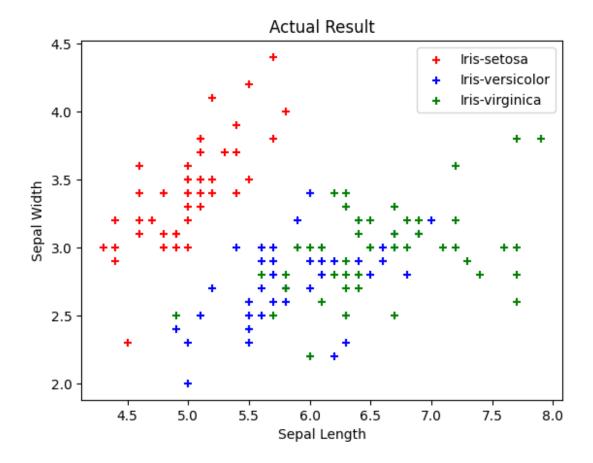
ML Assignment 4: Iris Dataset

```
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import silhouette_score, calinski_harabasz_score,
davies_bouldin_score
!gdown 17bW5DYVUVShliNePIuBfawk_y7F0q07F # iris dataset
```

Clustering in Iris Dataset

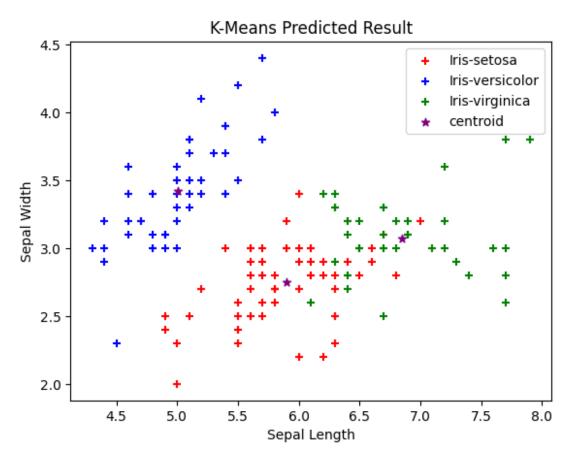
```
df iris = pd.read csv("iris.csv", names=['sepal length',
'sepal width', 'petal length', 'petal width', 'species'])
X = df_iris.drop('species', axis=1)
y = df iris.species
# Actual Clustering Result
newDf0 = df iris[df iris.species=="Iris-setosa"]
newDf1 = df iris[df iris.species=="Iris-versicolor"]
newDf2 = df iris[df iris.species=="Iris-virginica"]
plt.title("Actual Result")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.scatter(newDf0.sepal length, newDf0.sepal width, color="red",
marker="+", label="Iris-setosa")
plt.scatter(newDf1.sepal length, newDf1.sepal width, color="blue",
marker="+", label="Iris-versicolor")
plt.scatter(newDf2.sepal length, newDf2.sepal width, color="green",
marker="+", label="Iris-virginica")
plt.legend()
<matplotlib.legend.Legend at 0x7f3c26c6d3f0>
```



Partition Based: K-means Clustering in Iris Dataset

```
# Clustering using K-means algorithm
from sklearn.cluster import KMeans
km = KMeans(n clusters=3, n init=10)
y predicted = km.fit predict(X)
newDf = df iris
newDf["cluster"] = y predicted
newDf0 = newDf[newDf.cluster==0]
newDf1 = newDf[newDf.cluster==1]
newDf2 = newDf[newDf.cluster==2]
plt.title("K-Means Predicted Result")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.scatter(newDf0.sepal_length, newDf0.sepal_width, color="red",
marker="+", label="Iris-setosa")
plt.scatter(newDf1.sepal length, newDf1.sepal width, color="blue",
marker="+", label="Iris-versicolor")
plt.scatter(newDf2.sepal length, newDf2.sepal width, color="green",
marker="+", label="Iris-virginica")
plt.scatter(km.cluster centers [:,0], km.cluster centers [:,1],
```

```
color="purple", marker="*", label="centroid")
plt.legend()
<matplotlib.legend.Legend at 0x7f3c26651bd0>
```

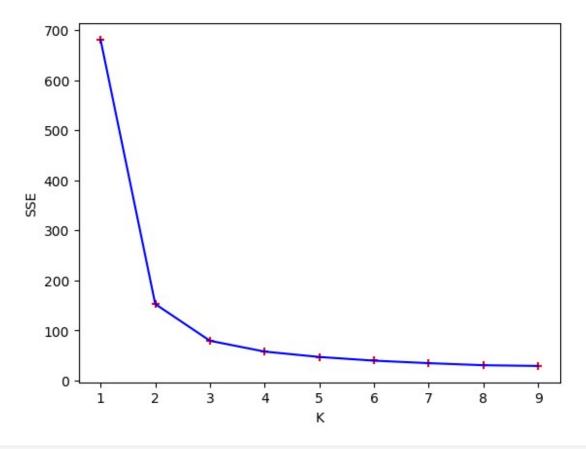


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7f3c26248a90>]
```



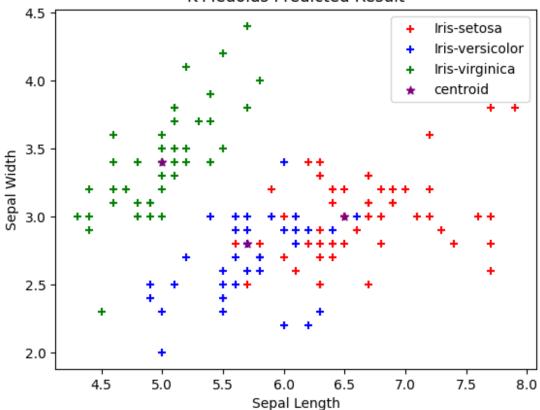
```
# Evaluating Metrics
silhouette_result = silhouette_score(X, km.labels_)
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski_result)
davies_result = davies_bouldin_score(X, km.labels_)
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion scores)
separation = SSB/N
```

```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
Silhouette Score: 0.34597762034129553
Calinski Harabasz Score: 401.8511977911363
Davies Bouldin Score: 1.0253347541939601
Cohesion Score: 0.04767664257307373
Separation Score: 0.08704928634167765
```

Partition Based: K-medoids Clustering in Iris Dataset

