Breast cancer prediction

In [1]: import pandas as pd from matplotlib import pyplot as plt %matplotlib inline

In [2]: df=pd.read_csv(r"C:\Users\Y.Saranya\Downloads\BreastCancerPrediction.csv")

Out[2]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness
0	842302	М	17.99	10.38	122.80	1001.0	0
1	842517	М	20.57	17.77	132.90	1326.0	0
2	84300903	M	19.69	21.25	130.00	1203.0	0
3	84348301	M	11.42	20.38	77.58	386.1	0
4	84358402	М	20.29	14.34	135.10	1297.0	0
564	926424	М	21.56	22.39	142.00	1479.0	C
565	926682	М	20.13	28.25	131.20	1261.0	0
566	926954	М	16.60	28.08	108.30	858.1	0
567	927241	М	20.60	29.33	140.10	1265.0	0
568	92751	В	7.76	24.54	47.92	181.0	0
569 rows × 33 columns							

Data preprocessing

In [3]: df.head

Out[3]:	<bou< th=""><th>nd method NDFrame</th><th>e.head o</th><th>f</th><th>id diagnosis</th><th>radius mean</th><th>texture m</th></bou<>	nd method NDFrame	e.head o	f	id diagnosis	radius mean	texture m
	ean	perimeter_mean	area_mea		Ü	_	_
	0	842302 <u>–</u>	м —	17.99	10.38	122.80	1001.0
	1	842517	М	20.57	17.77	132.90	1326.0
	2	84300903	М	19.69	21.25	130.00	1203.0
	3	84348301	М	11.42	20.38	77.58	386.1
	4	84358402	М	20.29	14.34	135.10	1297.0
	• •	• • •	• • •	• • •	• • •	• • •	• • •
	564	926424	М	21.56	22.39	142.00	1479.0
	565	926682	М	20.13	28.25	131.20	1261.0
	566	926954	М	16.60	28.08	108.30	858.1
	567	927241	М	20.60	29.33	140.10	1265.0
	568	92751	В	7.76	24.54	47.92	181.0
		smoothness_mean	compac	tness mean	concavity mean	concave poi	nts mean
	\	_	•	_	7 –	•	_
	0	0.11840		0.27760	0.30010		0.14710
	1	0.08474		0.07864	0.08690		0.07017
	2	0.10960		0.15990	0.19740		0.12790
	3	0.14250		0.28390	0.24140		0.10520
	4	0.10030		0.13280	0.19800		0.10430
	 564	0.11100		 0.11590	0.24390		 0.13890
	565	0.09780		0.11390	0.14400		0.09791
	566	0.08455		0.10340	0.09251		0.05302
	567	0.11780		0.10230	0.35140		0.15200
	568	0.05263		0.27700	0.00000		0.00000
	300	0.03263		0.04302	0.0000		0.00000
		texture_wor	rst per:	imeter_worst	area_worst	smoothness <u>w</u> o	rst \
	0	17	. 33	184.60	2019.0	0.16	220
	1	23	.41	158.86	1956.0	0.12	380
	2	25	.53	152.56	1709.0	0.14	440
	3	26	.50	98.87	7 567.7	0.20	980
	4	16	.67	152.26	1575.0	0.13	740
		•••	• • •	455.46		0.44	
	564		.40	166.16		0.14	
	565		. 25	155.00		0.11	
	566		.12	126.76		0.11	
	567		.42	184.66		0.16	
	568	30	. 37	59.16	268.6	0.08	996
		compactness_wors	st conc	avity_worst	concave points	s_worst symm	etry_worst
	\						
	0	0.6656	50	0.7119		0.2654	0.4601
	1	0.1866	50	0.2416		0.1860	0.2750
	2	0.424	50	0.4504		0.2430	0.3613
	3	0.8663	30	0.6869		0.2575	0.6638
	4	0.205	90	0.4000		0.1625	0.2364
	 564		• •	0.4107		 0 2216	0 2060
	565	0.211				0.2216 0.1628	0.2060
		0.192		0.3215		0.1628	0.2572
	566 567	0.3094		0.3403		0.1418	0.2218
	567	0.8683		0.9387		0.2650	0.4087
	568	0.0644	+4	0.0000		0.0000	0.2871
		fractal dimension	on worst	Unnamed: 3	32		

fractal_dimension_worst Unnamed: 32 0.11890 NaN

0

1	0.08902	NaN
2	0.08758	NaN
3	0.17300	NaN
4	0.07678	NaN
• •	• • •	
564	0.07115	NaN
565	0.06637	NaN
566	0.07820	NaN
567	0.12400	NaN
568	0.07039	NaN

