

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
from sklearn.cluster import KMeans
from sklearn import preprocessing
import sklearn.cluster as cluster
import sklearn.metrics as metrics
from sklearn.preprocessing import MinMaxScaler
```

```
data=pd.read_csv("/content/iris.csv")
```

data

	Id	SepallLengthCm	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
...
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 6 columns

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```
data.info()
```

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

```
data.head(10)
```

	Id	SepallLengthCm	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
5	6	5.4	3.9	1.7	0.4	Iris-setosa
6	7	4.6	3.4	1.4	0.3	Iris-setosa
7	8	5.0	3.4	1.5	0.2	Iris-setosa
8	9	4.4	2.9	1.4	0.2	Iris-setosa
9	10	4.9	3.1	1.5	0.1	Iris-setosa

```
data.describe()
```

	Id	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	

data.isnull()

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	False	False	False	False	False	False
1	False	False	False	False	False	False
2	False	False	False	False	False	False
3	False	False	False	False	False	False
4	False	False	False	False	False	False
...
145	False	False	False	False	False	False
146	False	False	False	False	False	False
147	False	False	False	False	False	False
148	False	False	False	False	False	False
149	False	False	False	False	False	False

150 rows × 6 columns

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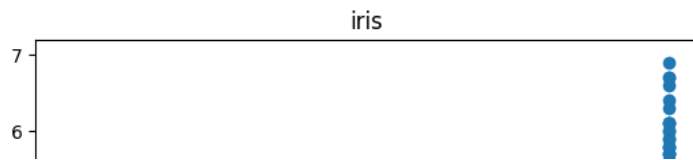
```
Id      0
SepalLengthCm  0
SepalWidthCm  0
PetalLengthCm  0
PetalWidthCm  0
Species      0
dtype: int64
```

data.columns

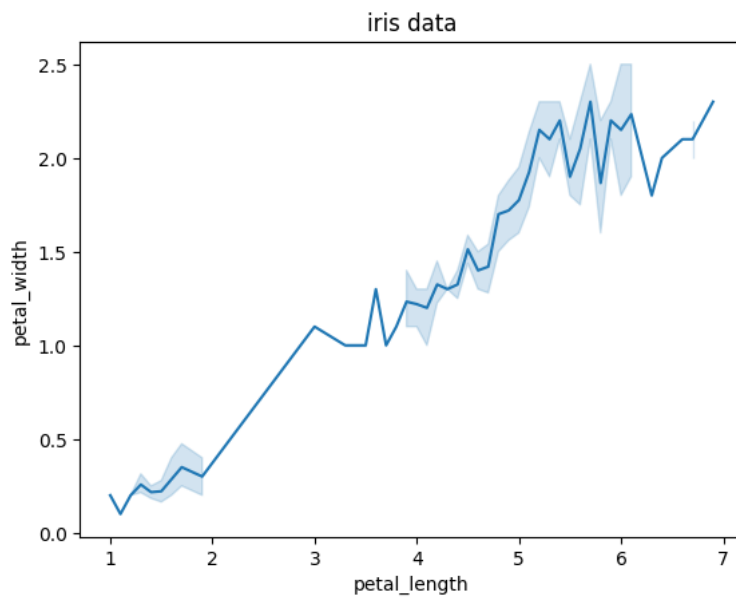
```
Index(['Id', 'SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm',
      'Species'],
      dtype='object')
```

data.rename(columns={'id':'id','SepalLengthCm':'sepal_length','SepalWidthCm':'sepal_width','PetalLengthCm':'petal_length','PetalWidthCm':

```
plt.scatter(data['Species'], data['petal_length'])
plt.title("iris")
plt.xlabel('Species')
plt.ylabel('petal_length')
plt.show()
```



```
sns.lineplot(x="petal_length", y="petal_width" ,data=data)
plt.title('iris data')
plt.show()
```



```
data['Species'].value_counts()
```

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```
Name: Species, dtype: int64
```

```
print(data["Species"].unique())
```

```
['Iris-setosa' 'Iris-versicolor' 'Iris-virginica']
```

```
plt.figure(figsize=(12, 6))
sns.histplot(data.sepal_length)
plt.xlabel('f')
plt.ylabel('Sepal Length')
plt.title('Histogram of Sepal Length (Cm)', size=16)
```

Text(0.5, 1.0, 'Histogram of Sepal Length (Cm)')

Histogram of Sepal Length (Cm)

