```
In [3]: import numpy as np
import pandas as pd
from sklearn.cluster import KMeans as km
import matplotlib.pyplot as plt
import seaborn as sns

df=pd.read_csv(r"C:\Users\THANUJA\Downloads\IRIS.csv")
df.head()
```

Out[3]:

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

Out[4]:

	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

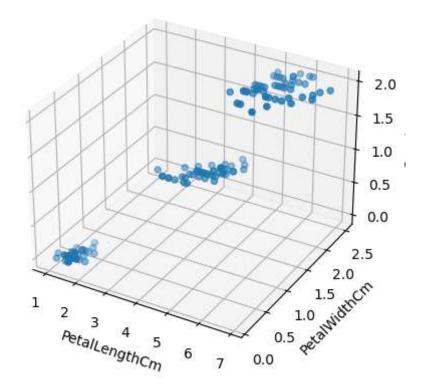
In [5]: df.describe

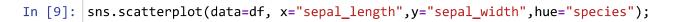
Out[5]:	<pre><bound h="" method="" petal="" pre="" widt<=""></bound></pre>	d NDFrame.desc	ribe of	sepal_length	sepal_width	petal_lengt
	0 <u> </u>	5.1	3.5	1.4	0.2	0
	1	4.9	3.0	1.4	0.2	0
	2	4.7	3.2	1.3	0.2	0
	3	4.6	3.1	1.5	0.2	0
	4	5.0	3.6	1.4	0.2	0
	• •	• • •	• • •	• • •		• •
	145	6.7	3.0	5.2	2.3	2
	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

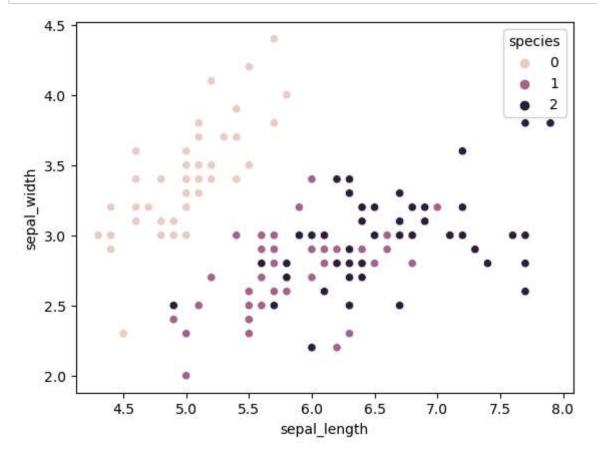
[150 rows x 5 columns]>

```
In [6]: df.isna().sum()
Out[6]: sepal_length
                         0
        sepal_width
                         0
        petal length
                        0
        petal_width
                         0
        species
                         0
        dtype: int64
In [8]: 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()
```

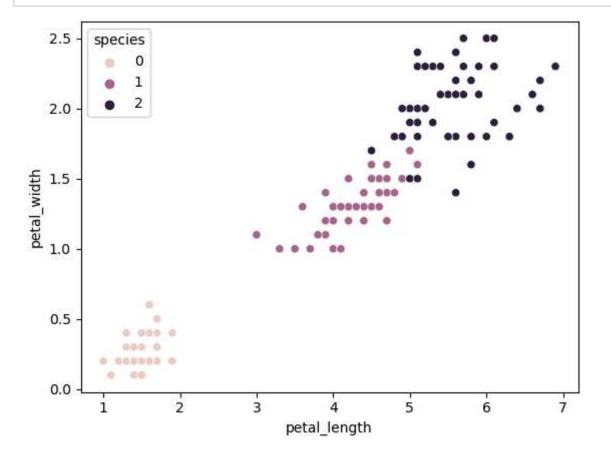
3d scatter plot Example







In [10]: sns.scatterplot(data=df, x="petal_length", y="petal_width",hue="species");



```
In [12]: from sklearn.cluster import KMeans
k_rng=range(1,10)
sse=[]

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

```
C:\Users\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster\ kmeans.py:1412:
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\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster\_kmeans.py:1436:
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(
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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\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:1436: 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(

C:\Users\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:1412: 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)

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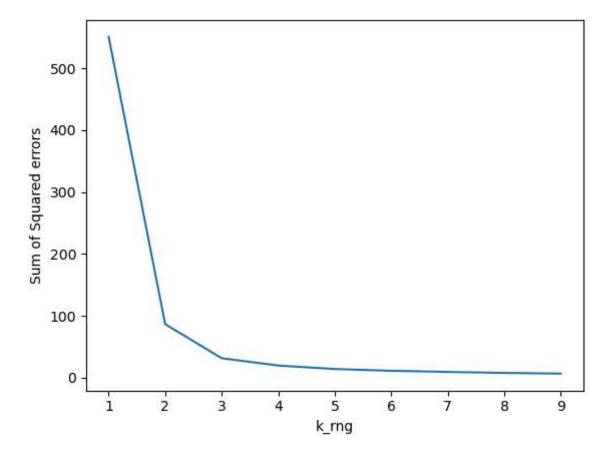
In [13]:

sse

Out[13]: [550.6434666666669, 86.40394533571003, 31.38775897435897, 19.499400899685114, 13.933308757908755, 11.067828739411809, 9.207570634920636, 7.631802244955954, 6.472894541406307]

```
In [14]: plt.xlabel('k_rng')
    plt.ylabel("Sum of Squared errors")
    plt.plot(k_rng, sse)
```

Out[14]: [<matplotlib.lines.Line2D at 0x2e6dee838d0>]



C:\Users\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:1412:
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C:\Users\THANUJA\anaconda3\Lib\site-packages\sklearn\cluster_kmeans.py:1436: 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(

```
In [16]: df['cluster']=y_predicted
    df.head(150)
```

Out[16]:		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

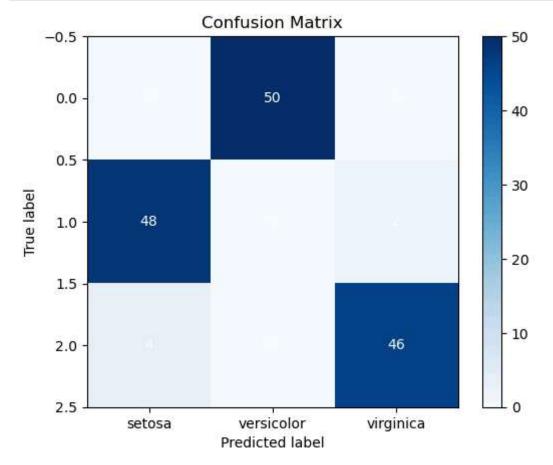
150 rows × 6 columns

```
In [17]: from sklearn.metrics import confusion_matrix
  cm=confusion_matrix(df.species, df.cluster)
  cm
```

```
In [19]: 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)):
        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 [ ]:
```