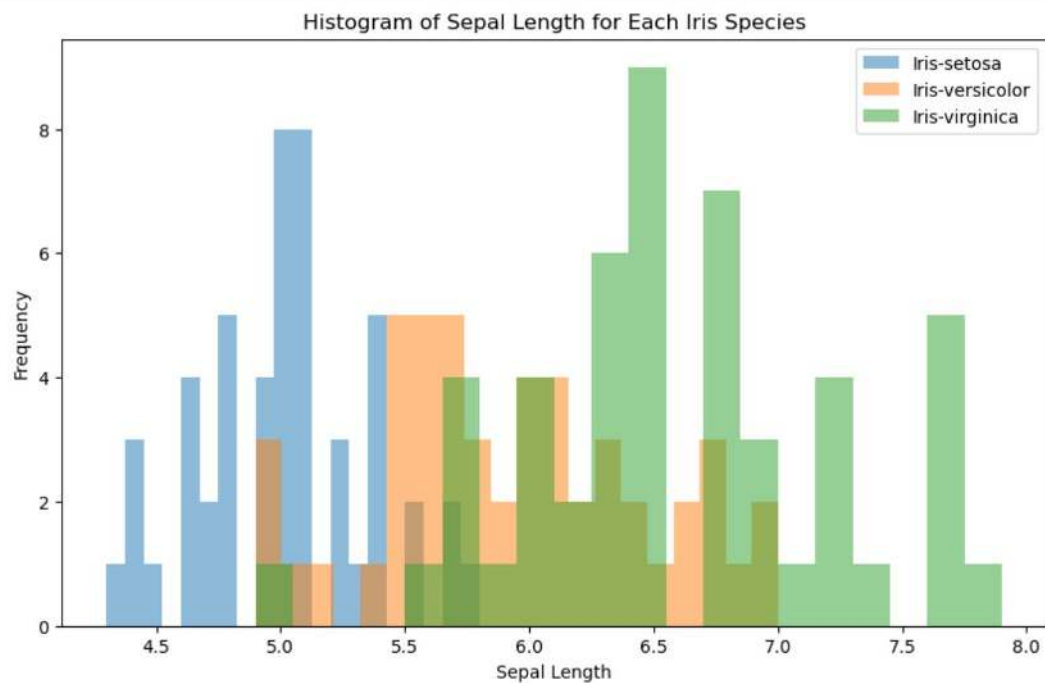
Q.2 B) Write a Python program to create a histogram of the three species of the Iris data.



```

In [6]: 1 import pandas as pd
2 import matplotlib.pyplot as plt
3
4 # Load the Iris dataset
5 iris_data = pd.read_csv('iris.csv')
6
7 # Create a histogram for each species
8 plt.figure(figsize=(10, 6))
9
10 # Iterate through each species and create a histogram
11 for species in iris_data['Species'].unique():
12     subset = iris_data[iris_data['Species'] == species]
13     plt.hist(subset['SepalLengthCm'], bins=20, alpha=0.5, label=species)
14
15 # Add Labels and title
16 plt.title('Histogram of Sepal Length for Each Iris Species')
17 plt.xlabel('Sepal Length')
18 plt.ylabel('Frequency')
19 plt.legend()
20
21 # Show the plot
22 plt.show()
23

```



SLIP 24 same as SLIP 21