[569 rows x 33 columns]>

In [4]: df.tail()

Out[4]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_m
564	926424	М	21.56	22.39	142.00	1479.0	0.1
565	926682	М	20.13	28.25	131.20	1261.0	90.0
566	926954	М	16.60	28.08	108.30	858.1	30.0
567	927241	М	20.60	29.33	140.10	1265.0	0.1′
568	92751	В	7.76	24.54	47.92	181.0	0.0

5 rows × 33 columns

In [5]: df.drop(['Unnamed: 32'],axis=1)

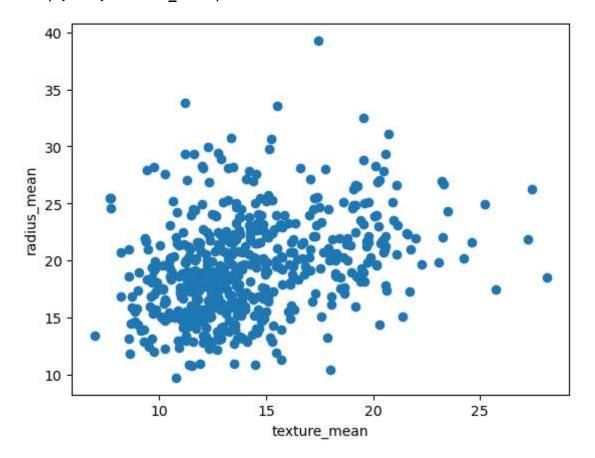
Out[5]:

is	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mea
М	17.99	10.38	122.80	1001.0	0.11840	0.2776
М	20.57	17.77	132.90	1326.0	0.08474	0.0786
М	19.69	21.25	130.00	1203.0	0.10960	0.1599
M	11.42	20.38	77.58	386.1	0.14250	0.2839
M	20.29	14.34	135.10	1297.0	0.10030	0.1328
		•••				
М	21.56	22.39	142.00	1479.0	0.11100	0.1159
М	20.13	28.25	131.20	1261.0	0.09780	0.1034
М	16.60	28.08	108.30	858.1	0.08455	0.1023
М	20.60	29.33	140.10	1265.0	0.11780	0.2770
В	7.76	24.54	47.92	181.0	0.05263	0.0436

Data visualization

```
In [6]: plt.scatter(df["radius_mean"],df["texture_mean"])
    plt.xlabel("texture_mean")
    plt.ylabel("radius_mean")
```

Out[6]: Text(0, 0.5, 'radius_mean')



```
In [7]: from sklearn.cluster import KMeans
km=KMeans()
km
```

Out[7]:

▼ KMeans KMeans()

```
In [8]: y_predicted=km.fit_predict(df[["texture_mean","radius_mean"]])
y_predicted
```