```
correlation = data.corr()
correlation
```

<ipython-input-54-521f87fcc686>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a f
correlation = data.corr()

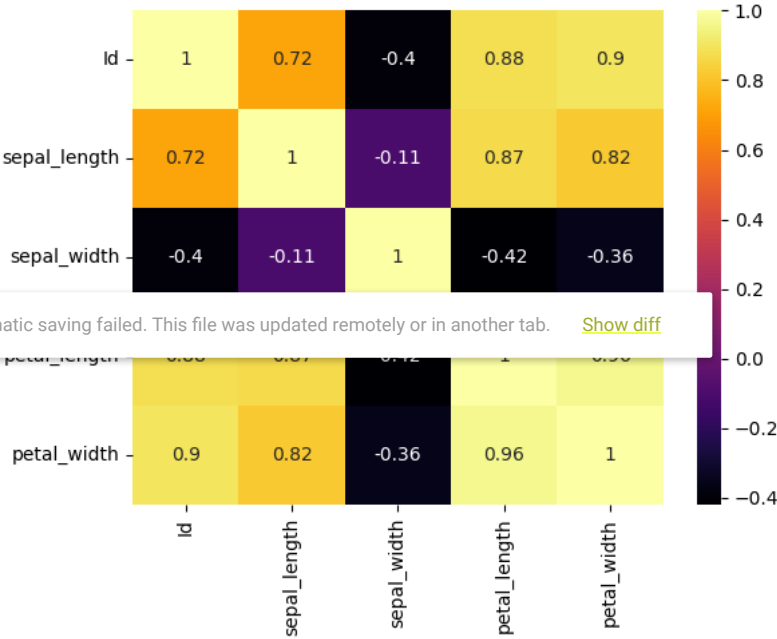
	Id	sepal_length	sepal_width	petal_length	petal_width
Id	1.000000	0.716676	-0.397729	0.882747	0.899759
sepal_length	0.716676	1.000000	-0.109369	0.871754	0.817954
sepal_width	-0.397729	-0.109369	1.000000	-0.420516	-0.356544
petal_length	0.882747	0.871754	-0.420516	1.000000	0.962757
petal_width	0.899759	0.817954	-0.356544	0.962757	1.000000



```
sns.heatmap(data.corr(),annot=True,cmap='inferno')
```

<ipython-input-55-c46cf576d981>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a f
sns.heatmap(data.corr(),annot=True,cmap='inferno')

<Axes: >




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```
sns.pairplot(data)
```

The figure displays a 6x6 matrix of plots illustrating the relationships between six variables: sepal_width, sepal_length, petal_length, petal_width, sepal_area, and petal_area. The diagonal elements are histograms showing the distribution of each variable. The off-diagonal elements are scatter plots showing the pairwise relationships between the variables. The variables are ordered as sepal_width, sepal_length, petal_length, petal_width, sepal_area, and petal_area from top-left to bottom-right.

```
scale = scaler.fit_transform(data[["sepal_length", "sepal_width", "petal_length", "petal_width"]])

data_scale = pd.DataFrame(scale, columns = ["sepal_length", "sepal_width", "petal_length", "petal_width"]);
```

	sepal_length	sepal_width	petal_length	petal_width	
0	0.222222	0.625000	0.067797	0.041667	
1	0.166667	0.416667	0.067797	0.041667	
2	0.111111	0.500000	0.050847	0.041667	
3	0.083333	0.458333	0.084746	0.041667	
4	0.194444	0.666667	0.067797	0.041667	

	sepal_length	sepal_width	petal_length	petal_width
0	0.222222	0.625000	0.067797	0.041667
1	0.166667	0.416667	0.067797	0.041667
2	0.111111	0.500000	0.050847	0.041667
3	0.083333	0.458333	0.084746	0.041667
4	0.194444	0.666667	0.067797	0.041667
..
145	0.666667	0.416667	0.711864	0.916667
146	0.555556	0.208333	0.677966	0.750000
147	0.611111	0.416667	0.711864	0.791667
148	0.527778	0.583333	0.745763	0.916667
149	0.444444	0.416667	0.694915	0.708333

[illegible]

```

km.cluster_centers_

array([[5.00566038, 3.36037736, 1.56226415, 0.28867925],
       [6.30103093, 2.88659794, 4.95876289, 1.69587629]])

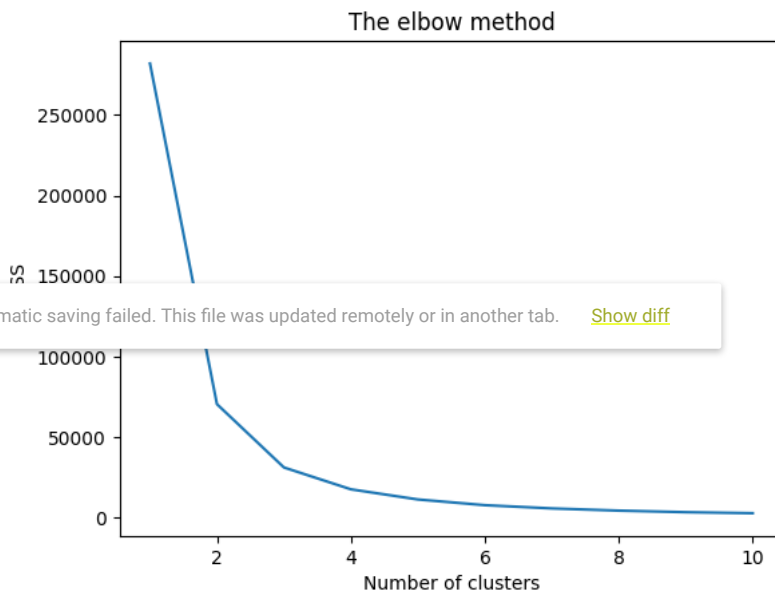
X= data.iloc[:, [0, 1, 2, 3]].values

from sklearn.cluster import KMeans
wcss = []

for i in range(1, 11):
    kmeans = KMeans(n_clusters = i, init = 'k-means++',
                    max_iter = 300, n_init = 10, random_state = 0)
    kmeans.fit(x)
    wcss.append(kmeans.inertia_)

# Plotting the results onto a line graph,
# `allowing us to observe 'The elbow'
plt.plot(range(1, 11), wcss)
plt.title('The elbow method')
plt.xlabel('Number of clusters')
plt.ylabel('WCSS') # Within cluster sum of squares
plt.show()

```



```

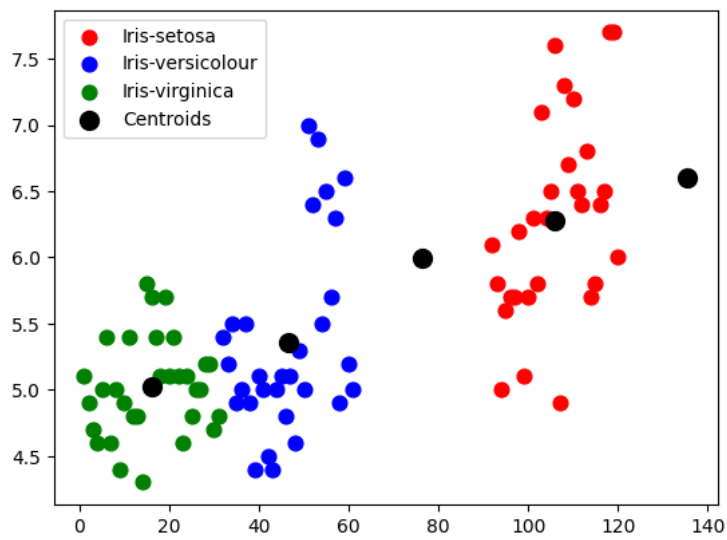
kmeans = KMeans(n_clusters = 5, init = "k-means++", random_state = 42)
y_kmeans = kmeans.fit_predict(X)

/usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change fr
warnings.warn(

plt.scatter(X[y_kmeans == 0, 0], X[y_kmeans == 0, 1], s = 60, c = 'red', label = 'Iris-setosa')
plt.scatter(X[y_kmeans == 1, 0], X[y_kmeans == 1, 1], s = 60, c = 'blue', label = 'Iris-versicolour')
plt.scatter(X[y_kmeans == 2, 0], X[y_kmeans == 2, 1], s = 60, c = 'green', label = 'Iris-virginica')
plt.scatter(kmeans.cluster_centers_[0, 0], kmeans.cluster_centers_[0, 1], s = 100, c = 'black', label = 'Centroids')
plt.legend()

plt.show()

```



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