

This is the Iris dataset. The dataset 150 data points.

It includes three iris species with 50 samples each as well as some properties about each flower. One flower species is linearly separable from the other two, but the other two are not linearly separable from each other.

The columns in this dataset are:

Id  
SepalLengthCm  
SepalWidthCm  
PetalLengthCm  
PetalWidthCm  
Species

1.Firstly, treat all outliers and missing values in the dataset.

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns
# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df) * 100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
```

```

'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25) # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75) # Gives 75th
Percentile or Q3

IQR = Q3 - Q1

UpperBound = Q3 + 1.5 * IQR
LowerBound = Q1 - 1.5 * IQR

OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
print(len(OutlierData))

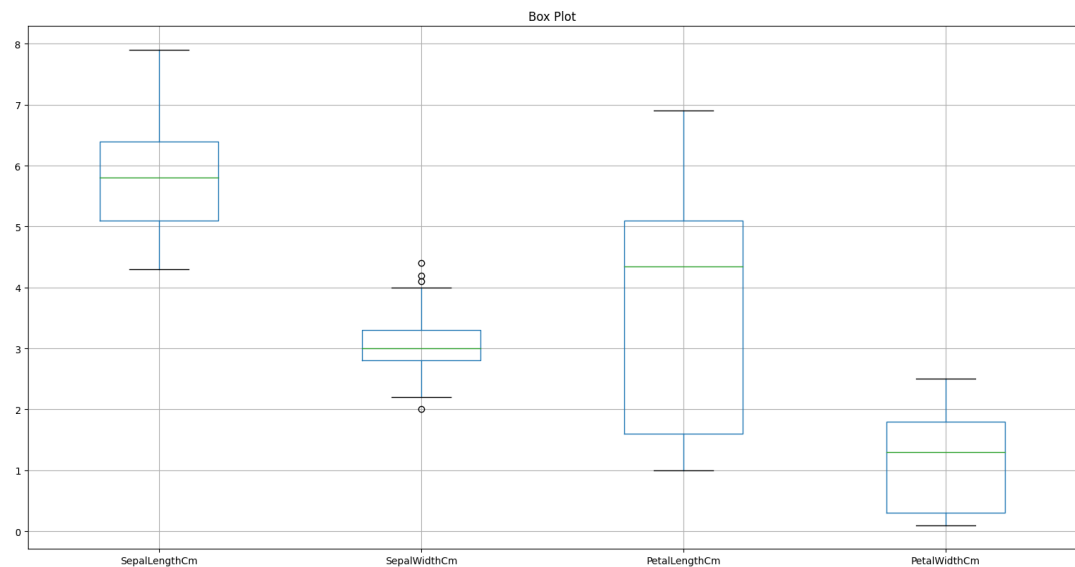
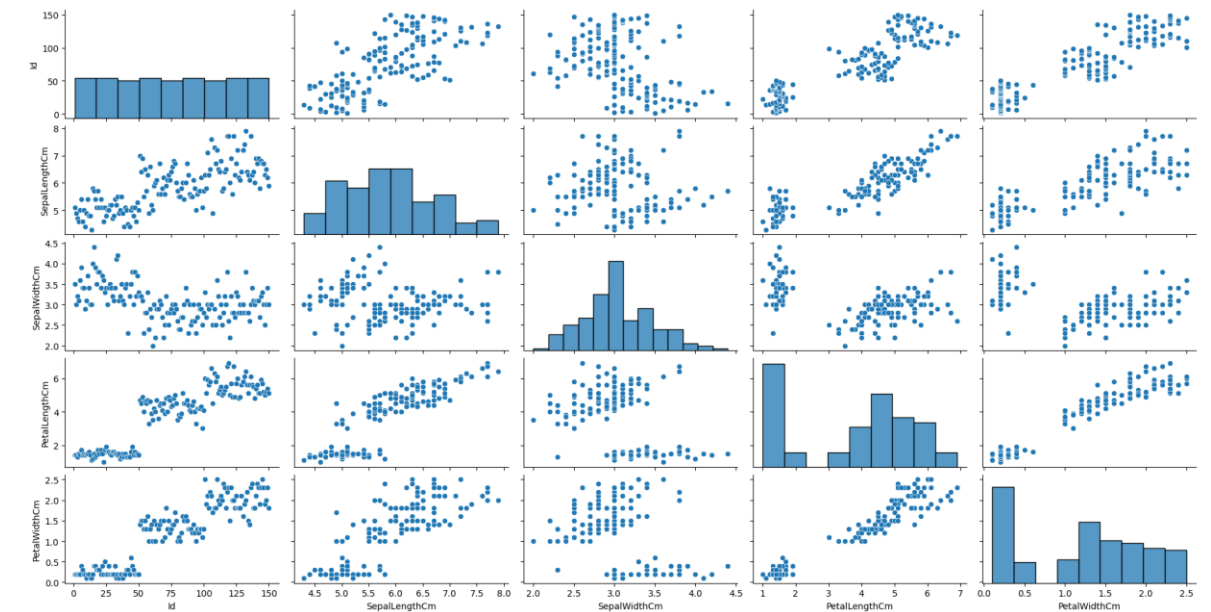
```

File - Day13Q1

```

1 C:\Users\tejas\PycharmProjects\pythonProject\venv\Scripts\python.exe C:\Users\
  tejas\PycharmProjects\pythonProject\START\Day13Q1.py
2 <class 'pandas.core.frame.DataFrame'>
3 RangeIndex: 150 entries, 0 to 149
4 Data columns (total 6 columns):
5 #   Column          Non-Null Count  Dtype
6 ---  ---
7 0    Id              150 non-null   int64
8 1    SepalLengthCm   150 non-null   float64
9 2    SepalWidthCm    150 non-null   float64
10 3    PetalLengthCm   150 non-null   float64
11 4    PetalWidthCm    150 non-null   float64
12 5    Species         150 non-null   object
13 dtypes: float64(4), int64(1), object(1)
14 memory usage: 7.2+ KB
15 Id              0.0
16 SepalLengthCm   0.0
17 SepalWidthCm    0.0
18 PetalLengthCm   0.0
19 PetalWidthCm    0.0
20 Species         0.0
21 dtype: float64
22 0
23 4
24 4
25 4
26
27 Process finished with exit code 0
28

```



2. Complete all basic data descriptive statistics by *Species*

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns

# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
```

```

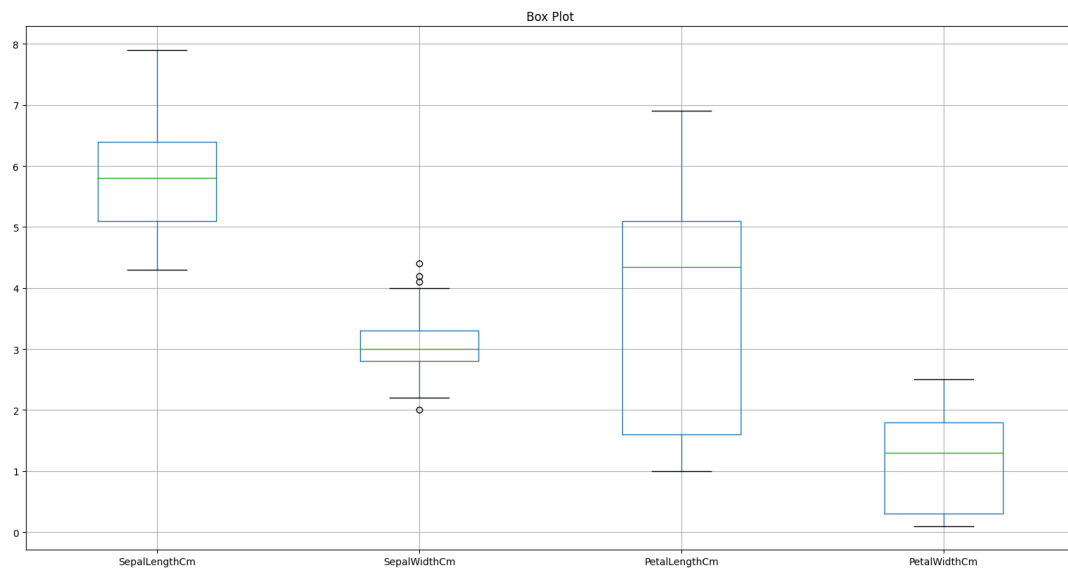
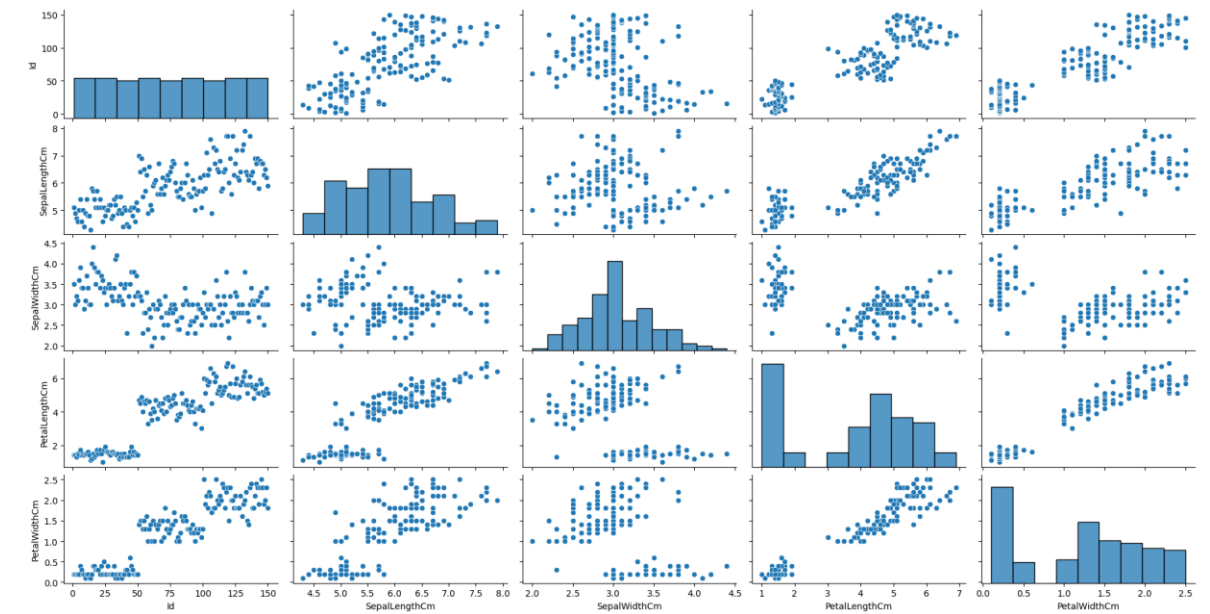
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df)
* 100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm',
'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25) # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75) # Gives 75th
Percentile or Q3

    IQR = Q3 - Q1

    UpperBound = Q3 + 1.5 * IQR
    LowerBound = Q1 - 1.5 * IQR

    OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
    print(len(OutlierData))
# group the dataset based on the "Species" column
descriptive_stats =
df.groupby('Species').describe()
print(descriptive_stats)

```



File - Day13Q2

```
1 C:\Users\tejas\PycharmProjects\pythonProject\venv\Scripts\python.exe C:\Users\
  tejas\PycharmProjects\pythonProject\START\Day13Q2.py
2 <class 'pandas.core.frame.DataFrame'>
3 RangeIndex: 150 entries, 0 to 149
4 Data columns (total 6 columns):
5 #   Column          Non-Null Count  Dtype
6 ---  ---
7 0   Id               150 non-null    int64
8 1   SepalLengthCm    150 non-null    float64
9 2   SepalWidthCm     150 non-null    float64
10 3   PetalLengthCm    150 non-null    float64
11 4   PetalWidthCm     150 non-null    float64
12 5   Species          150 non-null    object
13 dtypes: float64(4), int64(1), object(1)
14 memory usage: 7.2+ KB
15 Id               0.0
16 SepalLengthCm    0.0
17 SepalWidthCm     0.0
18 PetalLengthCm    0.0
19 PetalWidthCm     0.0
20 Species          0.0
21 dtype: float64
22 0
23 4
24 4
25 4
26
27
28
29
30
31
32
33 [3 rows x 40 columns]
34
35 Process finished with exit code 0
36
```

	Id	...	PetalWidthCm						
	count	mean	std	min	25%	...	min	25%	
50%	75%	max							
Species						...			
Iris-setosa	50.0	25.5	14.57738	1.0	13.25	...	0.1	0.2	0
Iris-versicolor	50.0	75.5	14.57738	51.0	63.25	...	1.0	1.2	1
Iris-virginica	50.0	125.5	14.57738	101.0	113.25	...	1.4	1.8	2

3. Use the *Sepal Length*, *Sepal Width*, *Petal Length* and *Petal Width* to find K-Means clusters.

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns
from sklearn.cluster import KMeans
```

```

# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df) *
100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25) # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75) # Gives 75th
Percentile or Q3