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:87
0: 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 warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:138 2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

```
Out[8]: array([4, 6, 6, 1, 6, 4, 0, 5, 5, 5, 0, 7, 5, 5, 2, 0, 0, 6, 4, 4, 3,
               4, 6, 0, 4, 5, 0, 5, 4, 7, 1, 7, 7, 0, 0, 5, 1, 5, 5, 5, 5, 7, 1,
               5, 0, 1, 1, 3, 5, 5, 4, 1, 0, 5, 1, 6, 5, 1, 3, 3, 1, 5, 3, 5, 5,
               1, 1, 1, 4, 6, 3, 7, 4, 1, 0, 3, 0, 7, 1, 1, 4, 7, 7, 3, 0, 5, 7,
               5, 4, 5, 5, 4, 1, 0, 7, 1, 1, 3, 0, 5, 3, 1, 1, 1, 4, 1, 1,
               1, 5, 0, 1, 3, 5, 3, 4, 5, 0, 3, 0, 6, 4, 4, 4, 5, 0, 4, 7, 3, 0,
               0, 4, 0, 5, 1, 3, 4, 3, 3, 0, 1, 4, 3, 3, 1, 0, 4, 1,
               4, 1, 0, 0, 3, 3, 1, 0, 6, 5, 6, 0, 3, 0, 7, 4, 3, 1, 4, 3, 3, 3,
                     5, 3, 6, 7, 0, 3, 5, 3, 0, 1, 1, 4, 5, 5, 1, 2,
               6, 0, 1, 0, 7, 5, 1, 4, 1, 0, 5, 4, 6, 1, 6, 7, 5, 4, 1, 1,
               4, 4, 1, 0, 4, 4, 3, 4, 5, 5, 0, 2, 2, 7, 3, 5, 7, 6,
                          1, 4, 5, 3, 7, 1, 6, 0, 6, 4, 7, 4, 5, 2, 7,
               0, 7, 1, 5, 4, 1, 4, 3, 6, 3, 7, 1, 3, 0, 1, 4, 7, 3, 6,
                        1, 0, 0, 4, 1, 3, 4, 3, 1, 1, 5, 6, 1, 7, 1, 1,
               4, 4, 1, 4, 3, 3, 1, 1, 3, 0, 1, 1, 3, 6, 3, 6, 3, 1,
               4, 1, 1, 3, 1, 0, 4, 0, 1, 7, 4, 1, 3, 6, 3, 3, 1, 4,
                                                                     3,
               6, 5, 3, 1, 1, 4, 3, 1, 1, 5, 1, 0, 4, 6, 7, 1, 6, 6, 5, 4,
                          4, 1, 3, 3, 5, 1, 4, 5, 3, 4, 3, 7, 3, 1,
                                                                        6,
                  1, 3, 1,
                          0, 3, 1, 4, 3, 1, 4, 5, 0, 1, 1, 1, 1, 5, 2,
               3, 5, 1, 4, 3, 1, 1, 1, 3, 5, 1, 1, 5, 1, 6, 6, 4, 1, 1, 4, 1, 4,
               1, 7, 4, 1, 0, 5, 7, 4, 0, 6, 5, 7, 2, 4, 1, 2, 2, 5,
               2, 1, 1, 1, 5, 1, 7, 1, 1, 2, 4, 2, 3, 4, 5, 4, 3, 0, 1, 1, 4, 1,
               4, 4, 4, 6, 3, 0, 5, 4, 0, 3, 5, 0, 1, 1, 0, 6, 4, 5, 4, 6, 3, 3,
               1, 1, 4, 5, 3, 4, 5, 4, 0, 1, 0, 6, 1, 4, 3, 6, 1, 1, 3, 3, 1, 3,
               4, 3, 1, 1, 4, 6, 1, 6, 5, 5, 5, 5, 3, 5, 5, 2, 5, 5, 3, 1, 1, 5,
               5, 5, 2, 5, 2, 2, 1, 2, 5, 5, 2, 2, 2, 7, 6, 7, 2, 7, 5])
```

```
In [9]: df["cluster"]=y_predicted
df.head()
```

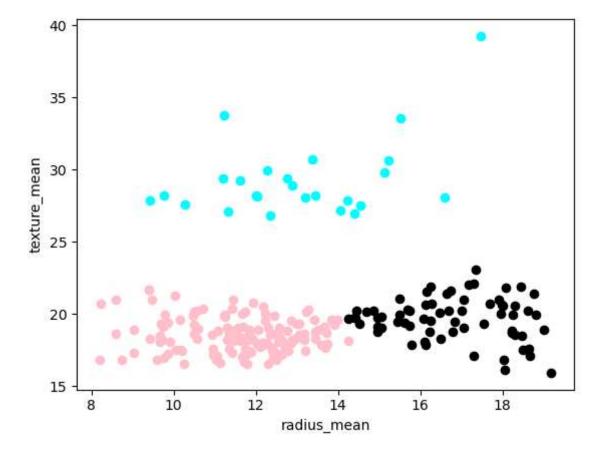
Out[9]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_rr
0	842302	М	17.99	10.38	122.80	1001.0	0.1′
1	842517	М	20.57	17.77	132.90	1326.0	30.0
2	84300903	М	19.69	21.25	130.00	1203.0	0.10
3	84348301	М	11.42	20.38	77.58	386.1	0.14
4	84358402	М	20.29	14.34	135.10	1297.0	0.10

5 rows × 34 columns

```
In [10]: df1=df[df.cluster==0]
    df2=df[df.cluster==1]
    df3=df[df.cluster==2]
    plt.scatter(df1["radius_mean"],df1["texture_mean"],color="black")
    plt.scatter(df2["radius_mean"],df2["texture_mean"],color="pink")
    plt.scatter(df3["radius_mean"],df3["texture_mean"],color="cyan")
    plt.xlabel("radius_mean")
    plt.ylabel("texture_mean")
```

Out[10]: Text(0, 0.5, 'texture_mean')



Out[11]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_rr
0	842302	М	17.99	0.022658	122.80	1001.0	0.1′
1	842517	М	20.57	0.272574	132.90	1326.0	30.0
2	84300903	М	19.69	0.390260	130.00	1203.0	0.10
3	84348301	М	11.42	0.360839	77.58	386.1	0.14
4	84358402	М	20.29	0.156578	135.10	1297.0	0.10

5 rows × 34 columns

In [12]: scaler.fit(df[["radius_mean"]])
 df["radius_mean"]=scaler.transform(df[["radius_mean"]])
 df.head()

Out[12]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_rr
0	842302	М	0.521037	0.022658	122.80	1001.0	0.1′
1	842517	М	0.643144	0.272574	132.90	1326.0	30.0
2	84300903	М	0.601496	0.390260	130.00	1203.0	0.10
3	84348301	М	0.210090	0.360839	77.58	386.1	0.14
4	84358402	М	0.629893	0.156578	135.10	1297.0	0.10

5 rows × 34 columns

→

```
In [13]: y_predicted=km.fit_predict(df[["radius_mean","texture_mean"]])
y_predicted
```

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:87
0: 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 warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:138 2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

```
Out[13]: array([3, 1, 1, 0, 1, 3, 1, 2, 2, 4, 2, 3, 7, 2, 2, 4, 2, 2, 1, 3, 3, 5,
                3, 6, 2, 1, 2, 1, 2, 1, 7, 0, 7, 7, 3, 2, 2, 0, 4, 2, 2, 0, 7, 2,
                2, 1, 5, 0, 5, 2, 0, 3, 0, 1, 2, 0, 1, 2, 0, 5, 5, 0, 2, 5, 4, 2,
                0, 0, 0, 3, 1, 5, 7, 3, 3, 2, 3, 1, 7, 0, 0, 3, 6, 7, 5, 1, 2, 7,
                2, 3, 2, 2, 3, 0, 2, 7, 0, 0, 5, 2, 4, 5, 0, 0, 0, 3, 0, 0,
                0, 0, 2, 0, 5, 0, 5, 3, 2, 1, 5, 1, 6, 3, 3, 3, 4, 1, 3, 7, 5, 2,
                2, 3, 1, 2, 0, 5, 3, 5, 5, 2, 0, 3, 5, 5, 0, 2, 3, 3, 2, 0,
                3, 0, 1, 1, 5, 5, 0, 1, 1, 2, 6, 2, 5, 1, 7, 3, 5, 2, 3, 5, 5, 5,
                0, 2, 2, 3, 6, 7, 2, 5, 2, 5, 1, 0, 0, 3, 2, 2, 0, 4, 2,
                1, 2, 0, 1, 6, 2, 0, 3, 0, 1, 2, 3, 1, 0, 6, 7, 2, 3, 0, 0, 1, 7,
                3, 3, 0, 2, 3, 3, 5, 3, 4, 2, 1, 4, 4, 7, 5, 2, 6, 1, 4,
                0, 2, 7, 0, 3, 3, 4, 5, 7, 0, 1, 1, 1, 3, 7, 3, 2, 4, 7,
                1, 7, 0, 2, 3, 0, 3, 5, 6, 5, 7, 0, 5, 1, 3, 3, 7, 5, 1, 2,
                         0, 2, 2, 3, 0, 3, 3, 5, 0, 3, 0, 1, 0, 7, 0,
                3, 3, 0, 3, 3, 5, 0, 0, 5, 1, 0, 0, 5, 1, 3, 1, 5,
                3, 0, 0, 5, 0, 1, 3, 1, 0, 6, 3, 5, 5, 1, 5, 5, 0, 3,
                6, 4, 5, 0, 0, 3, 5, 0, 0, 2, 0, 1, 3, 1, 7, 0, 1, 6, 2, 3,
                3, 3, 0, 4, 3, 0, 5, 5, 2, 0, 3, 2, 5, 3, 5, 7, 5, 5,
                2, 0, 5, 0, 1, 5, 0, 3, 3, 0, 3, 2, 1, 0, 0, 0, 0, 2, 4,
                5, 0, 0, 3, 5, 2, 0, 0, 5, 0, 0, 0, 2, 0, 1, 1, 3, 2,
                0, 7, 3, 0, 1, 4, 7, 3, 2, 1,
                                              0, 7, 4, 3, 0, 4, 4, 4,
                4, 0, 0, 2, 2, 0, 7, 0, 0, 4, 3, 4, 5, 3, 2, 3, 5, 2, 0, 2, 3, 3,
                3, 3, 3, 1, 5, 1, 2, 3, 1, 5, 2, 2, 0, 0, 1, 1, 3, 4, 3, 6, 5, 5,
                0, 0, 3, 2, 5, 3, 2, 3, 2, 0, 1, 1, 0, 3, 5, 6, 0, 2, 5, 5, 2, 5,
                3, 5, 0, 0, 3, 1, 0, 1, 2, 4, 4, 4, 5, 4, 4, 4, 2, 2, 5, 5, 0, 4,
                0, 0, 4, 0, 4, 4, 0, 4, 2, 4, 4, 4, 4, 7, 6, 7, 7, 7, 4])
```

```
In [14]: df["New Cluster"]=y_predicted
    df.head()
```

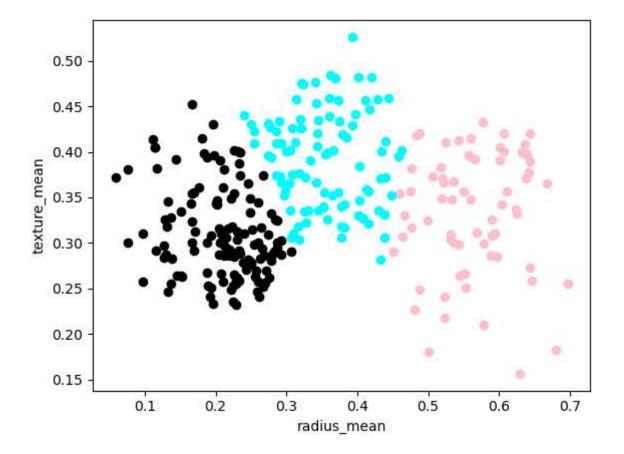
Out[14]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_rr
0	842302	М	0.521037	0.022658	122.80	1001.0	0.1′
1	842517	М	0.643144	0.272574	132.90	1326.0	30.0
2	84300903	М	0.601496	0.390260	130.00	1203.0	0.10
3	84348301	М	0.210090	0.360839	77.58	386.1	0.14
4	84358402	М	0.629893	0.156578	135.10	1297.0	0.10

