

## 54. K-Means Clustering (Practical)

```
In [1]: import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [2]: dataset = pd.read_csv(r'Data/iris_raw.csv')
dataset.head(3)
```

```
Out[2]:
```

	sepal_length	sepal_width	petal_length	petal_width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2

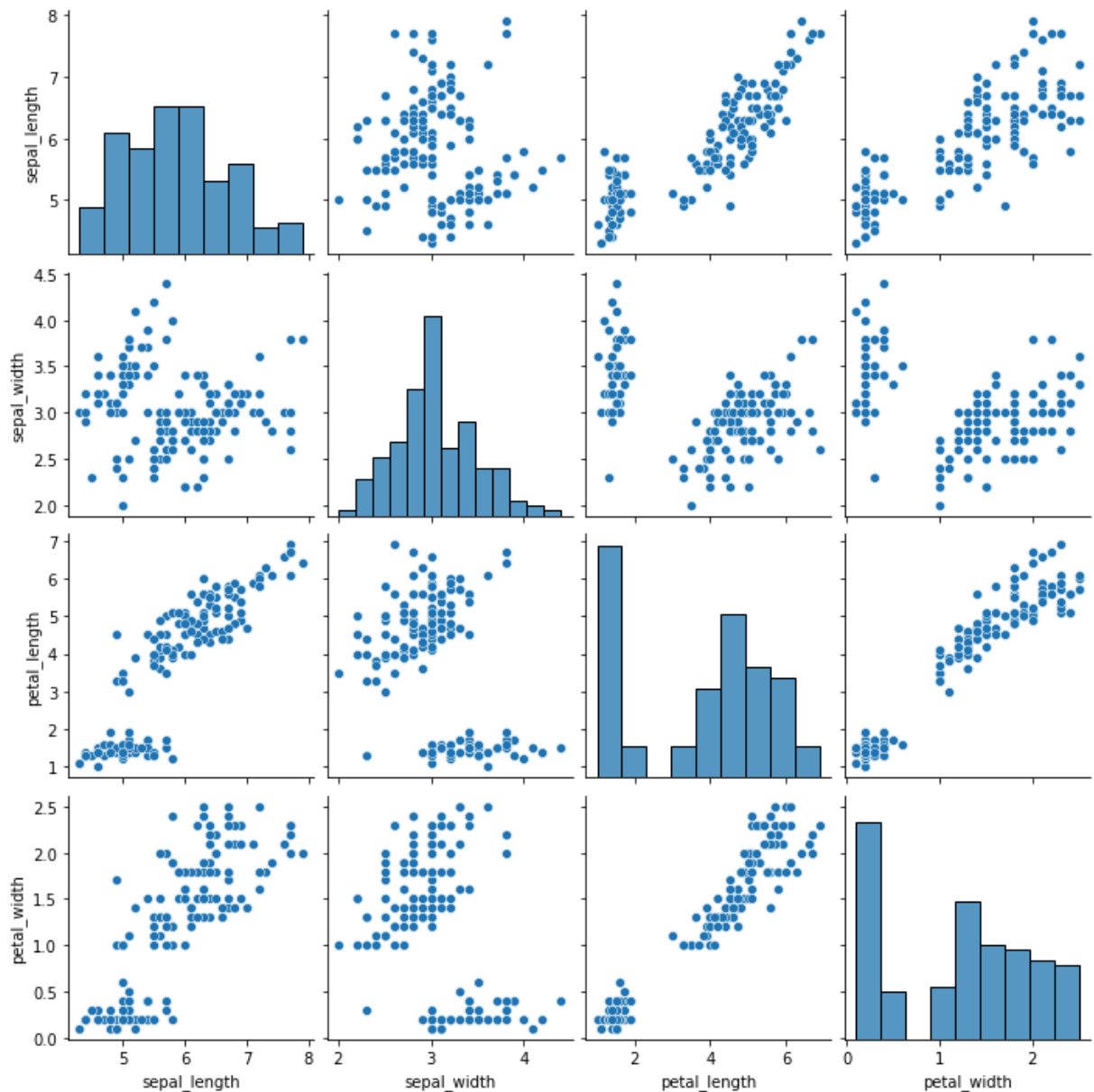
### 54.1 Making Clusters of Data

- Use K-mean clustering when **your data is linearly separable**

#### Check the data if it is linearly separable

```
In [3]: sns.pairplot(data=dataset)
plt.show()
```

```
C:\Users\rashi\AppData\Local\Programs\Python\Python39\lib\site-packages\seaborn\axis
grid.py:123: UserWarning: The figure layout has changed to tight
self._figure.tight_layout(*args, **kwargs)
```



- In supervised learning, the data is split into training and testing data
- In unsupervised learning, data is not split into training and testing data b/c the data is unlabelled