```
!pip install scikit-learn-extra
# Clustering using K-medoids algorithm
from sklearn extra.cluster import KMedoids
km = KMedoids(n clusters=3)
y predicted = km.fit predict(X)
newDf = df iris
newDf["cluster"] = y predicted
newDf0 = newDf[newDf.cluster==0]
newDf1 = newDf[newDf.cluster==1]
newDf2 = newDf[newDf.cluster==2]
plt.title("K-Medoids Predicted Result")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.scatter(newDf0.sepal length, newDf0.sepal width, color="red",
marker="+", label="Iris-setosa")
plt.scatter(newDf1.sepal length, newDf1.sepal width, color="blue",
marker="+", label="Iris-versicolor")
plt.scatter(newDf2.sepal length, newDf2.sepal_width, color="green",
marker="+", label="Iris-virginica")
plt.scatter(km.cluster centers [:,0], km.cluster centers [:,1],
color="purple", marker="*", label="centroid")
plt.legend()
<matplotlib.legend.Legend at 0x7f3c262b73d0>
```

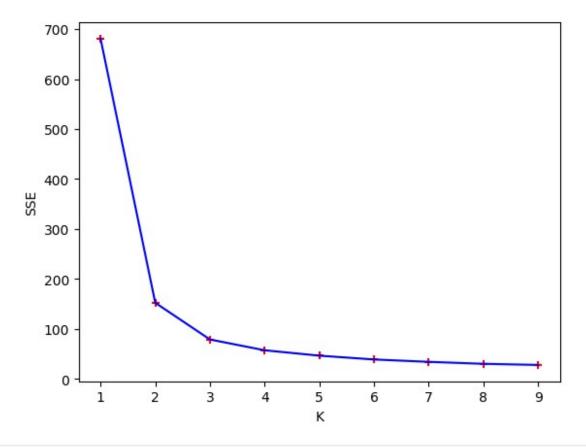
K-Medoids Predicted Result



```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3
[<matplotlib.lines.Line2D at 0x7f3c261579a0>]
```



```
# Evaluating Metrics
silhouette_result = silhouette_score(X, km.labels_)
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski_result)
davies_result = davies_bouldin_score(X, km.labels_)
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion scores)
separation = SSB/N
```

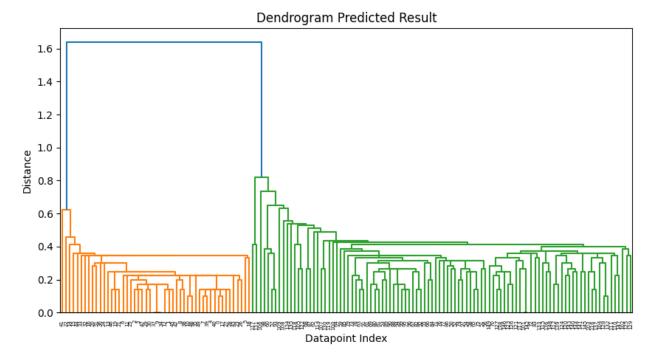
```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
Silhouette Score: 0.3435514923440904
Calinski Harabasz Score: 411.2774030706613
Davies Bouldin Score: 0.973553876855317
Cohesion Score: 0.04662882885901002
Separation Score: 0.06856001899335233
```

Hierarchical: Dendrogram Clustering in Iris Dataset

```
# Clustering using Dendrogram Clustering algorithm
from scipy.cluster.hierarchy import dendrogram, linkage, fcluster
Z = linkage(X, method='single')

# Create and plot the dendrogram
plt.figure(figsize=(10, 5))
dn = dendrogram(Z)

plt.title('Dendrogram Predicted Result')
plt.xlabel('Datapoint Index')
plt.ylabel('Distance')
plt.show()
```



```
# Evaluating Metrics
labels = fcluster(Z, 3, criterion='maxclust')
```

```
silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

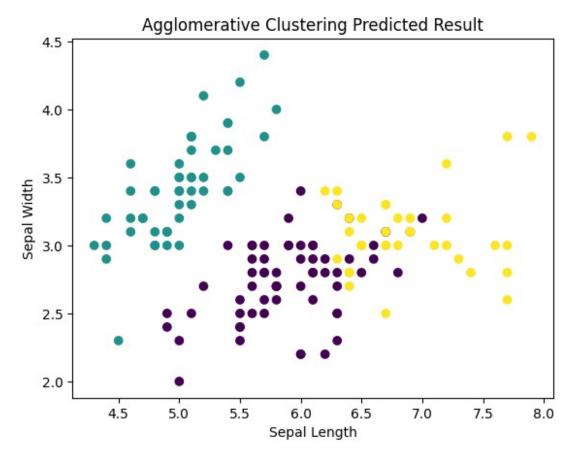
davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)

Silhouette Score: 0.5118387098922373
Calinski Harabasz Score: 277.4926776474616
Davies Bouldin Score: 0.4474384341989626
```

Hierarchical: AGNES Clustering in Iris Dataset

```
# Clustering using AGNES Clustering algorithm
from sklearn.cluster import AgglomerativeClustering
agg_cluster = AgglomerativeClustering(n_clusters=3, linkage='ward')
agg_cluster.fit(X)

plt.scatter(df_iris.sepal_length, df_iris.sepal_width,
c=agg_cluster.labels_, cmap='viridis')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.title('Agglomerative Clustering Predicted Result')
plt.show()
```



```
# Evaluating Metrics
labels = fcluster(Z, 3, criterion='maxclust')

from sklearn.metrics import silhouette_score
silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

from sklearn.metrics import calinski_harabasz_score
calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

from sklearn.metrics import davies_bouldin_score
davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)

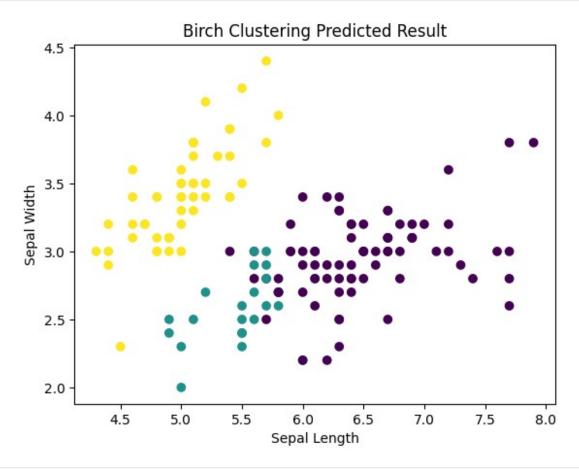
Silhouette Score: 0.5118387098922373
Calinski Harabasz Score: 277.4926776474616
Davies Bouldin Score: 0.4474384341989626
```

Hierarchical: BIRCH Clustering in Iris Dataset

Clustering using BIRCH Clustering algorithm from sklearn.cluster import Birch

```
birch_cluster = Birch(n_clusters=3)
birch_cluster.fit(X)

plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.scatter(df_iris.sepal_length, df_iris.sepal_width,
c=birch_cluster.labels_, cmap='viridis')
plt.title('Birch Clustering Predicted Result')
plt.show()
```



```
# Evaluating Metrics
labels = birch_cluster.fit_predict(X)

silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)
```

Silhouette Score: 0.5016992571068448

Calinski Harabasz Score: 457.54177598067594 Davies Bouldin Score: 0.6262973013286385

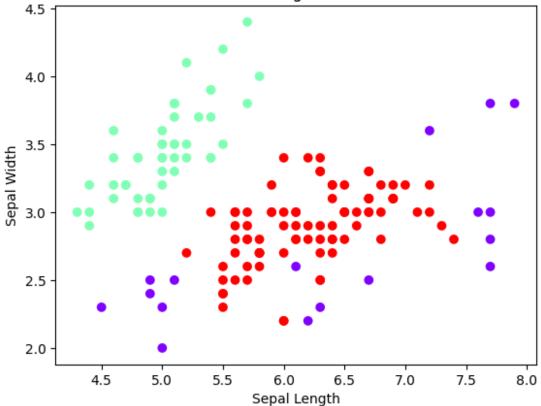
Density Based: DBSCAN Clustering in Iris Dataset

```
# Clustering using DBSCAN Clustering algorithm
from sklearn.cluster import DBSCAN

dbscan = DBSCAN(eps=0.5, algorithm='auto', metric='euclidean')
y = dbscan.fit_predict(X)

plt.scatter(df_iris.sepal_length, df_iris.sepal_width,
c=dbscan.labels_, cmap='rainbow')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.title('DBScan Clustering Predicted Result')
plt.show()
```

DBScan Clustering Predicted Result



```
# Evaluating Metrics
silhouette_result = silhouette_score(X, dbscan.labels_)
print("Silhouette Score: ", silhouette_result)
```

```
calinski_result = calinski_harabasz_score(X, dbscan.labels_)
print("Calinski Harabasz Score: ", calinski_result)

davies_result = davies_bouldin_score(X, dbscan.labels_)
print("Davies Bouldin Score: ", davies_result)

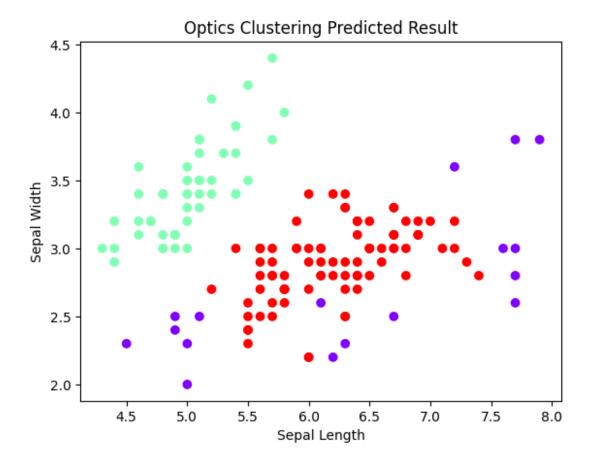
Silhouette Score: 0.485842354600955
Calinski Harabasz Score: 219.87022703461665
Davies Bouldin Score: 7.222826995273629
```

Density Based: Optics Clustering in Iris Dataset

```
# Clustering using Optics Clustering algorithm
from sklearn.cluster import OPTICS

optics_cluster = OPTICS(min_samples=5, xi=0.05,
cluster_method='dbscan')
optics_cluster.fit(X)

plt.scatter(df_iris.sepal_length, df_iris.sepal_width,
c=dbscan.labels_, cmap='rainbow')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.title('Optics Clustering Predicted Result')
plt.show()
```

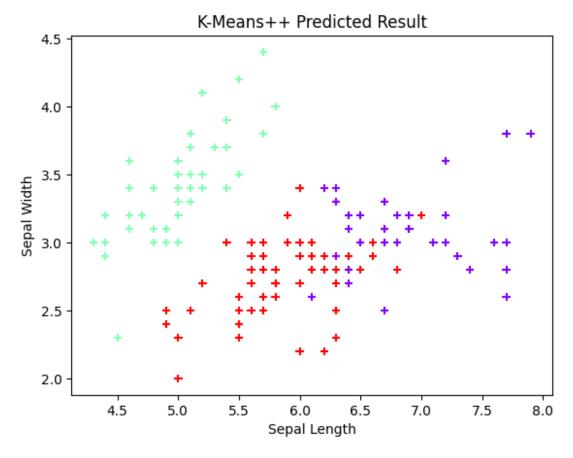


K-means++ Clustering in Iris Dataset

```
# Clustering using K-means++ algorithm
from sklearn.cluster import KMeans
km = KMeans(init='k-means++', n_clusters=3, n_init=10, max_iter=300,
random_state=42)
km = KMeans(n_clusters=3, n_init=10)

y_predicted = km.fit_predict(X)

plt.title("K-Means++ Predicted Result")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.scatter(df_iris.sepal_length, df_iris.sepal_width, c=km.labels_,
cmap='rainbow', marker="+")
plt.show()
```

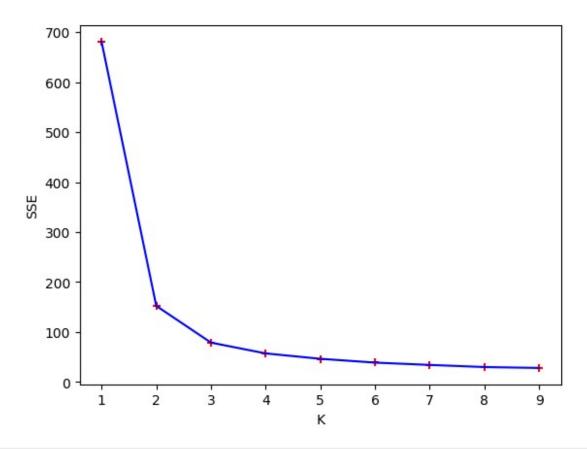


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7f3c25f7cd60>]
```



```
# Evaluating Metrics
silhouette_result = silhouette_score(X, km.labels_)
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski_result)
davies_result = davies_bouldin_score(X, km.labels_)
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion scores)
separation = SSB/N
```

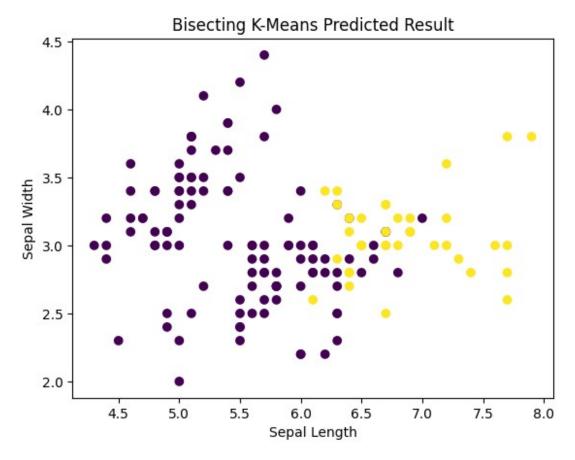
```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")

Silhouette Score: 0.35859288904946124
Calinski Harabasz Score: 407.19542232102003
Davies Bouldin Score: 0.958208320286649

Cohesion Score: 0.04707687224812225
Separation Score: 0.09227952602952604
```

Bisecting K-means Clustering in Iris Dataset

```
# Clustering using Bisecting K-means algorithm
from sklearn.cluster import KMeans
km = KMeans(n clusters=1, n init=10, random state=0).fit(X)
K=3
for i in range(K-1):
    largest cluster = np.argmax(np.bincount(km.labels ))
    largest_cluster_mask = (km.labels_ == largest_cluster)
    X split = X[largest cluster mask]
    km.labels [largest cluster mask] = KMeans(n clusters=2, n init=10,
random state=0).fit(X split).labels
plt.title("Bisecting K-Means Predicted Result")
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.scatter(df iris.sepal length, df iris.sepal width, c=km.labels ,
cmap='viridis')
plt.show()
```

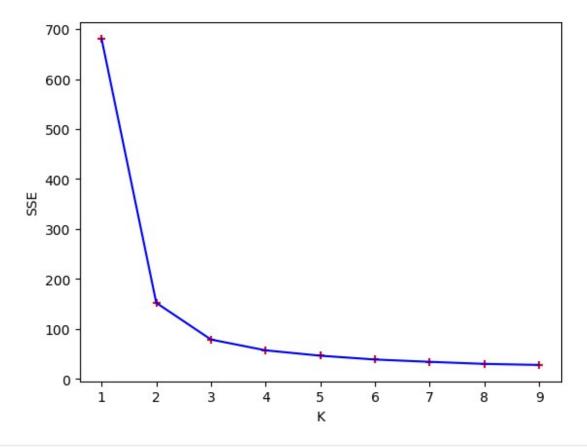


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7f3c2665c7c0>]
```



```
# Evaluating Metrics
silhouette_result = silhouette_score(X, km.labels_)
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski_result)
davies_result = davies_bouldin_score(X, km.labels_)
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion scores)
separation = SSB/N
```

```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
```

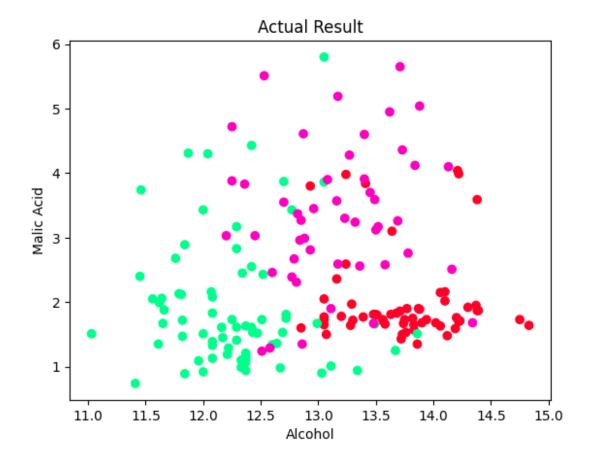
Silhouette Score: 0.34093973210740597

Calinski Harabasz Score: 409.94078898037156 Davies Bouldin Score: 1.0060550389051572

Cohesion Score: 0.046774595314776485 Separation Score: 0.06472950617283951

ML Assignment 4: Wine Dataset

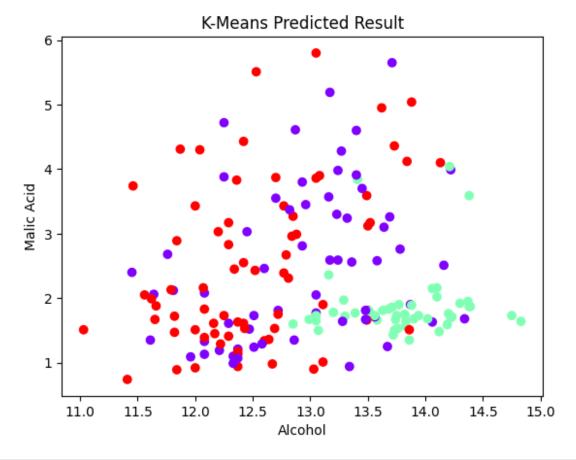
```
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import silhouette score, calinski harabasz score,
davies bouldin score
!gdown 1w57c07HLl33Dcwxae j41Gm8SiqJYpih
                                           # wine dataset
df = pd.read csv("wine.csv",
names=['class',"Alcohol","Malicacid","Ash","Alcalinity of ash","Magnes
ium", "Total phenols", "Flavanoids", "Nonflavanoid phenols", "Proanthocyan
ins", "Color intensity", "Hue", "OD280 OD315 of diluted wines", "Proline"]
X = df.drop('class',axis=1)
v = df["class"]
df.head()
          Alcohol Malicacid
                                     Alcalinity of ash
   class
                              Ash
                                                        Magnesium \
0
       1
            14.23
                        1.71 2.43
                                                  15.6
                                                              127
       1
            13.20
                                                  11.2
                                                              100
1
                        1.78 2.14
2
       1
            13.16
                        2.36 2.67
                                                  18.6
                                                              101
3
       1
            14.37
                        1.95
                              2.50
                                                  16.8
                                                              113
4
       1
            13.24
                        2.59 2.87
                                                  21.0
                                                              118
   Total phenols Flavanoids
                              Nonflavanoid phenols Proanthocyanins \
0
            2.80
                        3.06
                                               0.28
                                                                2.29
1
            2.65
                        2.76
                                               0.26
                                                                1.28
2
            2.80
                        3.24
                                               0.30
                                                                2.81
3
            3.85
                        3.49
                                               0.24
                                                                2.18
4
            2.80
                        2.69
                                               0.39
                                                                1.82
                          OD280 OD315 of diluted_wines
   Color intensity
                     Hue
                                                         Proline
0
              5.64
                   1.04
                                                   3.92
                                                            1065
1
              4.38 1.05
                                                   3.40
                                                            1050
2
              5.68
                   1.03
                                                   3.17
                                                            1185
3
              7.80
                    0.86
                                                   3.45
                                                            1480
              4.32 1.04
                                                   2.93
                                                             735
# Actual Clustering Result
plt.title("Actual Result")
plt.xlabel('Alcohol')
plt.ylabel('Malic Acid')
plt.scatter(df.Alcohol, df.Malicacid, c=df["class"],
cmap='gist rainbow')
<matplotlib.collections.PathCollection at 0x7a82044b9960>
```



Partition Based: K-means Clustering in Wine Dataset

```
# Clustering using K-means algorithm
from sklearn.cluster import KMeans
km = KMeans(init="random", n_clusters=3, n_init=10, max_iter=300,
random_state=42)
y_predicted = km.fit_predict(X)

plt.title("K-Means Predicted Result")
plt.xlabel("Alcohol")
plt.ylabel("Malic Acid")
plt.scatter(df.Alcohol, df.Malicacid, c=km.labels_, cmap='rainbow')
plt.show()
```

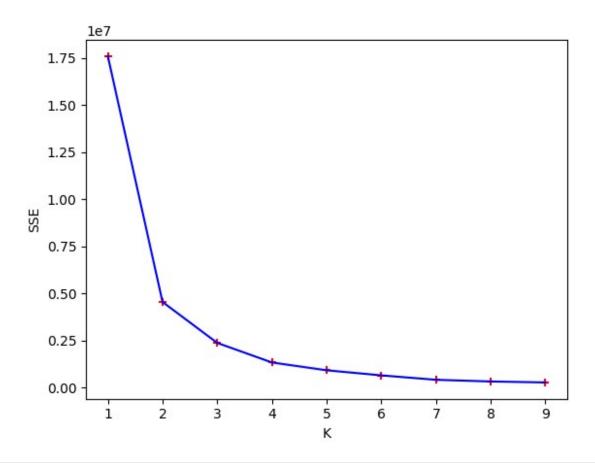


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7a8204383100>]
```



```
# Evaluating Metrics
silhouette result = silhouette score(X, km.labels )
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski result)
davies result = davies bouldin score(X, km.labels )
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion_scores)
separation = SSB/N
```

```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
Silhouette Score: 0.5307235924738344
Calinski Harabasz Score: 1350.458318826902
Davies Bouldin Score: 0.5163732495928284
Cohesion Score: 117.09374643108792
Separation Score: 607.9526439562346
```

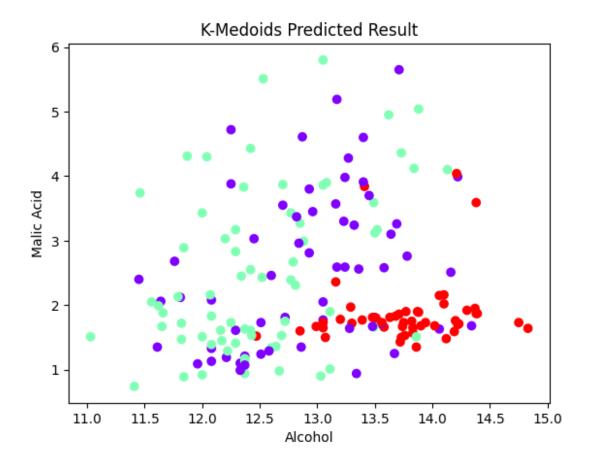
Partition Based: K-medoids Clustering in Wine Dataset

```
!pip install scikit-learn-extra

# Clustering using K-medoids algorithm
from sklearn_extra.cluster import KMedoids
km = KMedoids(n_clusters=3)

y_predicted = km.fit_predict(X)

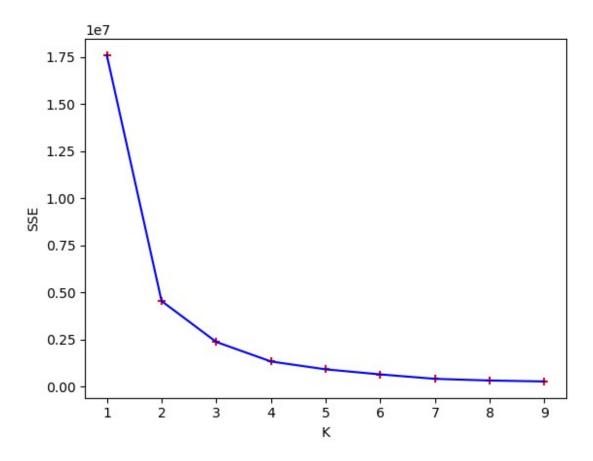
plt.title("K-Medoids Predicted Result")
plt.xlabel("Alcohol")
plt.ylabel("Malic Acid")
plt.scatter(df.Alcohol, df.Malicacid, c=km.labels_, cmap='rainbow')
plt.show()
```



```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3
[<matplotlib.lines.Line2D at 0x7a8254c14490>]
```



```
# Evaluating Metrics
silhouette result = silhouette score(X, km.labels )
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski result)
davies result = davies bouldin score(X, km.labels )
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion_scores)
separation = SSB/N
```

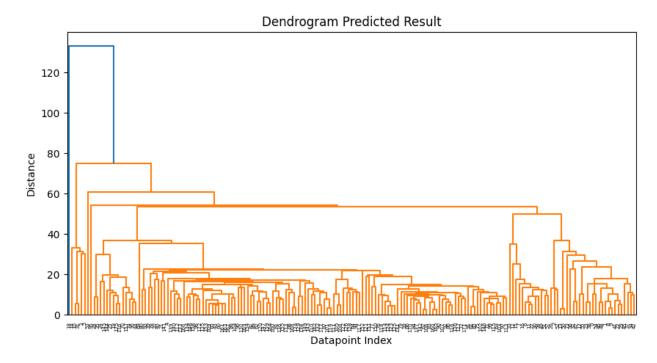
```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
Silhouette Score: 0.5382358200331198
Calinski Harabasz Score: 1340.298246818952
Davies Bouldin Score: 0.5274536247334654
Cohesion Score: 117.96759730604572
Separation Score: 617.91903929374
```

Hierarchical: Dendrogram Clustering in Wine Dataset

```
# Clustering using Dendrogram Clustering algorithm
from scipy.cluster.hierarchy import dendrogram, linkage, fcluster
Z = linkage(X, method='single')

# Create and plot the dendrogram
plt.figure(figsize=(10, 5))
dn = dendrogram(Z)

plt.title('Dendrogram Predicted Result')
plt.xlabel('Datapoint Index')
plt.ylabel('Distance')
plt.show()
```



```
# Evaluating Metrics
labels = fcluster(Z, 3, criterion='maxclust')
from sklearn.metrics import silhouette_score
```

```
silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

from sklearn.metrics import calinski_harabasz_score
calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

from sklearn.metrics import davies_bouldin_score
davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)

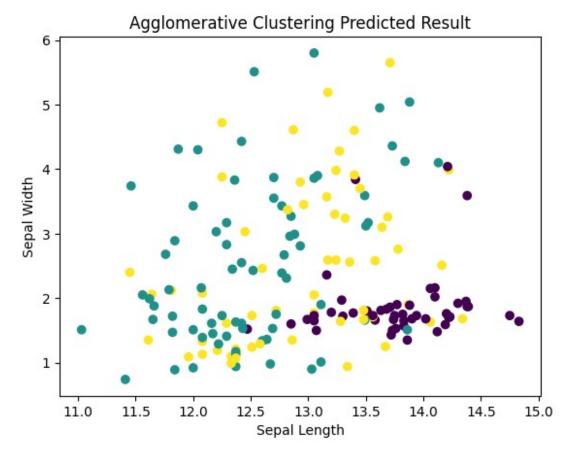
Silhouette Score: 0.4879820335189063
Calinski Harabasz Score: 24.42036238154286
Davies Bouldin Score: 0.30814096183494405
```

Hierarchical: AGNES Clustering in Wine Dataset

```
# Clustering using AGNES Clustering algorithm
from sklearn.cluster import AgglomerativeClustering

agg_cluster = AgglomerativeClustering(n_clusters=3, linkage='ward')
agg_cluster.fit(X)

plt.scatter(df.Alcohol, df.Malicacid, c=agg_cluster.labels_,
cmap='viridis')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.title('Agglomerative Clustering Predicted Result')
plt.show()
```



```
# Evaluating Metrics
labels = fcluster(Z, 3, criterion='maxclust')
silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

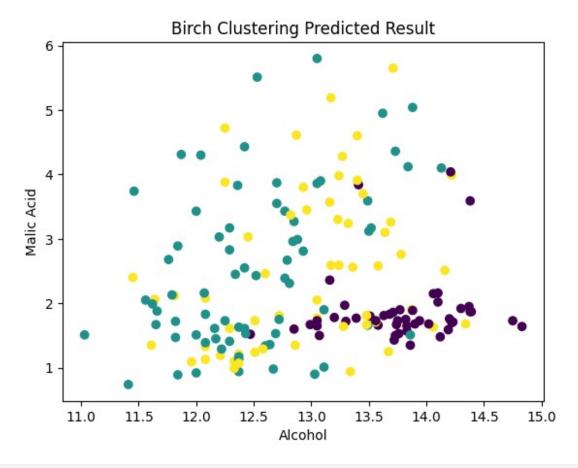
davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)

Silhouette Score: 0.4879820335189063
Calinski Harabasz Score: 24.42036238154286
Davies Bouldin Score: 0.30814096183494405
```

Hierarchical: BIRCH Clustering in Wine Dataset

```
# Clustering using BIRCH Clustering algorithm
from sklearn.cluster import Birch
birch_cluster = Birch(n_clusters=3)
birch_cluster.fit(X)
```

```
plt.xlabel('Alcohol')
plt.ylabel('Malic Acid')
plt.scatter(df.Alcohol, df.Malicacid, c=birch_cluster.labels_,
cmap='viridis')
plt.title('Birch Clustering Predicted Result')
plt.show()
```



```
# Evaluating Metrics
labels = birch_cluster.fit_predict(X)

from sklearn.metrics import silhouette_score
silhouette_result = silhouette_score(X, labels)
print("Silhouette Score: ", silhouette_result)

from sklearn.metrics import calinski_harabasz_score
calinski_result = calinski_harabasz_score(X, labels)
print("Calinski Harabasz Score: ", calinski_result)

from sklearn.metrics import davies_bouldin_score
davies_result = davies_bouldin_score(X, labels)
print("Davies Bouldin Score: ", davies_result)
```

Silhouette Score: 0.5644796401732071 Calinski Harabasz Score: 552.851711505718 Davies Bouldin Score: 0.5357343073560251

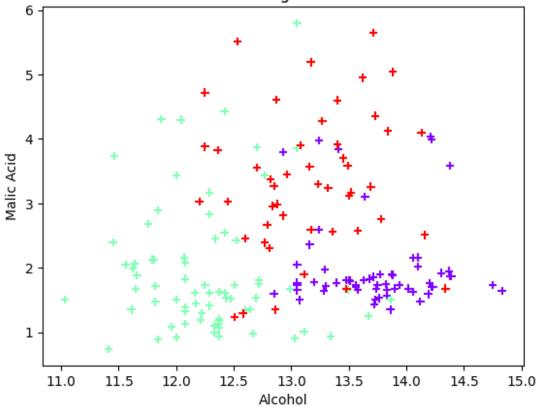
Density Based: DBSCAN Clustering in Wine Dataset

```
# Clustering using DBSCAN Clustering algorithm
from sklearn.cluster import DBSCAN

dbscan = DBSCAN(eps=0.5, algorithm='auto', metric='euclidean')
y = dbscan.fit_predict(X)

plt.title('DBScan Clustering Predicted Result')
plt.xlabel('Alcohol')
plt.ylabel('Malic Acid')
plt.scatter(df.Alcohol, df.Malicacid, c=df["class"], cmap='rainbow',
marker="+")
plt.show()
```

DBScan Clustering Predicted Result

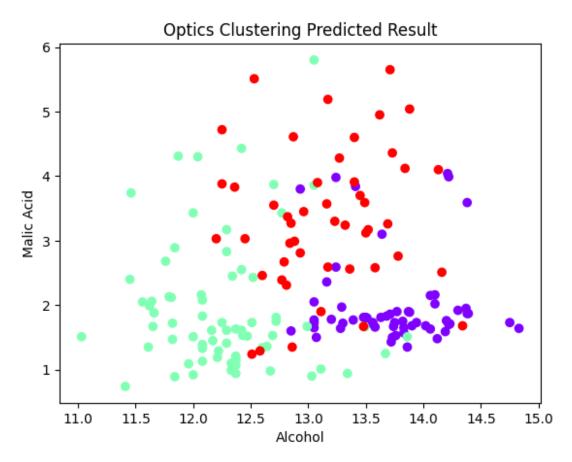


Density Based: Optics Clustering in Wine Dataset

Clustering using Optics Clustering algorithm
from sklearn.cluster import OPTICS

```
optics_cluster = OPTICS(min_samples=5, xi=0.05,
cluster_method='dbscan')
optics_cluster.fit(X)

plt.scatter(df.Alcohol, df.Malicacid, c=df["class"], cmap='rainbow')
plt.xlabel('Alcohol')
plt.ylabel('Malic Acid')
plt.title('Optics Clustering Predicted Result')
plt.show()
```



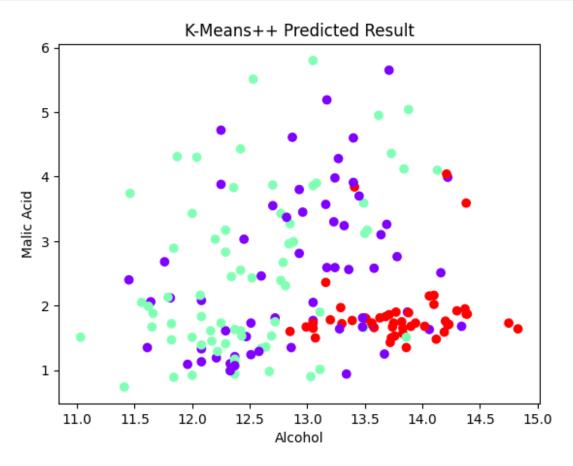
K-means++ Clustering in Wine Dataset

```
# Clustering using K-means++ algorithm
from sklearn.cluster import KMeans
km = KMeans(init='k-means++', n_clusters=3, n_init=10, max_iter=300,
random_state=42)
km = KMeans(n_clusters=3, n_init=10)

y_predicted = km.fit_predict(X)

plt.title("K-Means++ Predicted Result")
plt.xlabel("Alcohol")
```

```
plt.ylabel("Malic Acid")
plt.scatter(df.Alcohol, df.Malicacid, c=km.labels_, cmap='rainbow')
plt.show()
```

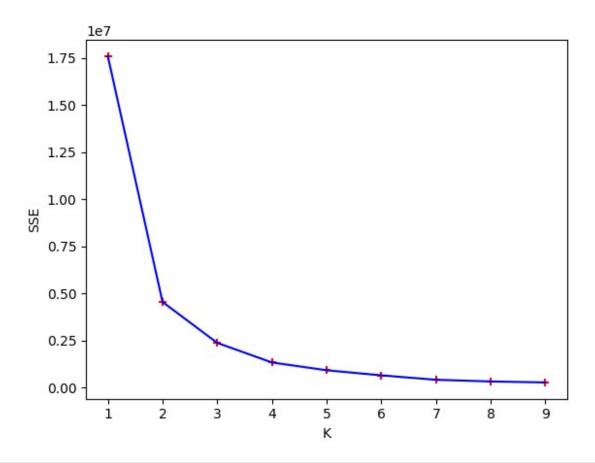


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7a8204295270>]
```



```
# Evaluating Metrics
from sklearn.metrics import silhouette score
silhouette result = silhouette score(X, km.labels )
print("Silhouette Score: ", silhouette result)
from sklearn.metrics import calinski harabasz score
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski result)
from sklearn.metrics import davies bouldin score
davies_result = davies_bouldin_score(X, km.labels_)
print("Davies Bouldin Score: ", davies_result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
```

```
cohesion = np.mean(cohesion_scores)
separation = SSB/N

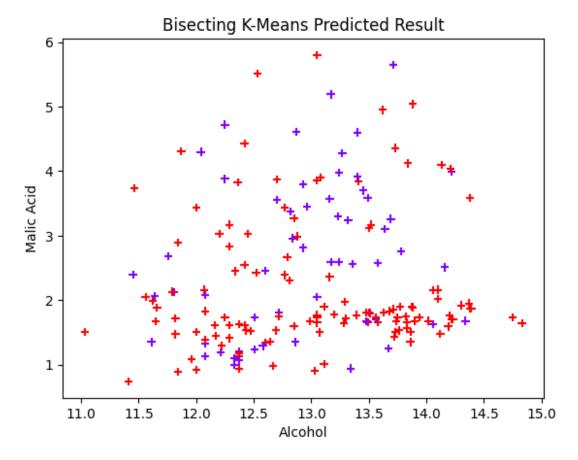
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")

Silhouette Score: 0.5287268772337207
Calinski Harabasz Score: 1354.5755834453266
Davies Bouldin Score: 0.530361984915425

Cohesion Score: 116.74330248636556
Separation Score: 395.5195298071477
```

Bisecting K-means Clustering in Wine Dataset

```
# Clustering using Bisecting K-means algorithm
from sklearn.cluster import KMeans
km = KMeans(n clusters=1, n init=10, random state=0).fit(X)
K=3
for i in range(K-1):
    largest cluster = np.argmax(np.bincount(km.labels ))
    largest_cluster mask = (km.labels_ == largest_cluster)
    X split = X[largest cluster mask]
    km.labels [largest cluster mask] = KMeans(n clusters=2, n init=10,
random state=0).fit(X split).labels
plt.title("Bisecting K-Means Predicted Result")
plt.xlabel("Alcohol")
plt.ylabel("Malic Acid")
plt.scatter(df.Alcohol, df.Malicacid, c=km.labels_, cmap='rainbow',
marker="+")
plt.show()
```

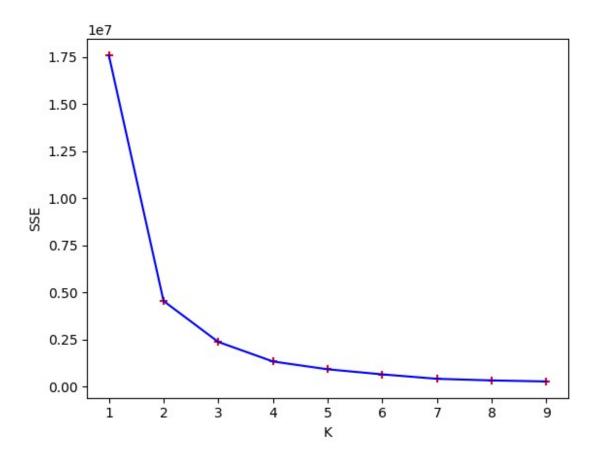


```
# Visualisation of SSE (Sum of Squared Errors) & Elbow Graph:
sse = []
k_range = range(1, 10)

for k in k_range:
    km = KMeans(n_clusters=k, n_init=10)
    km.fit_predict(X)
    sse.append(km.inertia_)

plt.xlabel("K")
plt.ylabel("SSE")
plt.scatter(k_range, sse, color="red", marker="+")
plt.plot(k_range, sse, color="blue")
# We can see here, our elbow is at K=3

[<matplotlib.lines.Line2D at 0x7a82041950f0>]
```



```
# Evaluating Metrics
silhouette result = silhouette score(X, km.labels )
print("Silhouette Score: ", silhouette_result)
calinski result = calinski harabasz score(X, km.labels )
print("Calinski Harabasz Score: ", calinski result)
davies result = davies bouldin score(X, km.labels )
print("Davies Bouldin Score: ", davies result)
# Evaluating Cohesion & Separation
labels = km.labels
centroids = km.cluster centers
SSE = np.sum((X - centroids[labels])**2)
overall centroid = np.mean(X, axis=0)
SSB = np.sum([np.sum((X[labels == i] - centroids[i])**2)) for i in
range(3)])
N = X.shape[0]
cohesion scores = SSE/N
cohesion = np.mean(cohesion_scores)
separation = SSB/N
```

```
print(f"\nCohesion Score: {cohesion}")
print(f"Separation Score: {separation}")
```

Silhouette Score: 0.527999057875864

Calinski Harabasz Score: 1349.5503166007632 Davies Bouldin Score: 0.5215893651849661

Cohesion Score: 117.17131504353662 Separation Score: 435.555516861042

CLUSTERING ALGORITHMS

Type of Algorithm	Algorithm	Dataset	Silhouette Score	Calinski Harabasz Score	Davies Bouldin Score	Cohesion	Separation
Partition Based	K-means	IRIS PLANT DATASET	0.34597762	401.8511978	1.025334754	0.047676643	0.087049286
		WINE DATASET	0.530723592	1350.458319	0.51637325	117.0937464	607.952644
	K-medoids	IRIS PLANT DATASET	0.343551492	411.2774031	0.973553877	0.046628829	0.068560019
		WINE DATASET	0.53823582	1340.298247	0.527453625	117.9675973	617.9190393
Hierarchical	Dendrogram	IRIS PLANT DATASET	0.51183871	277.4926776	0.447438434	-	-
		WINE DATASET	0.487982034	24.42036238	0.308140962	-	-
	AGNES	IRIS PLANT DATASET	0.51183871	277.4926776	0.447438434	-	-
		WINE DATASET	0.487982034	24.42036238	0.308140962	-	-
	BIRCH	IRIS PLANT DATASET	0.501699257	457.541776	0.626297301	-	-
		WINE DATASET	0.56447964	552.8517115	0.535734307	-	-
Density Based	DBSCAN	IRIS PLANT DATASET	0.485842355	219.870227	7.222826995	-	-
		WINE DATASET	-	-	-	-	-
	OPTICS	IRIS PLANT DATASET	-	-	-	-	-
		WINE DATASET	-	-	-	-	-
Additional	K-means++	IRIS PLANT DATASET	0.358592889	407.1954223	0.95820832	0.047076872	0.092279526
		WINE DATASET	0.528726877	1354.575583	0.530361985	116.7433025	395.5195298
	Bisecting K-means	IRIS PLANT DATASET	0.340939732	409.940789	1.006055039	0.046774595	0.064729506
		WINE DATASET	0.527999058	1349.550317	0.521589365	117.171315	435.5555169