5 rows × 35 columns

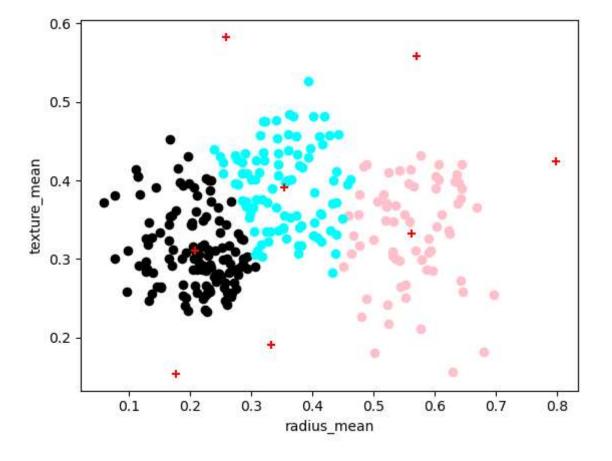
```
In [15]: df1=df[df["New Cluster"]==0]
    df2=df[df["New Cluster"]==1]
    df3=df[df["New Cluster"]==2]
    plt.scatter(df1["radius_mean"],df1["texture_mean"],color="black")
    plt.scatter(df2["radius_mean"],df2["texture_mean"],color="pink")
    plt.scatter(df3["radius_mean"],df3["texture_mean"],color="cyan")
    plt.xlabel("radius_mean")
    plt.ylabel("texture_mean")
```

Out[15]: Text(0, 0.5, 'texture_mean')



```
In [16]:
         km.cluster_centers_
Out[16]: array([[0.20878924, 0.31058452],
                [0.56287997, 0.33184226],
                [0.3534653, 0.39091896],
                [0.3331624, 0.18999839],
                [0.2590623, 0.58293879],
                [0.17652977, 0.15382448],
                [0.79840767, 0.42469846],
                [0.57132058, 0.55893025]])
In [17]: | df1=df[df["New Cluster"]==0]
         df2=df[df["New Cluster"]==1]
         df3=df[df["New Cluster"]==2]
         plt.scatter(df1["radius_mean"],df1["texture_mean"],color="black")
         plt.scatter(df2["radius_mean"],df2["texture_mean"],color="pink")
         plt.scatter(df3["radius_mean"],df3["texture_mean"],color="cyan")
         plt.scatter(km.cluster_centers_[:,0],km.cluster_centers_[:,1],color="red",mark
         plt.xlabel("radius mean")
         plt.ylabel("texture_mean")
```

Out[17]: Text(0, 0.5, 'texture_mean')



```
In [18]: k_rng=range(1,10)
sse=[]
```

Elbow method

```
breast cancer - Jupyter Notebook
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:87
0: 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
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:138
2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh
en there are less chunks than available threads. You can avoid it by setting
the environment variable OMP_NUM_THREADS=3.
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:87
0: 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
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\_kmeans.py:138
2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh
en there are less chunks than available threads. You can avoid it by setting
the environment variable OMP NUM THREADS=3.
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:87
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the environment variable OMP NUM THREADS=3.
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\_kmeans.py:87
0: 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
  warnings.warn(
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:138
```

2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:87
0: 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

warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:138 2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:87
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C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:138 2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

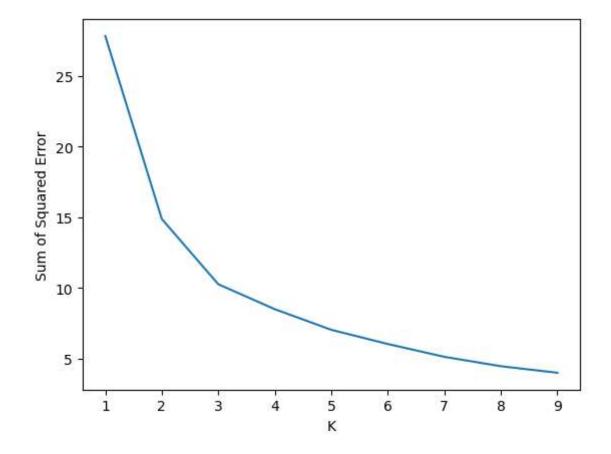
C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:87
0: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning warnings.warn(

C:\Users\Y.Saranya\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:138 2: UserWarning: KMeans is known to have a memory leak on Windows with MKL, wh en there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=3.

warnings.warn(

[27.817507595043075, 14.872296449956036, 10.252751496105198, 8.48893568609588 9, 7.030611949327338, 6.025284267841283, 5.118942965866768, 4.45468255928802 7, 3.991752934887513]

Out[20]: Text(0, 0.5, 'Sum of Squared Error')



CONCLUSION:- BASED ON THE ABOVE PROGRAM THE DATA HAS DIVIDED INTO SEVARAL THE DATA HAS DIVIDED INTO SEVARAL

In []:	