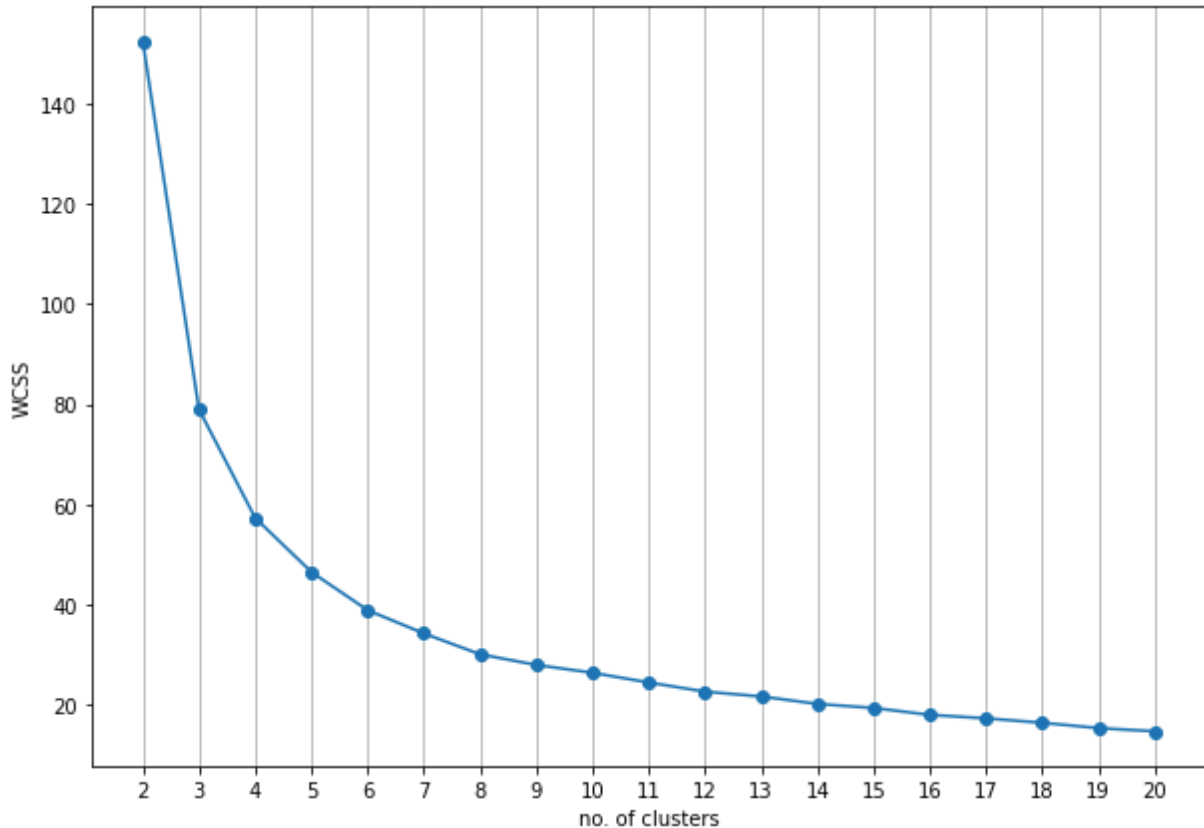
## 54.1.1 Find Number of clusters

```
In [7]: from sklearn.cluster import KMeans
```

```
In [14]: # Use a loop to find best number of clusters from 2 to 20
wcsc = []

for i in range(2,21):
    km = KMeans(n_clusters=i, init='k-means++')
    km.fit(dataset)
    wcsc.append(km.inertia_) # it assigns value of wcsc {Elbow graph}
```

```
In [29]: plt.figure(figsize=(10,7))
plt.plot([i for i in range(2,21)], wcss, marker='o')
plt.xlabel('no. of clusters')
plt.xticks([i for i in range(2,21)])
plt.ylabel('WCSS')
plt.grid(axis='x')
plt.show()
```



**Elbow point = 3**

**It means that will have 3 number of clusters**

```
In [ ]:
```

```
In [30]: kmn = KMeans(n_clusters=3)
kmn.fit_predict(dataset)
```

```
Out[30]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 1, 2, 2, 2, 2, 2, 1, 2, 2, 2,
2, 2, 2, 1, 1, 2, 2, 2, 2, 1, 2, 1, 2, 1, 2, 2, 1, 1, 2, 2, 2, 2,
2, 1, 2, 2, 2, 2, 1, 2, 2, 2, 1, 2, 2, 2, 1, 2, 2, 1])
```

```
In [32]: dataset['Predict'] = kmn.fit_predict(dataset)
```

```
In [33]: dataset
```

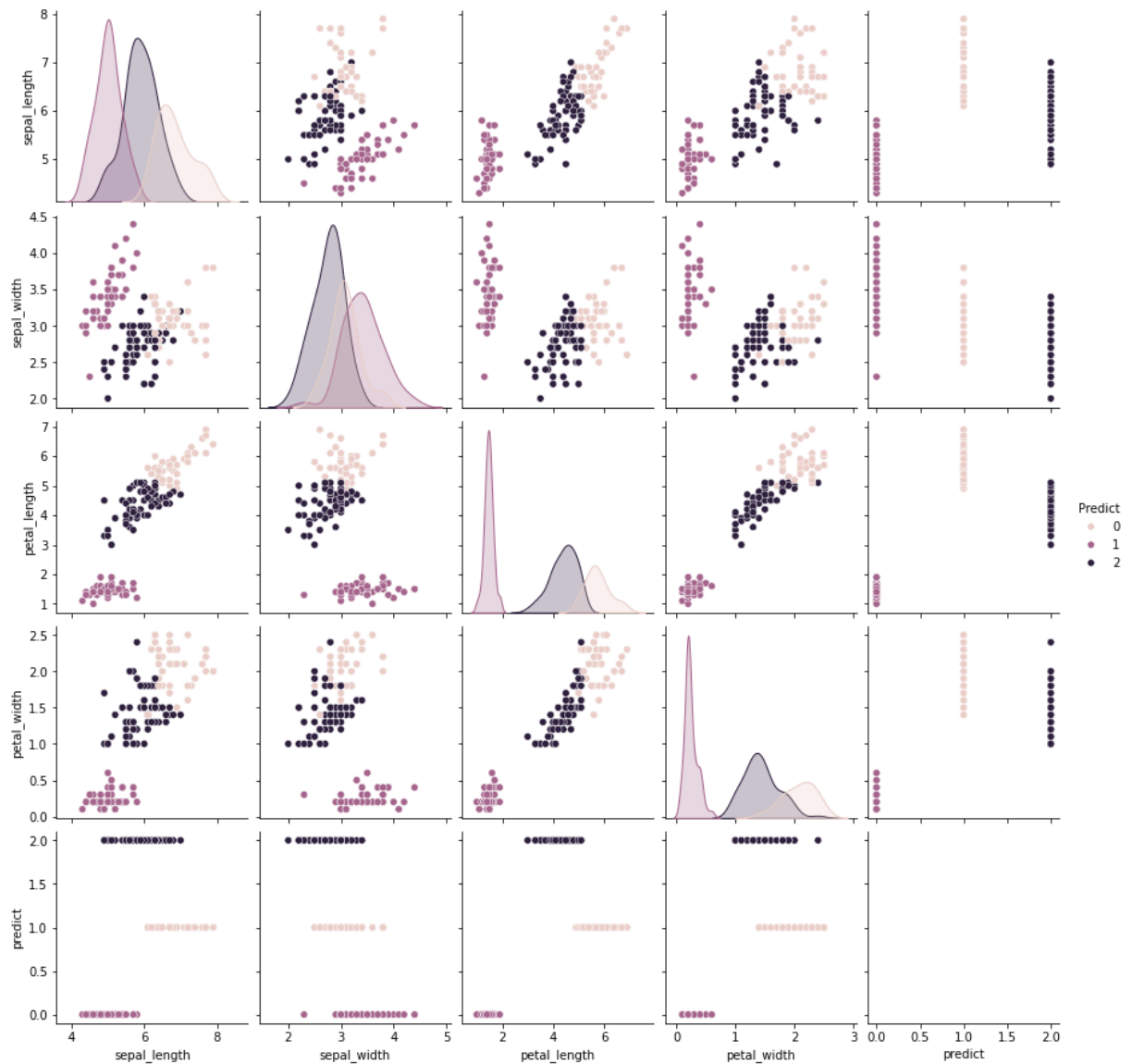
Out[33]:

	sepal_length	sepal_width	petal_length	petal_width	predict	Predict
<b>0</b>	5.1	3.5	1.4	0.2	0	1
<b>1</b>	4.9	3.0	1.4	0.2	0	1
<b>2</b>	4.7	3.2	1.3	0.2	0	1
<b>3</b>	4.6	3.1	1.5	0.2	0	1
<b>4</b>	5.0	3.6	1.4	0.2	0	1
<b>...</b>	...	...	...	...	...	...
<b>145</b>	6.7	3.0	5.2	2.3	1	0
<b>146</b>	6.3	2.5	5.0	1.9	2	2
<b>147</b>	6.5	3.0	5.2	2.0	1	0
<b>148</b>	6.2	3.4	5.4	2.3	1	0
<b>149</b>	5.9	3.0	5.1	1.8	2	2

150 rows × 6 columns

```
In [39]: sns.pairplot(data=dataset, hue='Predict')
plt.savefig(r"Generated_images/raw-iris-clustering-predict.jpg")
plt.show()
```

```
C:\Users\rashi\AppData\Local\Programs\Python\Python39\lib\site-packages\seaborn\axis
grid.py:123: UserWarning: The figure layout has changed to tight
self._figure.tight_layout(*args, **kwargs)
```