    IQR = Q3 - Q1

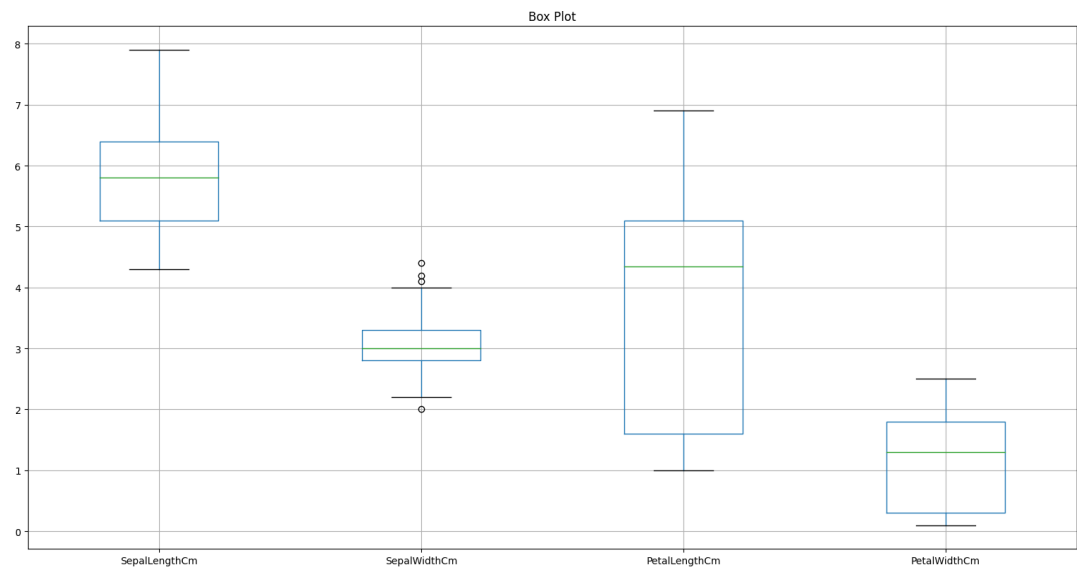
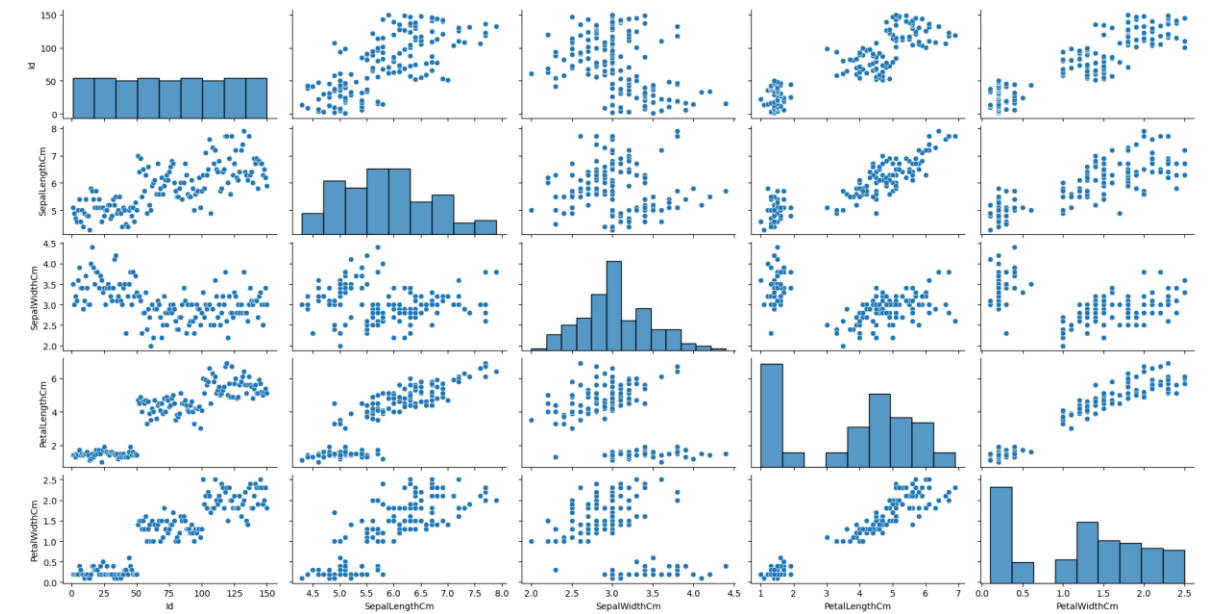
    UpperBound = Q3 + 1.5 * IQR
    LowerBound = Q1 - 1.5 * IQR

    OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
    print(len(OutlierData))
# group the dataset based on the "Species" column
descriptive_stats =
df.groupby('Species').describe()
print(descriptive_stats)
X = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
# specify the number of clusters
k = 3
# create a kmeans instance
km = KMeans(n_clusters=k, n_init=25,

```

```
random_state=1234)
# fit the data to the kmeans model
km.fit(X)
# get the cluster labels for each data point
cluster_labels = km.labels_
# the total within cluster sum of squares
clusterWCSS = km.inertia_
print(cluster_labels)
print(clusterWCSS)
```





[illegible]

Optimal Cluster number is 3 as we get the same score in elbow method and Calinski Harabasz Score.

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns
```

```

from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score
from sklearn.metrics import calinski_harabasz_score

# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df) *
100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25) # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75) # Gives 75th
Percentile or Q3

    IQR = Q3 - Q1

    UpperBound = Q3 + 1.5 * IQR
    LowerBound = Q1 - 1.5 * IQR

    OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
    print(len(OutlierData))
# group the dataset based on the "Species" column
descriptive_stats =
df.groupby('Species').describe()
print(descriptive_stats)
X = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]

```

```

# specify the number of clusters
k = 3
# create a kmeans instance
km = KMeans(n_clusters=k, n_init=25,
random_state=1234)
# fit the data to the kmeans model
km.fit(X)
# get the cluster labels for each data point
cluster_labels = km.labels_
# the total within cluster sum of squares
clusterWCSS = km.inertia_
print(cluster_labels)
print(clusterWCSS)
# tabulation of the size of the clusters
pd.Series(km.labels_).value_counts().sort_index()
# 'km.cluster_centers_' :gives the cluster
centroids
cluster_centers = pd.DataFrame(km.cluster_centers_,

columns=['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm'])
print(cluster_centers)
# to get the correct centroids, we need to un-scale
the data,
cluster_centers_unscaled = pd.DataFrame()
for i in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    cluster_centers_unscaled[i] =
(cluster_centers[i] * df[i].std()) + df[i].mean()
print(cluster_centers_unscaled)
wcss = []
for k in range(2, 11):
    km = KMeans(n_clusters=k, n_init=25,
random_state=1234)
    km.fit(X)
    wcss.append(km.inertia_)

plt.plot(range(2, 11), wcss)
plt.xlabel('Number of Clusters')
plt.ylabel('WCSS')
plt.title('Elbow Method')
plt.show()

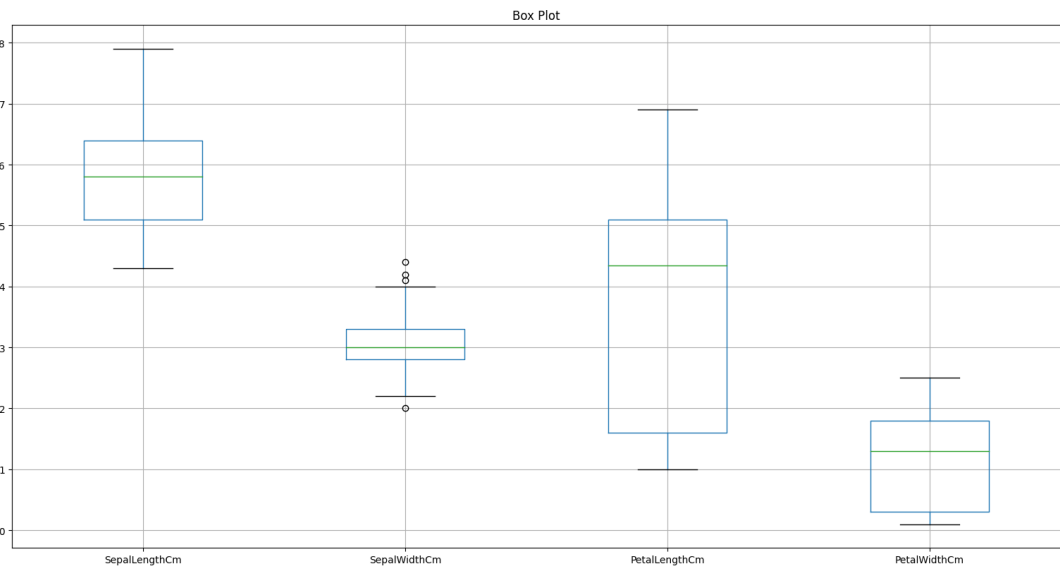
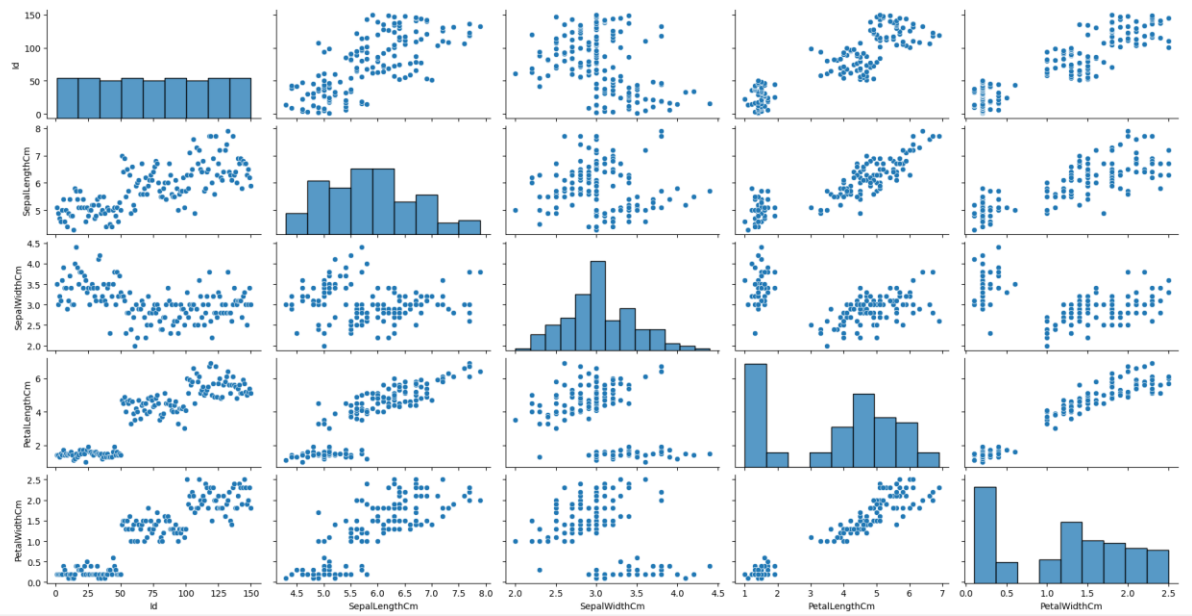
```

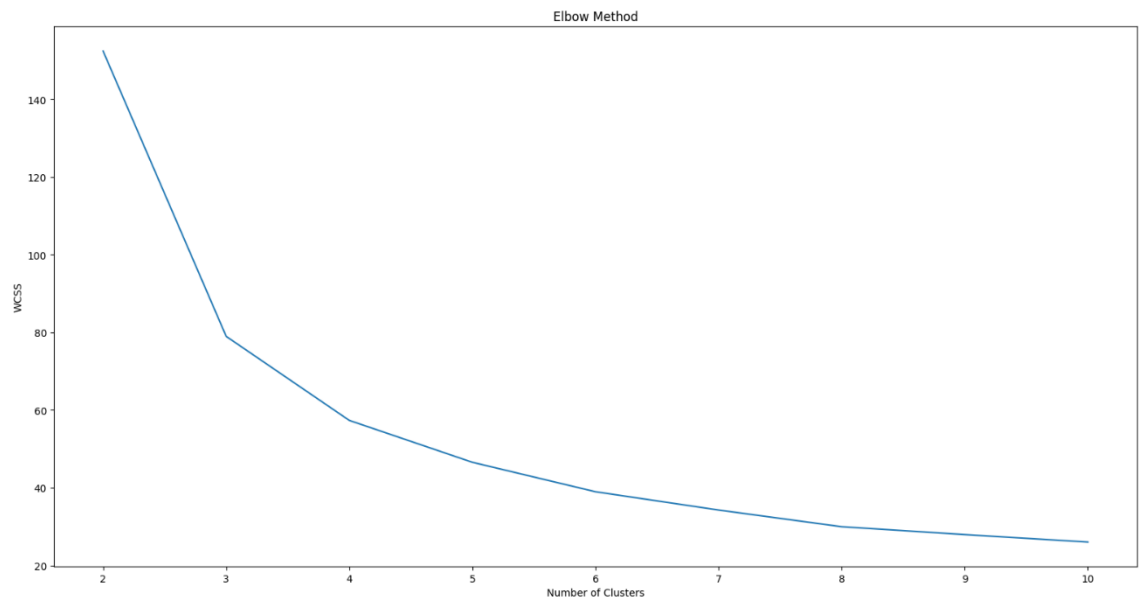
```
silhouette = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
random_state = 1234)
    km.fit(X)
    silhouette.append(silhouette_score(X,
km.labels_))

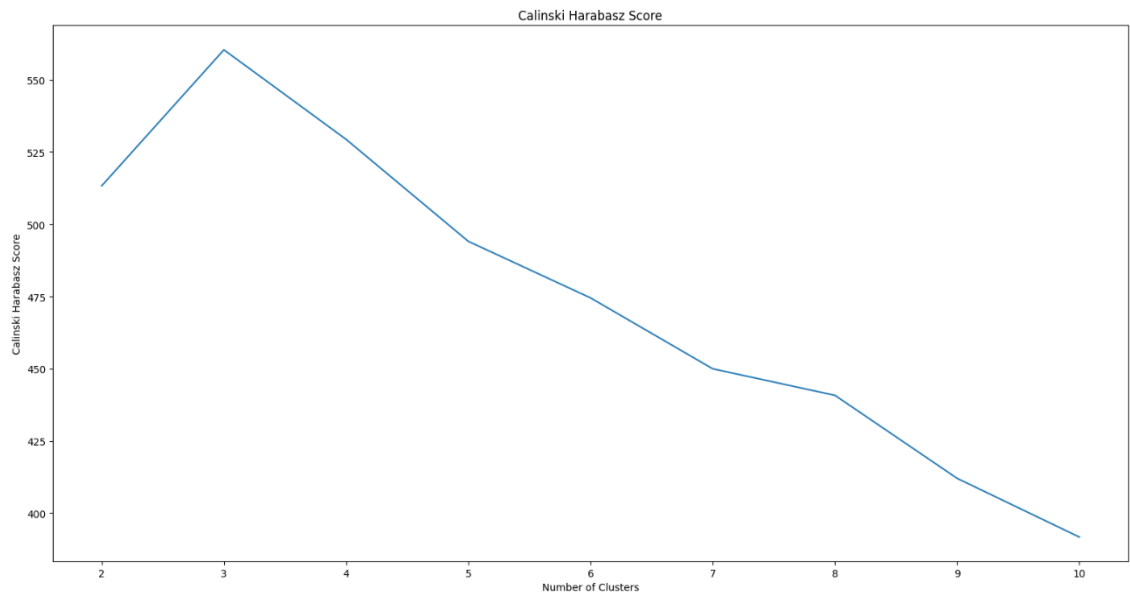
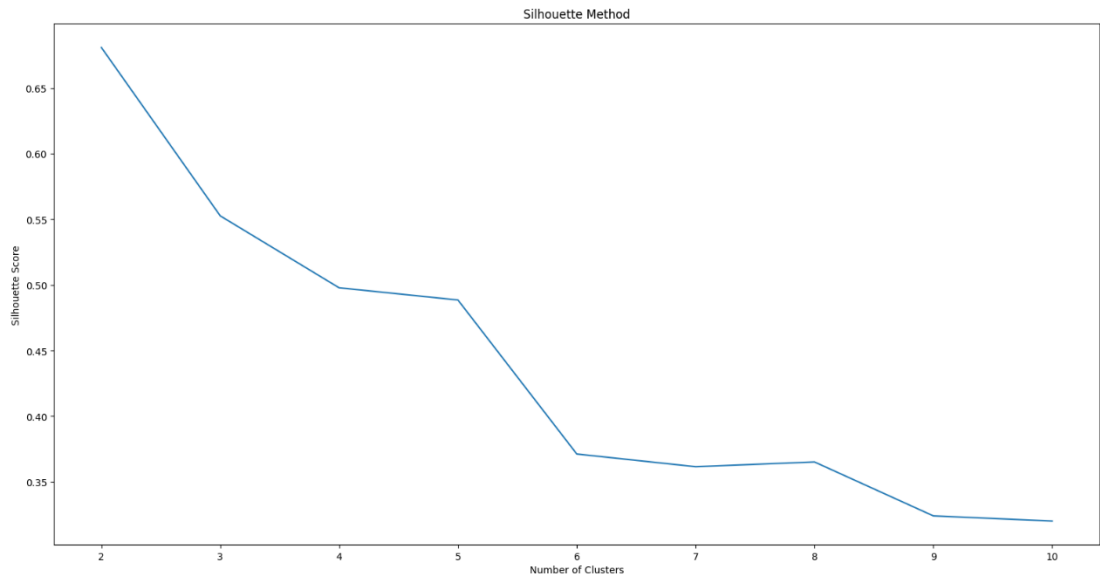
plt.plot(range(2, 11), silhouette)
plt.xlabel('Number of Clusters')
plt.ylabel('Silhouette Score')
plt.title('Silhouette Method')
plt.show()

calinski = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
random_state = 1234)
    km.fit(X)
    calinski.append(calinski_harabasz_score(X,
km.labels_))

plt.plot(range(2, 11), calinski)
plt.xlabel('Number of Clusters')
plt.ylabel('Calinski Harabasz Score')
plt.title('Calinski Harabasz Score')
plt.show()
```











```

import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score
from sklearn.metrics import calinski_harabasz_score

# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df) *
100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25) # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75) # Gives 75th
Percentile or Q3

    IQR = Q3 - Q1

    UpperBound = Q3 + 1.5 * IQR
    LowerBound = Q1 - 1.5 * IQR

    OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
    print(len(OutlierData))
# group the dataset based on the "Species" column
descriptive_stats =

```

```

df.groupby('Species').describe()
print(descriptive_stats)
X = df[['SepalLengthCm', 'SepalWidthCm',
        'PetalLengthCm', 'PetalWidthCm']]
# specify the number of clusters
k = 3
# create a kmeans instance
km = KMeans(n_clusters=k, n_init=25,
            random_state=1234)
# fit the data to the kmeans model
km.fit(X)
# get the cluster labels for each data point
cluster_labels = km.labels_
# the total within cluster sum of squares
clusterWCSS = km.inertia_
print(cluster_labels)
print(clusterWCSS)
# tabulation of the size of the clusters
pd.Series(km.labels_).value_counts().sort_index()
# 'km.cluster_centers_' :gives the cluster
centroids
cluster_centers = pd.DataFrame(km.cluster_centers_,

columns=['SepalLengthCm', 'SepalWidthCm',
        'PetalLengthCm', 'PetalWidthCm'])
print(cluster_centers)
# to get the correct centroids, we need to un-scale
the data,
cluster_centers_unscaled = pd.DataFrame()
for i in ['SepalLengthCm', 'SepalWidthCm',
        'PetalLengthCm', 'PetalWidthCm']:
    cluster_centers_unscaled[i] =
    (cluster_centers[i] * df[i].std()) + df[i].mean()
print(cluster_centers_unscaled)
wcss = []
for k in range(2, 11):
    km = KMeans(n_clusters=k, n_init=25,
                random_state=1234)
    km.fit(X)
    wcss.append(km.inertia_)

plt.plot(range(2, 11), wcss)

```

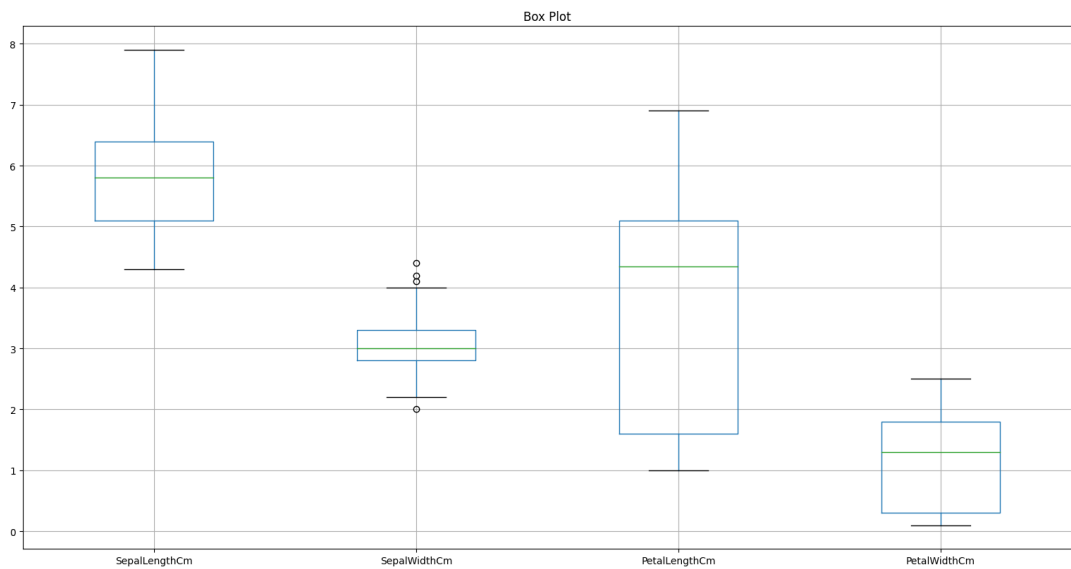
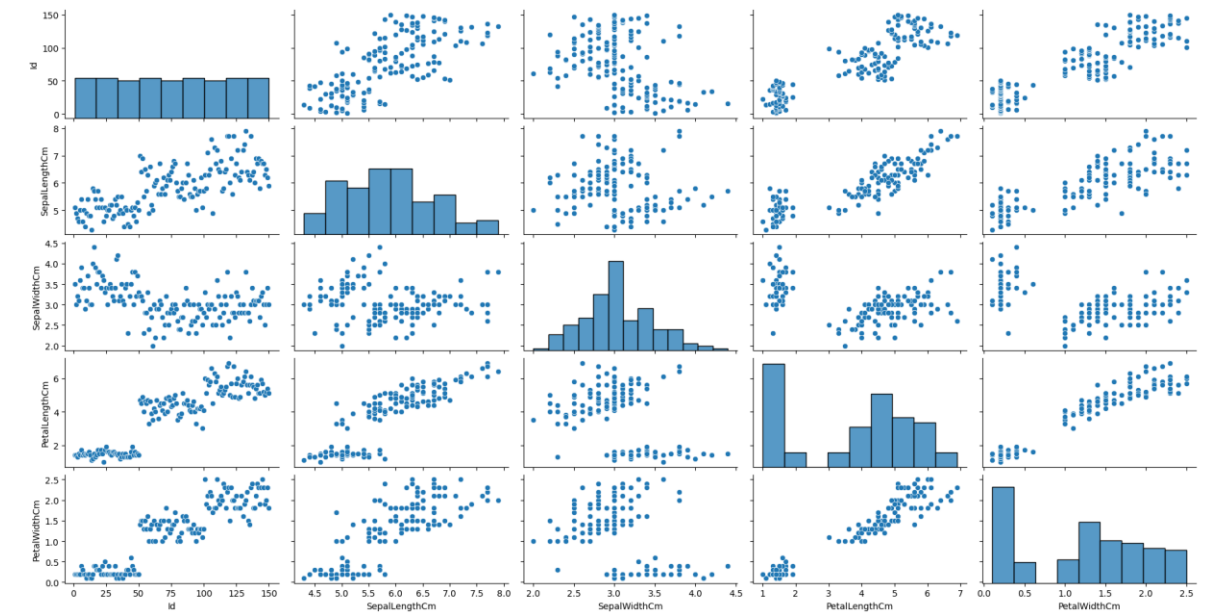
```

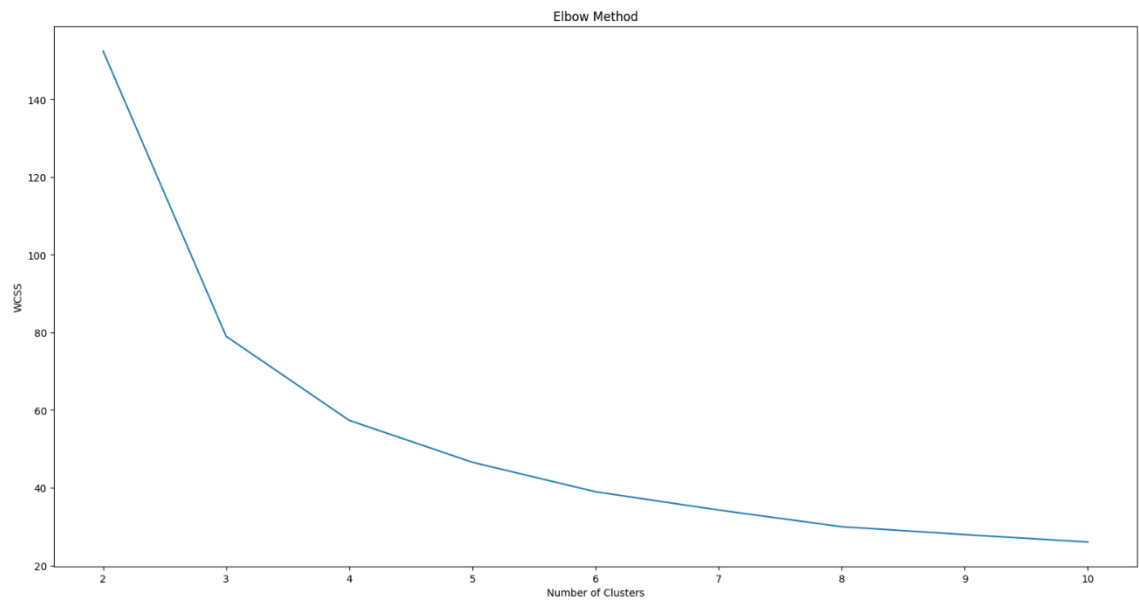
plt.xlabel('Number of Clusters')
plt.ylabel('WCSS')
plt.title('Elbow Method')
plt.show()
silhouette = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
random_state = 1234)
    km.fit(X)
    silhouette.append(silhouette_score(X,
km.labels_))

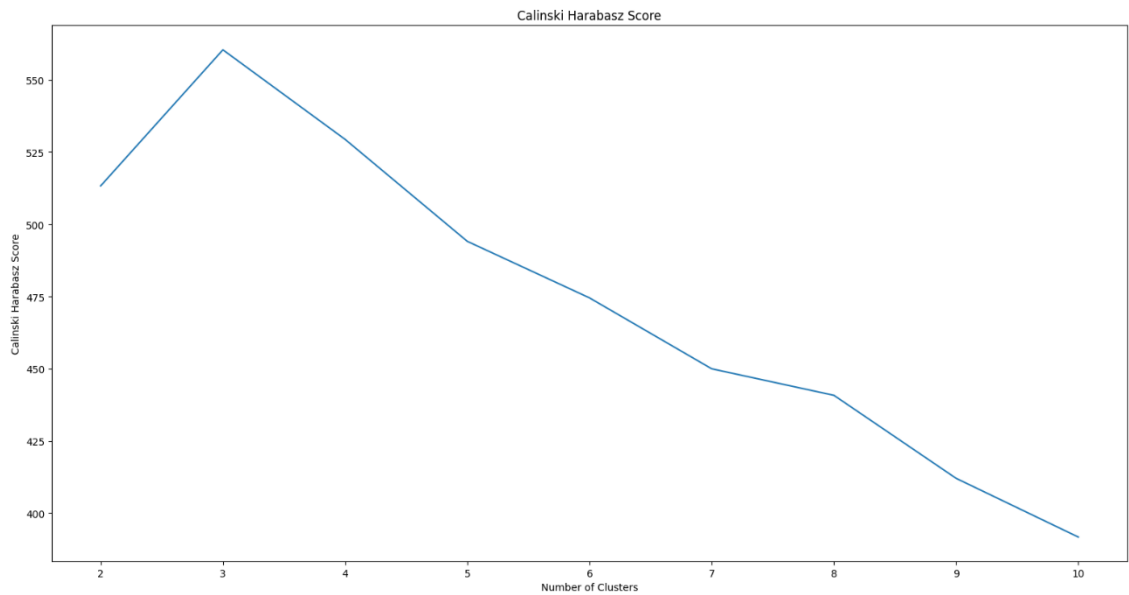
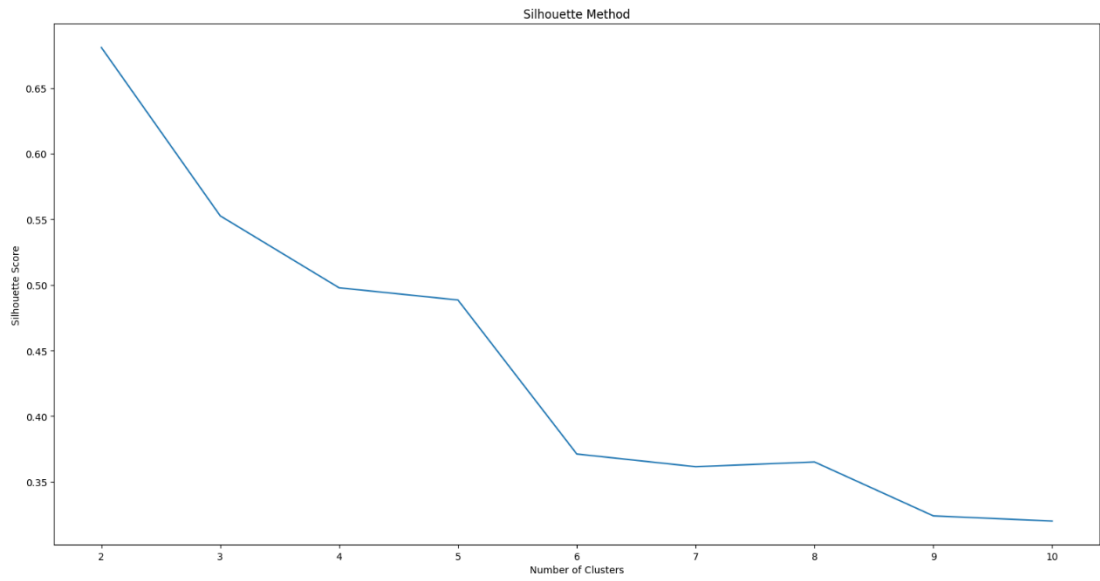
plt.plot(range(2, 11), silhouette)
plt.xlabel('Number of Clusters')
plt.ylabel('Silhouette Score')
plt.title('Silhouette Method')
plt.show()
calinski = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
random_state = 1234)
    km.fit(X)
    calinski.append(calinski_harabasz_score(X,
km.labels_))

plt.plot(range(2, 11), calinski)
plt.xlabel('Number of Clusters')
plt.ylabel('Calinski Harabasz Score')
plt.title('Calinski Harabasz Score')
plt.show()
cluster_df = pd.DataFrame({'Cluster':
cluster_labels, 'Species': df['Species']})
cross_tab = pd.crosstab(cluster_df['Cluster'],
cluster_df['Species'])
proportion = cross_tab.div(cross_tab.sum(axis=1),
axis=0) * 100
print(proportion)

```







[illegible]



File - Day13Q5

```
52 2          0.0      77.419355    22.580645
53
54 Process finished with exit code 0
55
```

6. Share your insights on the data based on the clusters [optional]

Petal width and petal length are better differentiators than sepal width and sepal length. In the petal width-petal length plot we can see that setosa has smallest petal width and petal length, versicolor lies between setosa and virginica, virginica has the largest petal width and petal length. But from the sepal width-sepal length plot we don't get much idea about the species differentiation, albeit we can observe that sepal length gives us the idea that versicolor and virginica are somehow related or have shared qualities, sepal width doesn't give us any idea AT ALL. So it's not a good differentiator

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
import seaborn as sns
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score
from sklearn.metrics import calinski_harabasz_score

# Load the dataset
df = pd.read_csv('Dataset_Day13.csv')
df.info()
# pair plot for additional insight
sns.pairplot(df)
plt.show()
# calculate missing-value percentage
missing_value_percent = df.isna().sum() / len(df) * 100
print(missing_value_percent)
df.boxplot(column=['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm'])
plt.title('Box Plot')
```

```

plt.show()
OutlierData = pd.DataFrame()
temp = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
for col in ['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']:
    Q1 = temp[col].quantile(0.25)    # Gives 25th
Percentile or Q1
    Q3 = temp[col].quantile(0.75)    # Gives 75th
Percentile or Q3

    IQR = Q3 - Q1

    UpperBound = Q3 + 1.5 * IQR
    LowerBound = Q1 - 1.5 * IQR

    OutlierData[col] = temp[col][(temp[col] <
LowerBound) | (temp[col] > UpperBound)]
    print(len(OutlierData))
# group the dataset based on the "Species" column
descriptive_stats =
df.groupby('Species').describe()
print(descriptive_stats)
X = df[['SepalLengthCm', 'SepalWidthCm',
'PetalLengthCm', 'PetalWidthCm']]
# specify the number of clusters
k = 3
# create a kmeans instance
km = KMeans(n_clusters=k, n_init=25,
random_state=1234)
# fit the data to the kmeans model
km.fit(X)
# get the cluster labels for each data point
cluster_labels = km.labels_
# the total within cluster sum of squares
clusterWCSS = km.inertia_
print(cluster_labels)
print(clusterWCSS)
# tabulation of the size of the clusters
pd.Series(km.labels_).value_counts().sort_index()
# 'km.cluster_centers_' :gives the cluster
centroids

```

```

cluster_centers = pd.DataFrame(km.cluster_centers_,
                                columns=['SepalLengthCm', 'SepalWidthCm',
                                         'PetalLengthCm', 'PetalWidthCm'])
print(cluster_centers)
# to get the correct centroids, we need to un-scale
the data,
cluster_centers_unscaled = pd.DataFrame()
for i in ['SepalLengthCm', 'SepalWidthCm',
          'PetalLengthCm', 'PetalWidthCm']:
    cluster_centers_unscaled[i] =
    (cluster_centers[i] * df[i].std()) + df[i].mean()
print(cluster_centers_unscaled)
wcss = []
for k in range(2, 11):
    km = KMeans(n_clusters=k, n_init=25,
                random_state=1234)
    km.fit(X)
    wcss.append(km.inertia_)

plt.plot(range(2, 11), wcss)
plt.xlabel('Number of Clusters')
plt.ylabel('WCSS')
plt.title('Elbow Method')
plt.show()
silhouette = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
                random_state = 1234)
    km.fit(X)
    silhouette.append(silhouette_score(X,
                                        km.labels_))

plt.plot(range(2, 11), silhouette)
plt.xlabel('Number of Clusters')
plt.ylabel('Silhouette Score')
plt.title('Silhouette Method')
plt.show()
calinski = []
for k in range(2, 11):
    km = KMeans(n_clusters = k, n_init = 25,
                random_state = 1234)

```

```
km.fit(X)
calinski.append(calinski_harabasz_score(X,
km.labels_))

plt.plot(range(2, 11), calinski)
plt.xlabel('Number of Clusters')
plt.ylabel('Calinski Harabasz Score')
plt.title('Calinski Harabasz Score')
plt.show()

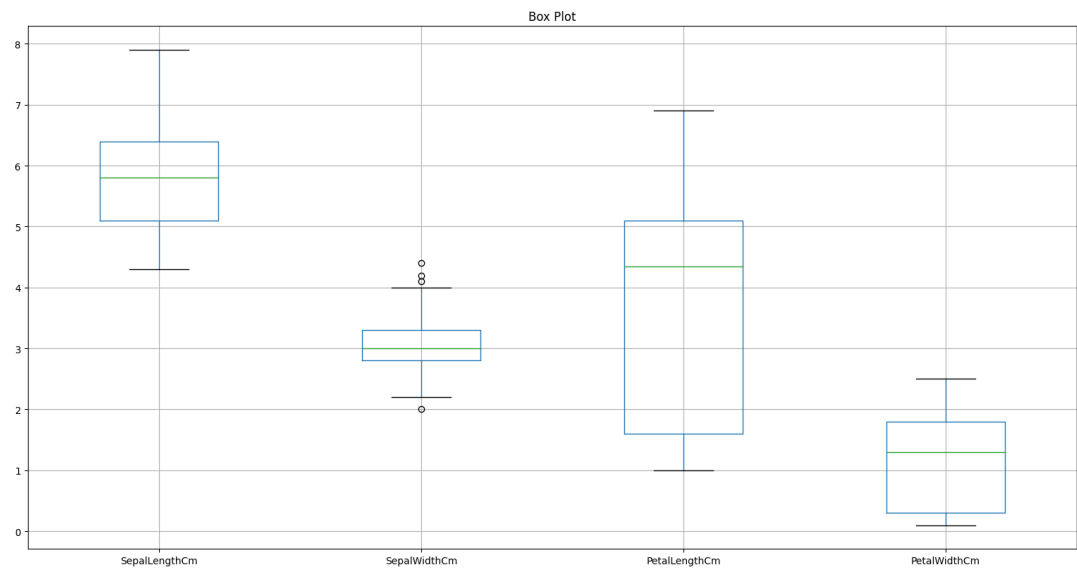
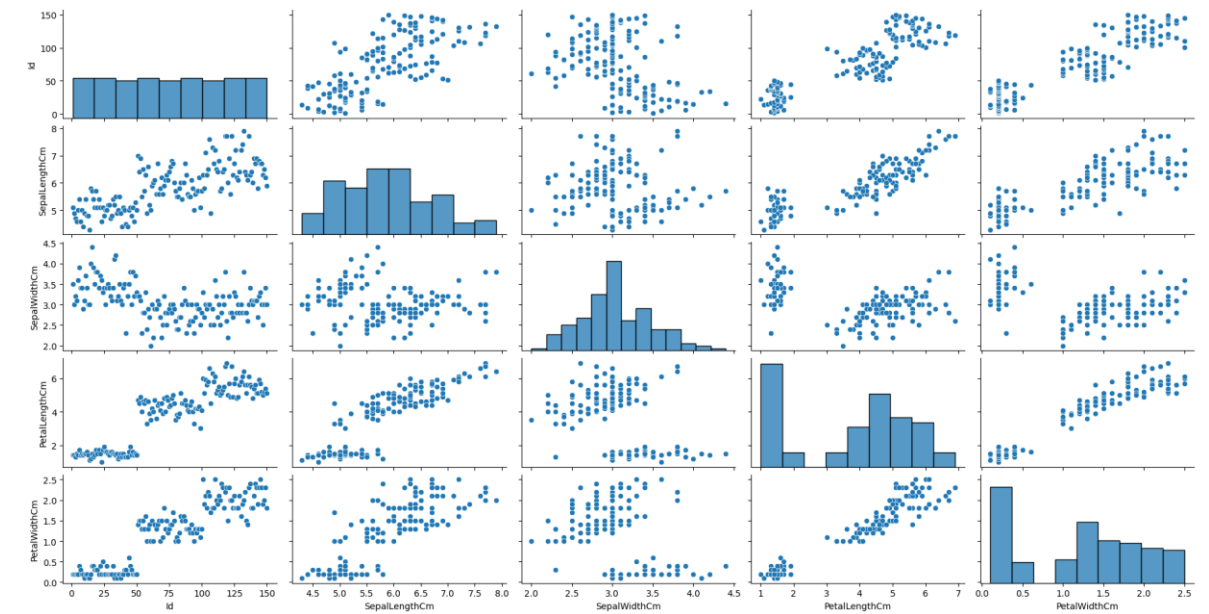
cluster_df = pd.DataFrame({'Cluster':
cluster_labels, 'Species': df['Species']})
cross_tab = pd.crosstab(cluster_df['Cluster'],
cluster_df['Species'])
proportion = cross_tab.div(cross_tab.sum(axis=1),
axis=0) * 100
print(proportion)

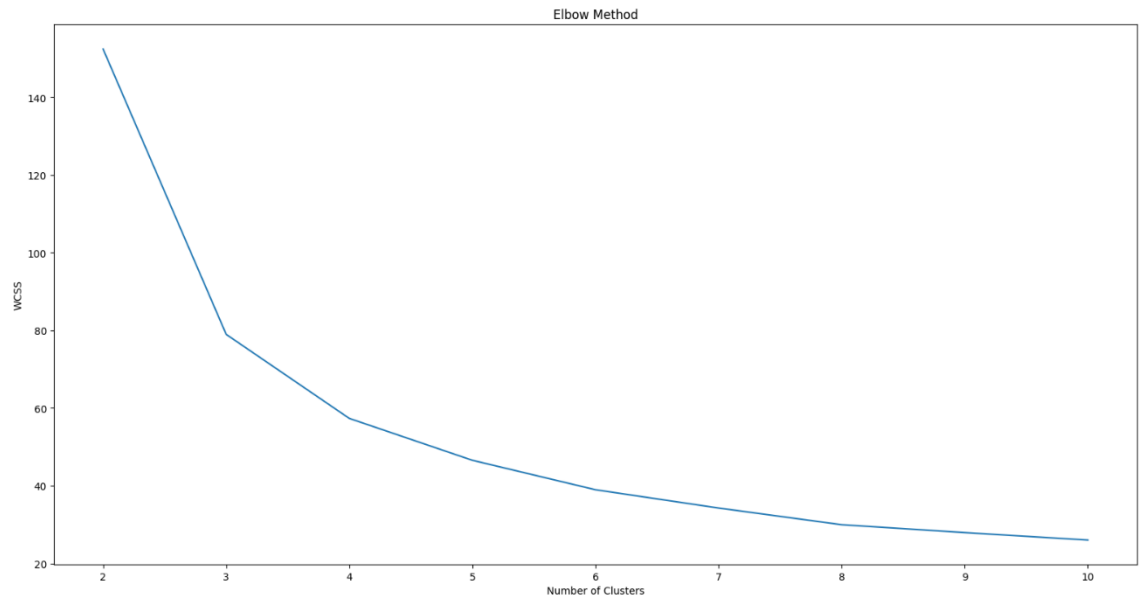
sns.scatterplot(x='PetalLengthCm',
y='PetalWidthCm',
hue='Species', data=df, )

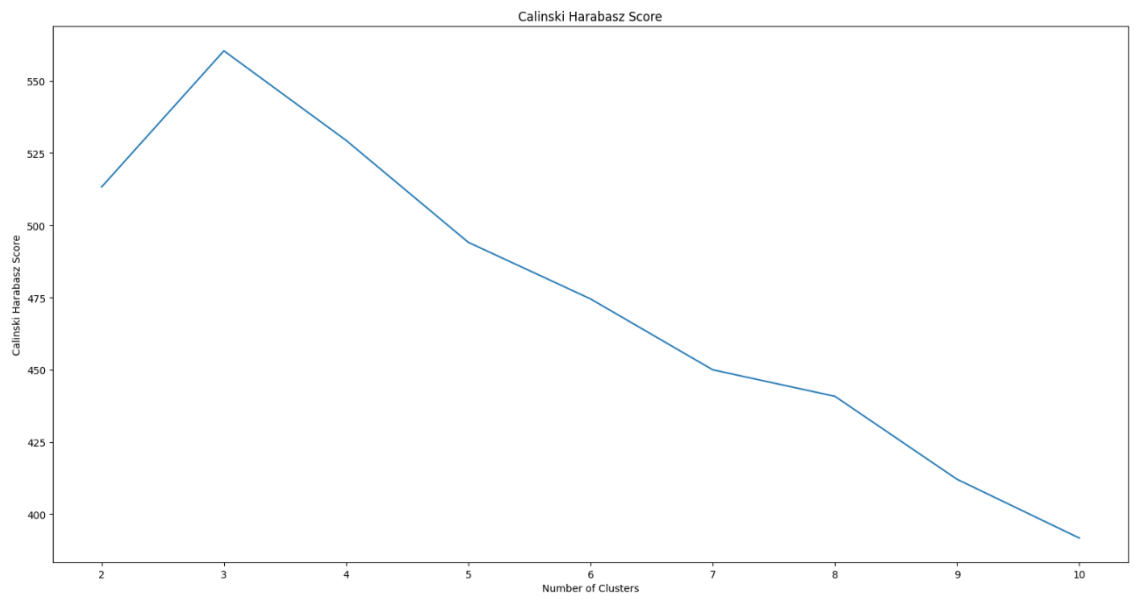
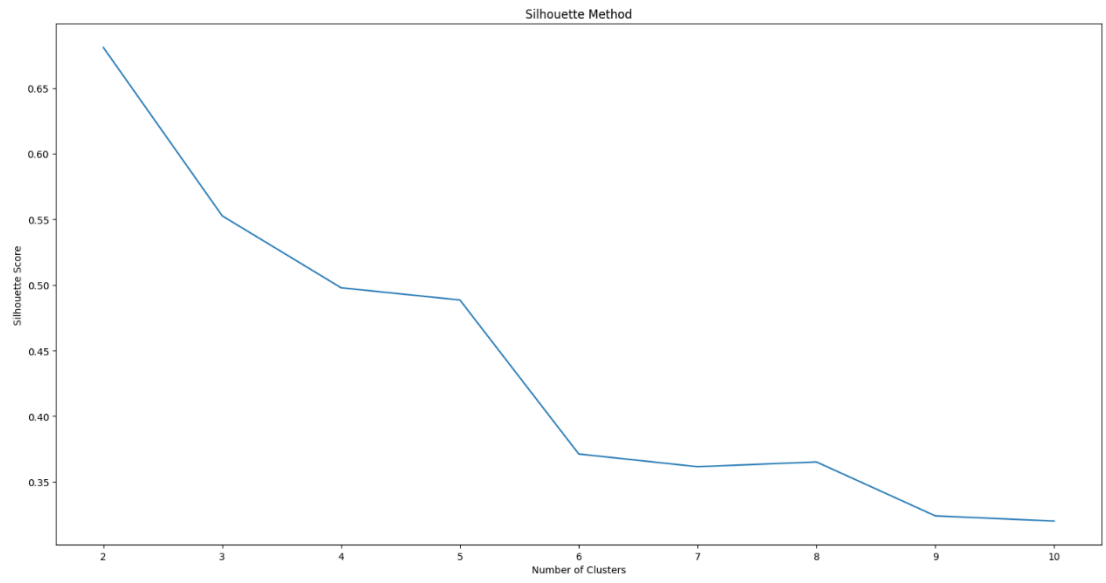
plt.show()

sns.scatterplot(x='SepalLengthCm',
y='SepalWidthCm',
hue='Species', data=df, )

plt.show()
```







[illegible]



File - Day13Q5

```
52 2          0.0      77.419355    22.580645
53
54 Process finished with exit code 0
55
```

