Assignment – 6

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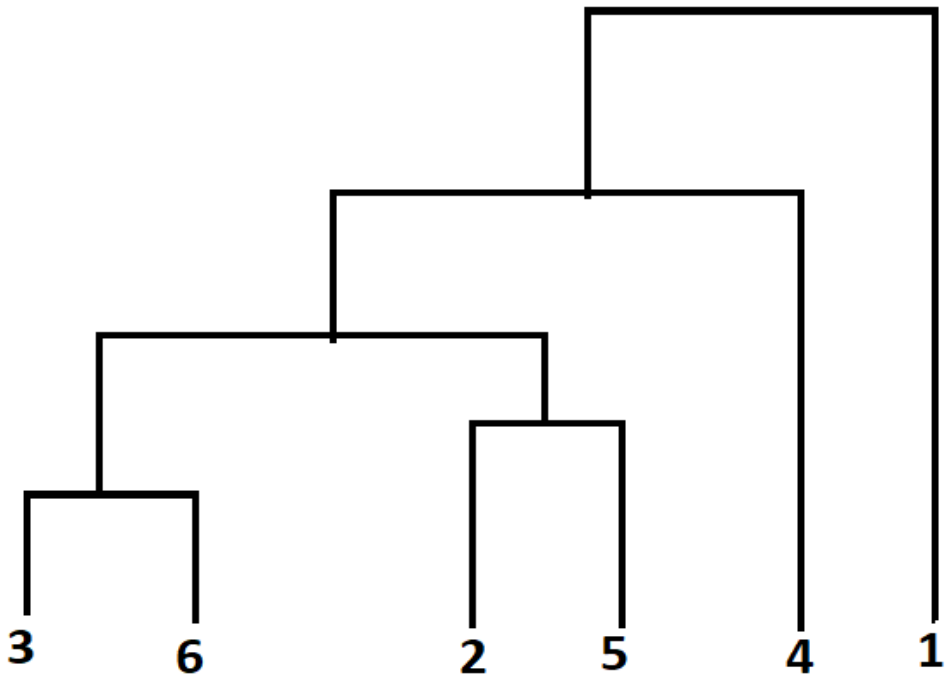
Question - 1:
I have done all the calculations and plotted the dendrograms in question1.docx file. Please refer to the question.docx/question1.pdf file for question 1.

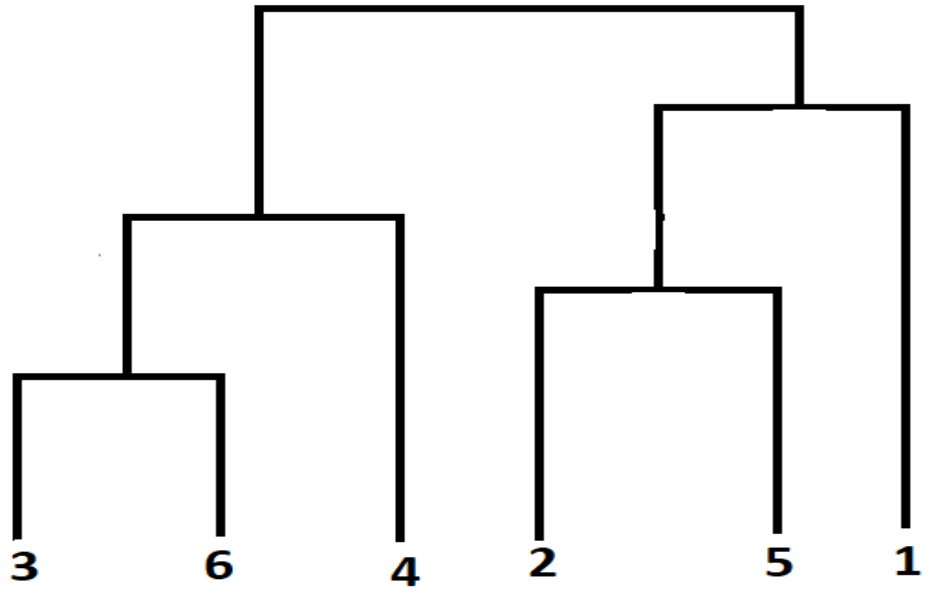
For first question, I have performed all the mathematical calculations for Single link, Complete link and Average link. Plotted the three dendrograms as shown below

For the Single link, I have used minimum distance to do the calculations and plot the dendrogram.
For the Complete link, I have used maximum distance to do the calculations and plot the dendrogram.
For the Average link, I have used average distance to do the calculations and plot the dendrogram.

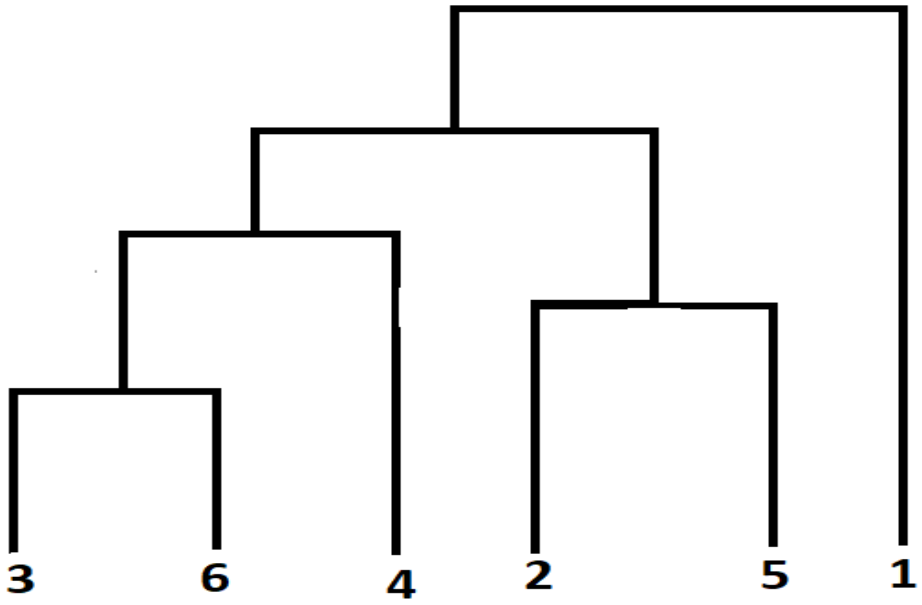
Dendrogram of Single link:



Dendrogram of Complete link:



Dendrogram of Average link:



Question - 2:

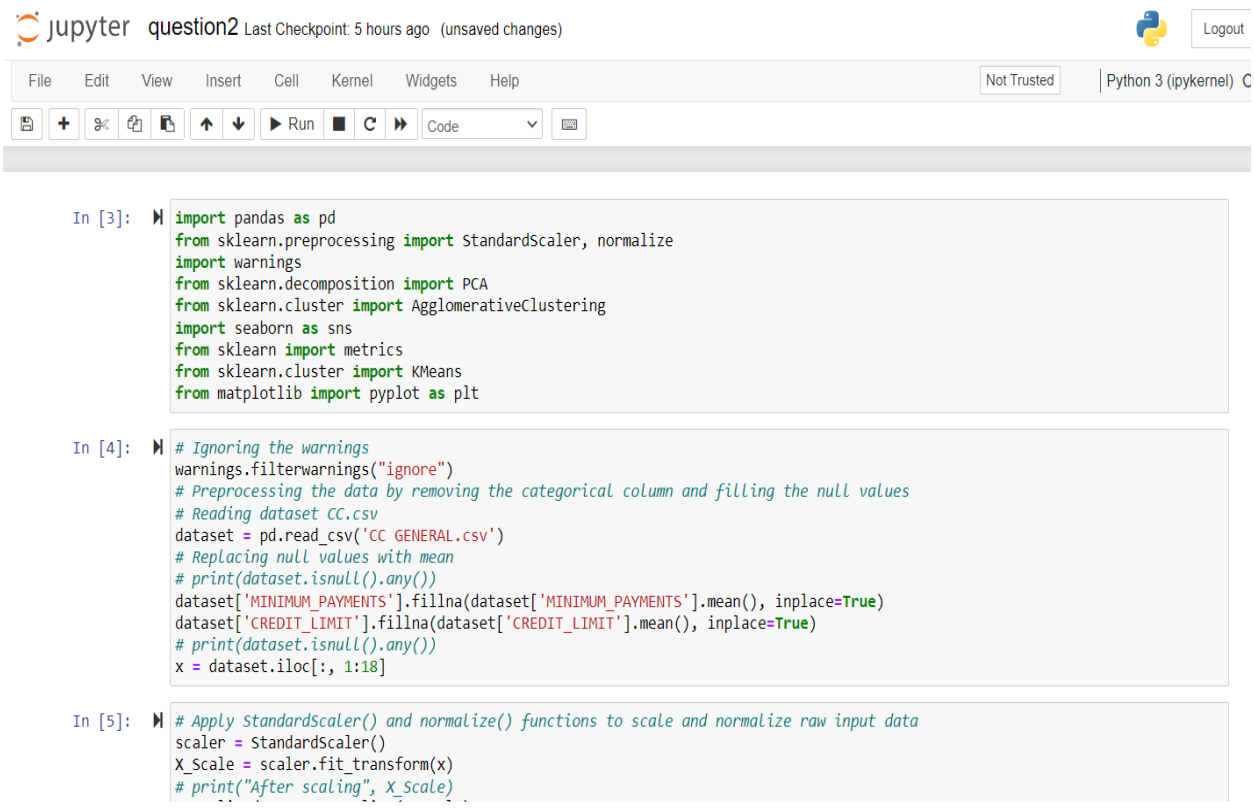
In the second question as explained in the class the followed the below steps

- Imported all the required modules and suppressed warnings.
- Read the data file and replaced the null values with mean and categorized the columns.
- Later applied StandardScaler and normalize functions.
- Used PCA(n=2) to reduce the input dimensions to two features.
- Applied Agglomerative Clustering with k=2,3,4 and 5 on finaldf.
- Visualized the results of Agglomerative clustering for k=2,3,4 and 5 using scatter plot.
- Evaluated the Silhouette scores for each k value i.e., 2,3,4 and 5 and plotted the results in a bar graph with respective to number of clusters.

I have commented the tasks as a heading in the code which are given to the question 2.

Adding the screenshots of the code and output for Question - 2

Code & Output:



Imported all the required modules and suppressed warnings, read the data file CC GENERAL.csv and replaced the null values with mean and categorized the columns in the above screenshot.

In [5]:
```
# Apply StandardScaler() and normalize() functions to scale and normalize raw input data
scaler = StandardScaler()
X_Scale = scaler.fit_transform(x)
# print("After scaling", X_Scale)
normalized_arr = normalize(X_Scale)
# print("After normalizing", normalized_arr)
# Using PCA with K=2
pca = PCA(n_components=2)
principalComponents = pca.fit_transform(x)
principalDf = pd.DataFrame(data=principalComponents, columns=['principal component 1', 'principal component 2'])
finalDf = pd.concat([principalDf, dataset[['TENURE']]], axis=1)
print('CC dataset finalDf after scaling and normalizing:')
print(finalDf)
```

```
CC dataset finalDf after scaling and normalizing:
      principal component 1  principal component 2  TENURE
0              -4326.383956             921.566884      12
1               4118.916676           -2432.846347      12
2               1497.907660           -1997.578692      12
3               1394.548556           -1488.743450      12
4              -3743.351874             757.342659      12
...                     ...                    ...     ...
8945           -4208.357938            1122.443274       6
8946           -4123.924001             951.683803       6
8947           -4379.444202             911.504566       6
8948           -4791.117744            1032.540944       6
8949           -3623.702749            1555.134769       6

[8950 rows x 3 columns]
```

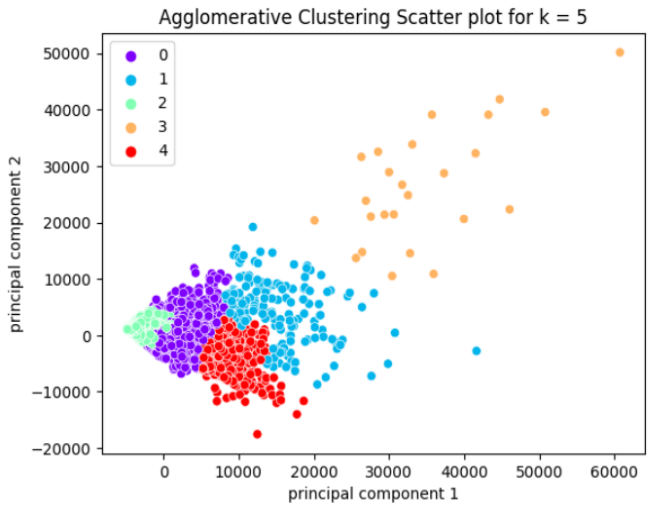
In [6]:
```
# Applying Agglomerative Clustering with k = 2,3,4 and 5 on reduced features and
# visualize result for each k value using scatter plot
# Agglomerative Clustering and scatter plot with k = 5
clustering_model_pca = AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
clustering_model_pca.fit(finalDf)
data_labels_pca = clustering_model_pca.labels_
```

Applied StandardScaler() function and normalize() function to the input data and then applied PCA(components=2) for reducing the data to two features in the above screenshot.

```
[8950 rows x 3 columns]
```

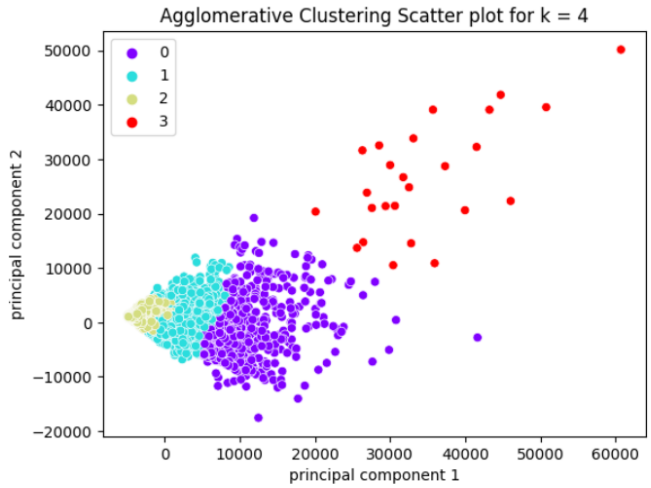
In [6]:
```
# Applying Agglomerative Clustering with k = 2,3,4 and 5 on reduced features and
# visualize result for each k value using scatter plot
# Agglomerative Clustering and scatter plot with k = 5
clustering_model_pca = AgglomerativeClustering(n_clusters=5, affinity='euclidean', linkage='ward')
clustering_model_pca.fit(finalDf)
data_labels_pca = clustering_model_pca.labels_
sns.scatterplot(x=finalDf['principal component 1'],
                y=finalDf['principal component 2'],
                hue=data_labels_pca,
                palette="rainbow").set_title('Agglomerative Clustering Scatter plot for k = 5')
```

Out[6]: Text(0.5, 1.0, 'Agglomerative Clustering Scatter plot for k = 5')



In [7]:
```
# Agglomerative Clustering and scatter plot with k = 4
clustering_model_pca = AgglomerativeClustering(n_clusters=4, affinity='euclidean', linkage='ward')
clustering_model_pca.fit(finalDf)
data_labels_pca = clustering_model_pca.labels_
sns.scatterplot(x=finalDf['principal component 1'],
                y=finalDf['principal component 2'],
                hue=data_labels_pca,
                palette="rainbow").set_title('Agglomerative Clustering Scatter plot for k = 4')
```

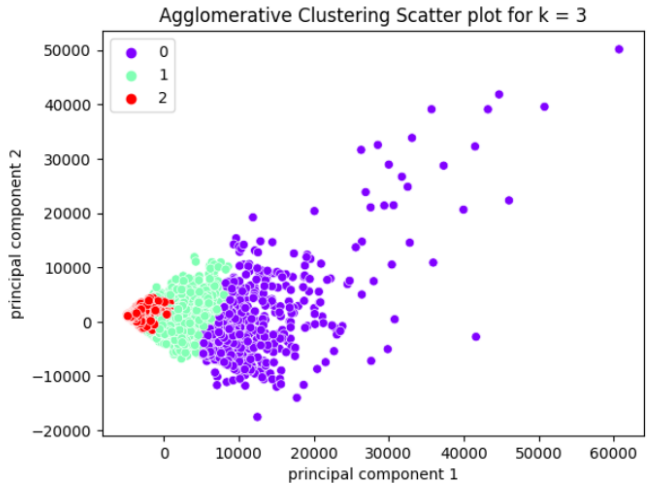
Out[7]: Text(0.5, 1.0, 'Agglomerative Clustering Scatter plot for k = 4')



In [8]:
```
# Agglomerative Clustering and scatter plot with k = 3
clustering_model_pca = AgglomerativeClustering(n_clusters=3, affinity='euclidean', linkage='ward')
```

In [8]:
```
# Agglomerative Clustering and scatter plot with k = 3
clustering_model_pca = AgglomerativeClustering(n_clusters=3, affinity='euclidean', linkage='ward')
clustering_model_pca.fit(finalDf)
data_labels_pca = clustering_model_pca.labels_
sns.scatterplot(x=finalDf['principal component 1'],
                y=finalDf['principal component 2'],
                hue=data_labels_pca,
                palette="rainbow").set_title('Agglomerative Clustering Scatter plot for k = 3')
```

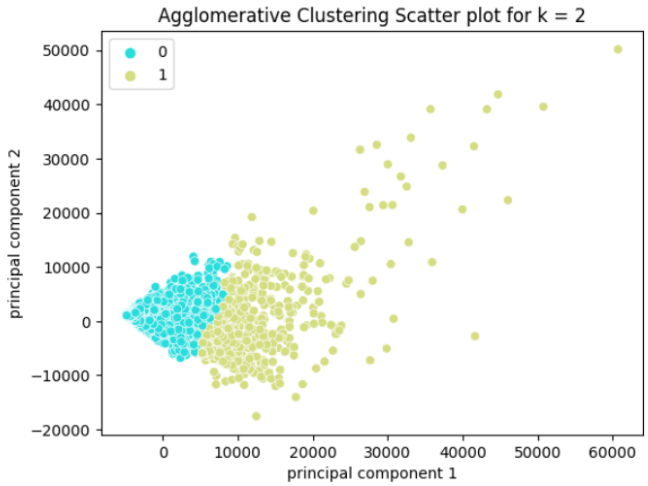
Out[8]: Text(0.5, 1.0, 'Agglomerative Clustering Scatter plot for k = 3')



In [9]:
```
# Agglomerative Clustering and scatter plot with k = 2
clustering_model_pca = AgglomerativeClustering(n_clusters=2, affinity='euclidean', linkage='ward')
```

In [9]:
```
# Agglomerative Clustering and scatter plot with k = 2
clustering_model_pca = AgglomerativeClustering(n_clusters=2, affinity='euclidean', linkage='ward')
clustering_model_pca.fit(finalDf)
data_labels_pca = clustering_model_pca.labels_
sns.scatterplot(x=finalDf['principal component 1'],
                y=finalDf['principal component 2'],
                hue=data_labels_pca,
                palette="rainbow").set_title('Agglomerative Clustering Scatter plot for k = 2')
```

Out[9]: Text(0.5, 1.0, 'Agglomerative Clustering Scatter plot for k = 2')



In [10]:
```
# Evaluating different variations using Silhouette Scores
# Silhouette Scores for 2 clusters
```

Applied Agglomerative Clustering with k=2,3,4 and 5 on finaldf and visualized the results using scatter plot in the above 4 screenshots.

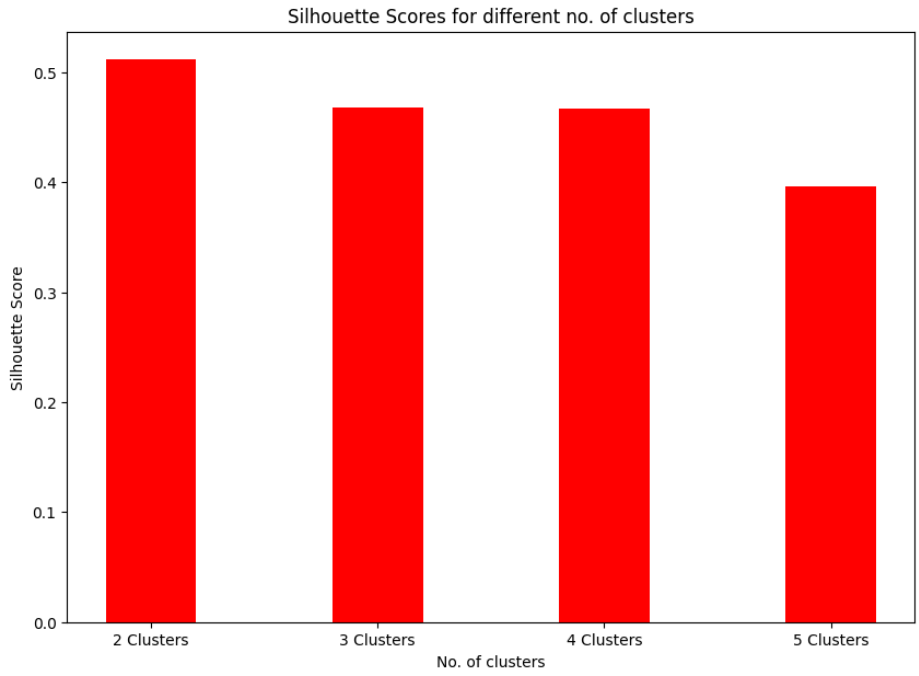
In [10]:
```
# Evaluating different variations using Silhouette Scores
# Silhouette Scores for 2 clusters
nclusters = 2
km = KMeans(n_clusters=nclusters)
km.fit(x)
y_cluster_kmeans = km.predict(x)
score2 = metrics.silhouette_score(x, y_cluster_kmeans)
print('Silhouette Score for 2 clusters =', score2)
# Silhouette Scores for 3 clusters
nclusters = 3
km = KMeans(n_clusters=nclusters)
km.fit(x)
y_cluster_kmeans = km.predict(x)
score3 = metrics.silhouette_score(x, y_cluster_kmeans)
print('Silhouette Score for 3 clusters =', score3)
# Silhouette Scores for 4 clusters
nclusters = 4
km = KMeans(n_clusters=nclusters)
km.fit(x)
y_cluster_kmeans = km.predict(x)
score4 = metrics.silhouette_score(x, y_cluster_kmeans)
print('Silhouette Score for 4 clusters =', score4)
# Silhouette Scores for 5 clusters
nclusters = 5
km = KMeans(n_clusters=nclusters)
km.fit(x)
y_cluster_kmeans = km.predict(x)
score5 = metrics.silhouette_score(x, y_cluster_kmeans)
print('Silhouette Score for 5 clusters =', score5)
```

```
Silhouette Score for 2 clusters = 0.5114773214237212
Silhouette Score for 3 clusters = 0.4676551448676235
Silhouette Score for 4 clusters = 0.46655763491298624
Silhouette Score for 5 clusters = 0.3964843861130676
```

In [11]:
```
# Visualizing different Silhouette Scores with a bar chart
silhouetteScores = [score2, score3, score4, score5]
clusters = ['2 Clusters', '3 Clusters', '4 Clusters', '5 Clusters']
```

Evaluated the Silhouette Scores for k=2,3,4 and 5 in the above screenshot.

```
In [11]:  # Visualizing different Silhouette Scores with a bar chart
          silhouetteScores = [score2, score3, score4, score5]
          clusters = ['2 Clusters', '3 Clusters', '4 Clusters', '5 Clusters']
          fig = plt.figure(figsize=(10, 7))
          plt.bar(clusters, silhouetteScores, color='red', width=0.4)
          plt.xlabel("No. of clusters")
          plt.ylabel("Silhouette Score")
          plt.title("Silhouette Scores for different no. of clusters")
          plt.show()
```



Visualized the Silhouette Scores for different number of clusters using bar graph in the above screenshot.

In GitHub Machine Learning repository, we have an Assignment 6 folder where I have uploaded document(Word & PDF) for question 1 and I have uploaded the code for question 2 in question2.ipynb file. I have uploaded screenshots of dendrograms of single link, complete link and average link for question1 and screenshots of code and output for question 2 in Screenshots folder. As I have explained the code for question 2, I haven't recorded any video for this assignment. I have uploaded word and pdf document of Assignment-6 in Word & PDF file of Assignment 6. The following is the link of the GitHub repository

GitHub link: https://github.com/ChethanKacham/MachineLearning.git

Video link: Explained the code in the class