## 54.2 Making raw data with original data

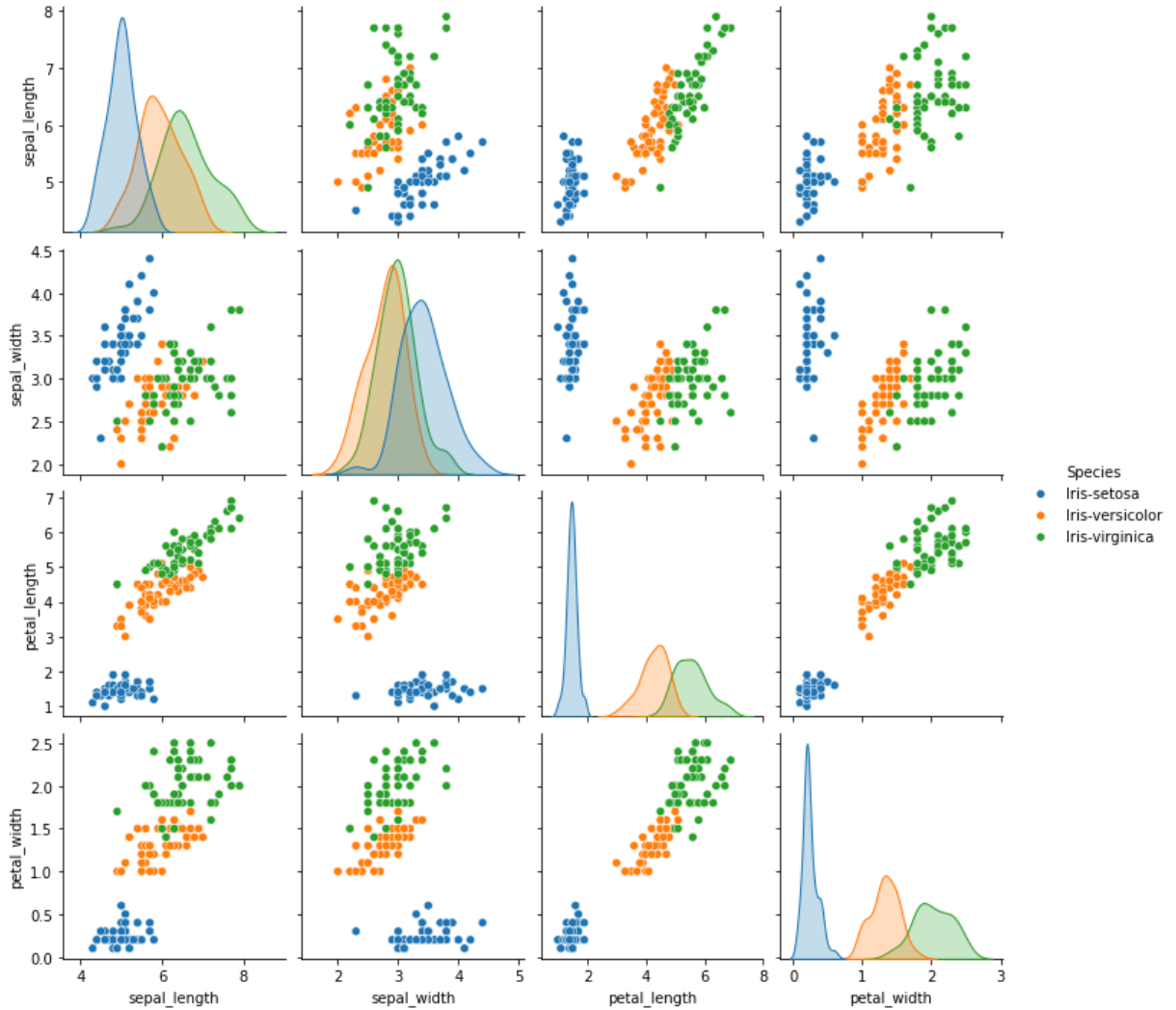
```
In [35]: org_dataset = pd.read_csv(r'Data/iris.csv')
org_dataset.head(3)
```

```
Out[35]:
```

	sepal_length	sepal_width	petal_length	petal_width	Species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa

```
In [40]: sns.pairplot(data=org_dataset, hue='Species')
plt.savefig(r"Generated_images/raw-iris-clustering-original-data.jpg")
plt.show()
```

```
C:\Users\rashi\AppData\Local\Programs\Python\Python39\lib\site-packages\seaborn\axis
grid.py:123: UserWarning: The figure layout has changed to tight
self._figure.tight_layout(*args, **kwargs)
```



In [ ]: