

```
In [9]: import numpy as np
import pandas as pd
from sklearn.cluster import KMeans as km
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [18]: df=pd.read_csv(r"C:\Users\lahar\Downloads\archive (7)\IRIS.csv")
df.head()
```

Out[18]:

	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
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa

```
In [19]: df['species'],categories = pd.factorize(df['species'])
df.head()
```

Out[19]:

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0

```
In [20]: df.describe
```

```
Out[20]: <bound method NDFrame.describe of
ngth petal_width species
0      5.1      3.5      1.4      0.2      0
1      4.9      3.0      1.4      0.2      0
2      4.7      3.2      1.3      0.2      0
3      4.6      3.1      1.5      0.2      0
4      5.0      3.6      1.4      0.2      0
..      ...      ...      ...      ...      ...
145     6.7      3.0      5.2      2.3      2
146     6.3      2.5      5.0      1.9      2
147     6.5      3.0      5.2      2.0      2
148     6.2      3.4      5.4      2.3      2
149     5.9      3.0      5.1      1.8      2

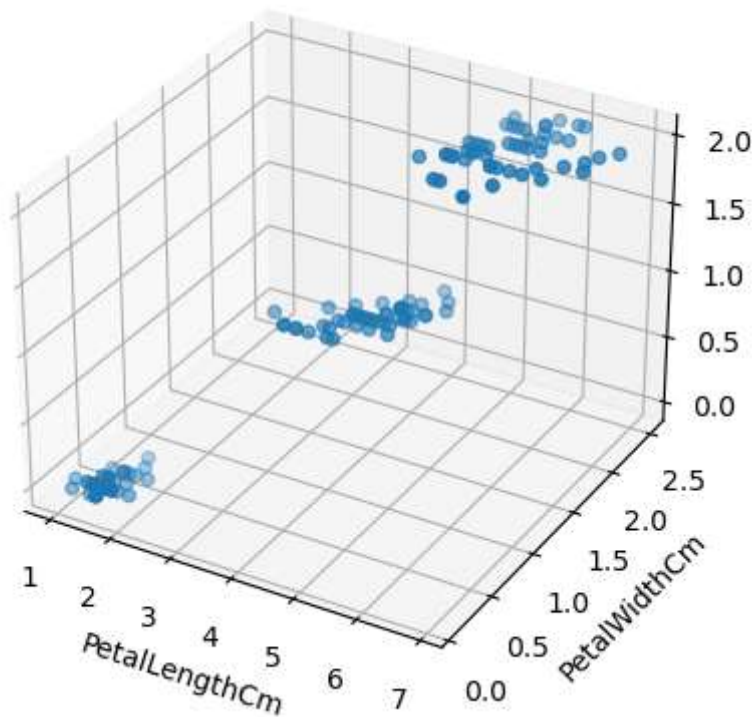
[150 rows x 5 columns]>
```

```
In [21]: df.isna().sum()
```

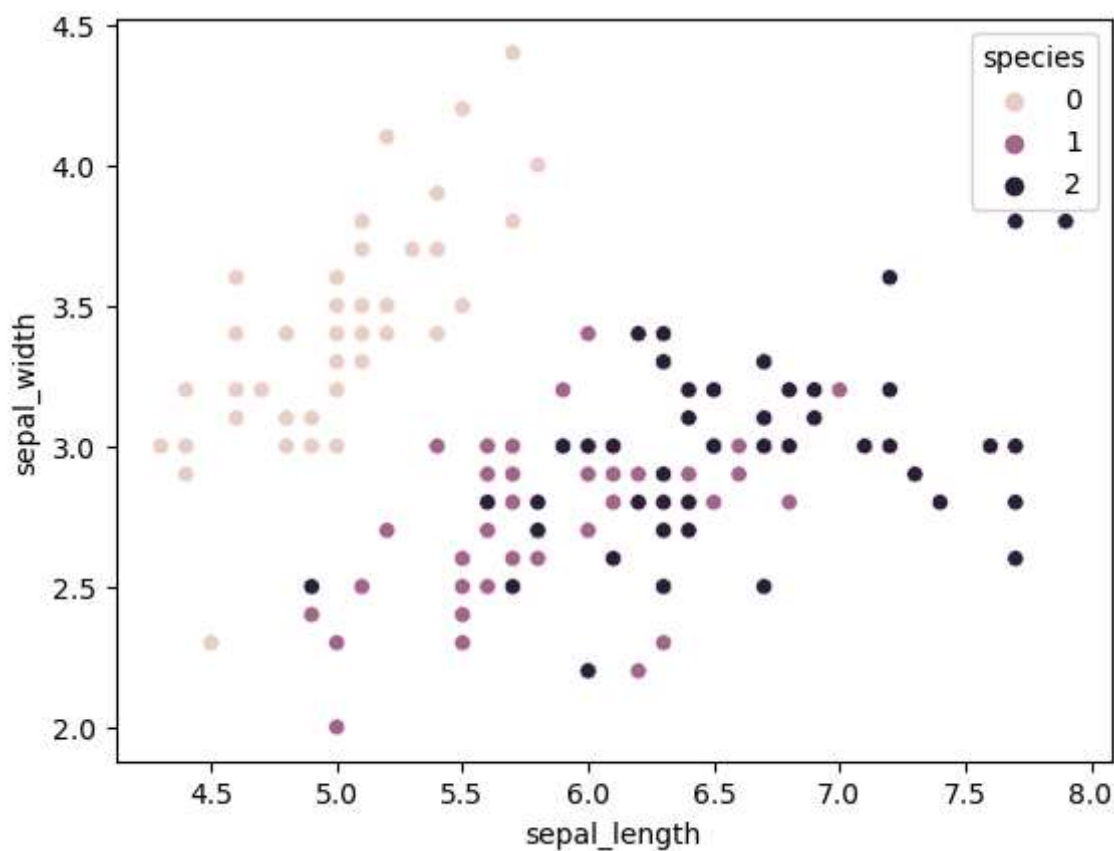
```
Out[21]: sepal_length    0  
sepal_width    0  
petal_length    0  
petal_width    0  
species        0  
dtype: int64
```

```
In [22]: from mpl_toolkits.mplot3d import Axes3D  
fig = plt.figure()  
ax=fig.add_subplot(111,projection='3d')  
ax.scatter(df.petal_length, df.petal_width, df.species)  
ax.set_xlabel('PetalLengthCm')  
ax.set_ylabel('PetalWidthCm')  
ax.set_zlabel('Species')  
plt.title('3D scatter plot Example')  
plt.show()
```

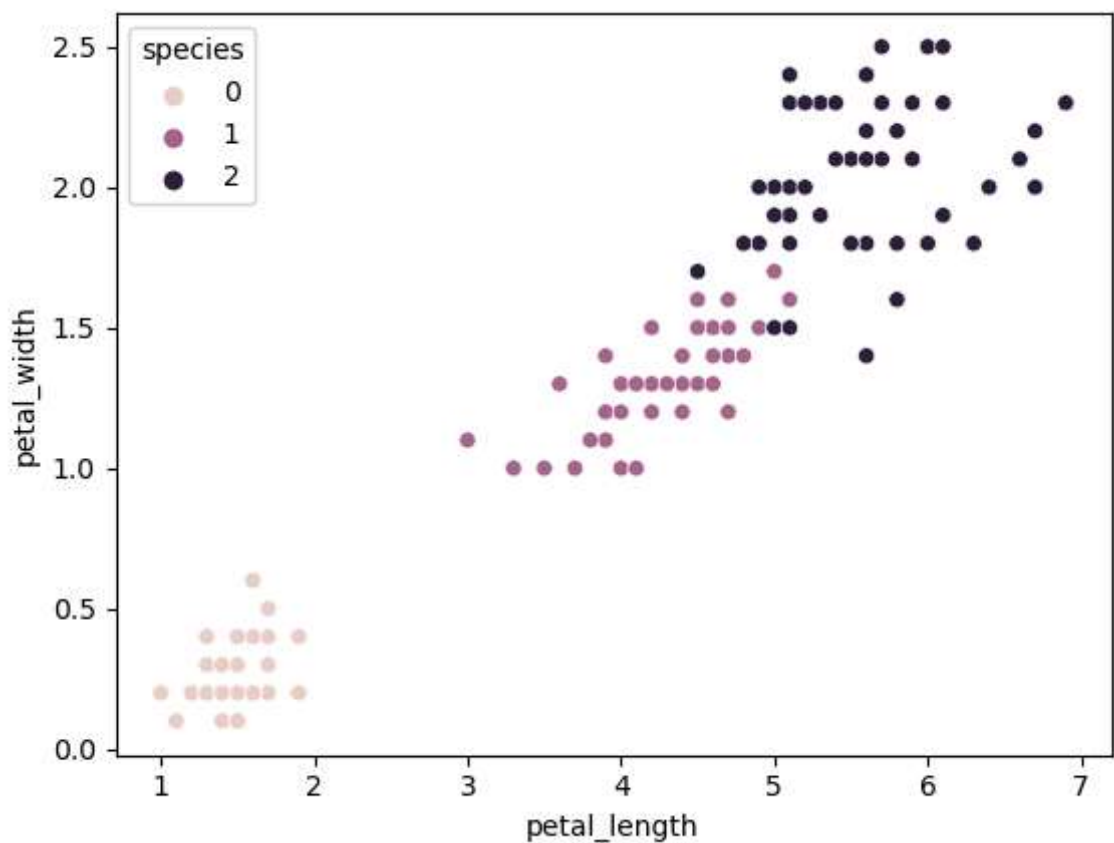
3D scatter plot Example



```
In [23]: sns.scatterplot(data=df, x="sepal_length", y="sepal_width", hue="species");
```



```
In [24]: sns.scatterplot(data=df, x="petal_length", y="petal_width", hue="species");
```



```
In [29]: k_rrg = range(1,10)
sse=[]

for k in k_rrg:
    km= KMeans(n_clusters=k)
    km.fit(df[['petal_length', 'petal_width']])
    sse.append(km.inertia_)
```

```

C:\Users\lahar\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:141
2: FutureWarning: The default value of `n_init` will change from 10 to 'au
to' in 1.4. Set the value of `n_init` explicitly to suppress the warning
    super()._check_params_vs_input(X, default_n_init=10)
C:\Users\lahar\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:143
6: UserWarning: KMeans is known to have a memory leak on Windows with MKL,
when there are less chunks than available threads. You can avoid it by set
ting the environment variable OMP_NUM_THREADS=1.
    warnings.warn(
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```

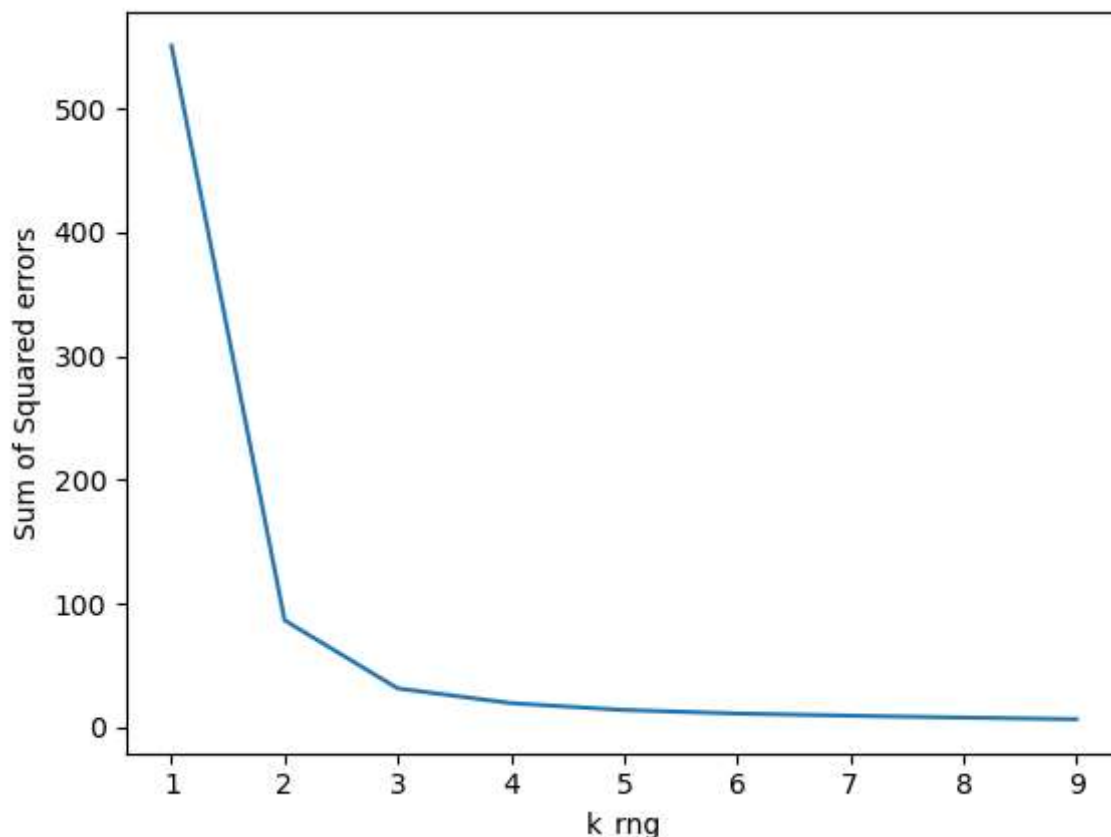
```
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C:\Users\lahar\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:141
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ting the environment variable OMP_NUM_THREADS=1.
warnings.warn(
```

In [30]: sse

```
Out[30]: [550.6434666666668,
86.40394533571005,
31.387758974358974,
19.49352336396547,
13.933308757908756,
11.10362366260479,
9.344727985739752,
7.836495238095239,
6.496659206692711]
```

```
In [33]: plt.xlabel('k_rng')
plt.ylabel("Sum of Squared errors")
plt.plot(k_rng, sse)
```

```
Out[33]: [<matplotlib.lines.Line2D at 0x24f97d09a50>]
```



```
In [34]: km = KMeans(n_clusters=3,random_state=0,)
y_predicted = km.fit_predict(df[['petal_length','petal_width']])
y_predicted
```

C:\Users\lahar\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:141
 2: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
 super()._check_params_vs_input(X, default_n_init=10)
 C:\Users\lahar\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:143
 6: UserWarning: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=1.
 warnings.warn(

```
Out[34]: array([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, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 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, 2, 0, 0, 0, 0, 0, 2, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2,
2, 2, 2, 2, 2, 2, 0, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

```
In [37]: df['cluster']=y_predicted  
df.head(150)
```

Out[37]:

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

150 rows × 6 columns

```
In [41]: from sklearn.metrics import confusion_matrix  
cm =confusion_matrix(df.species, df.cluster)  
cm
```

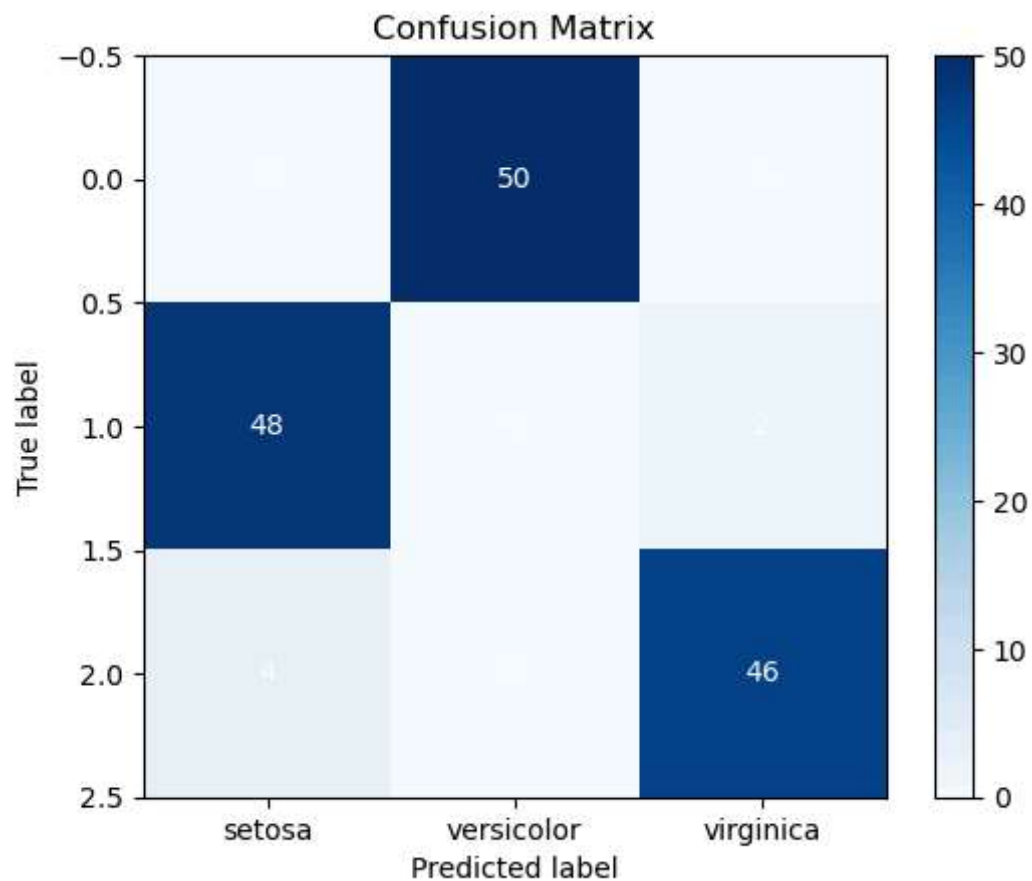
Out[41]: array([[0, 50, 0],
 [48, 0, 2],
 [4, 0, 46]], dtype=int64)


```
In [42]: true_labels = df.species
predicted_labels = df.cluster

cm = confusion_matrix(true_labels, predicted_labels)
class_labels = ['setosa', 'versicolor', 'virginica']

plt.imshow(cm, interpolation='nearest', cmap=plt.cm.Blues)
plt.title('Confusion Matrix')
plt.colorbar()
tick_marks = np.arange(len(class_labels))
plt.xticks(tick_marks, class_labels)

for i in range(len(class_labels)):
    for j in range(len(class_labels)):
        plt.text(j, i, str(cm[i][j]), ha='center', va='center', color='white')
plt.xlabel('Predicted label')
plt.ylabel('True label')
plt.show()
```



In []: