Mode=Single

March 29, 2022

```
[1]: # MLFA Assignment 6
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# 20CS30066
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```
import scipy.spatial
from scipy import stats
import numpy as np
import statistics
import pandas as pd
from tqdm import tqdm
import matplotlib.pyplot as plt
from collections import Iterable
from scipy.cluster.hierarchy import dendrogram
from sklearn.cluster import AgglomerativeClustering
```

/home/tfjuror/anaconda3/lib/python3.7/site-packages/ipykernel_launcher.py:10: DeprecationWarning: Using or importing the ABCs from 'collections' instead of from 'collections.abc' is deprecated since Python 3.3, and in 3.9 it will stop working

Remove the CWD from sys.path while we load stuff.

```
[3]: # loading csv
df = pd.read_csv('Twitter_data.csv')

# dropping rows with NaN
df = df.dropna(axis=0)

# shuffling dataframe
df = df.sample(frac=1)
```

/home/tfjuror/anaconda3/lib/python3.7/site-packages/IPython/core/interactiveshell.py:3063: DtypeWarning: Columns (0,1,3,4,9,11) have mixed types.Specify dtype option on import or set low memory=False.

interactivity=interactivity, compiler=compiler, result=result)

```
[4]: # converting categorical to numerical
     col_4 = {False: 0, True: 1}
     col_3 = {'en': 0, '_u': 1}
     days ={'Friday':1, 'Saturday':2, 'Sunday':3, 'Thursday':0}
     df.iloc[:,4] = df.iloc[:,4].map(col_4)
     df.iloc[:,3] = df.iloc[:,3].map(col_3)
     df.iloc[:,1] = df.iloc[:,1].map(days)
[5]: # train test split 90%-10%
     df test = df[-int(0.1*len(df)):]
     df = df[:-int(0.1*len(df))]
     print("Test set :",len(df))
     print("Train set :",len(df_test))
    Test set: 898
    Train set: 99
[6]: # droppping non-required columns
     X_columns = [ ' Hour', ' Lang', ' IsReshare', ' Reach',
            ' RetweetCount', ' Likes', ' Sentiment', ' LocationID']
     Y_columns = ['Day']
     X = df[X_columns]
     X.columns = X columns
     Y = df[Y_columns]
     X_test = df_test[X_columns]
     X_test.columns = X_columns
     Y_test = df_test[Y_columns]
[7]: print(X.shape)
     print(Y.shape)
    (898, 8)
    (898, 1)
[8]: # utility function to flatten nested lists
     def flatten(lis):
          for item in lis:
              if isinstance(item, Iterable) and not isinstance(item, str):
                  for x in flatten(item):
                      vield x
              else:
                  yield item
```

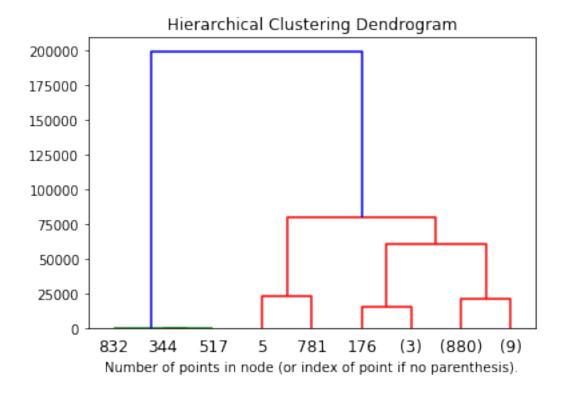
```
[9]: # utility function to find euclidean distance
      def euclidean_dist(x,y):
          return np.sqrt(np.sum(np.square(x-y)))
[10]: # utility function for train accuracy
      def train_accuracy(X,Y,clusters,labels_list):
          for k,cluster in enumerate(clusters):
              for idx in cluster:
                  if(Y.iloc[idx,0] == labels_list[k]):
                      correct+=1
          return correct/len(X)
[11]: # utility function for test accuracy
      def test_accuracy(X_test,Y_test,X,Y,clusters,labels_list):
          correct=0
          cluster_centres=[]
          label_map = np.zeros((len(X),))
          pred_labels=[]
          for k,cluster in enumerate(clusters):
              arr = np.array([X.iloc[idx,:] for idx in cluster])
              mean = np.mean(arr,axis=0)
              cluster_centres.append(mean)
              for idx in cluster:
                  label_map[idx]=labels_list[k]
          test_dist = scipy.spatial.distance.cdist(X_test,X)
          for i in range(len(X_test)):
              closest = np.argmin(np.array(test dist[i]))
              pred_labels.append(label_map[closest])
              if(label_map[closest] == Y_test.iloc[i,0]):
                  correct+=1
          return correct/len(X_test),pred_labels,label_map,cluster_centres
[12]: # mutual distances array calculated only once very efficiently
      dist_arr = scipy.spatial.distance.cdist(X,X)
      for i in range(len(X)):
          dist_arr[i,i]=1e9
[13]: # dictionary to store our dendogram
      levels = {}
[14]: # initialising all data points as separate clusters
      clusters =∏
      for i in range(len(X)):
          clusters.append([i])
      levels[str(len(clusters))] = clusters.copy()
```

```
[15]: # calculating the complete tree with single linkage
      def generator():
          while len(clusters)>1:
              yield
      for _ in tqdm(generator()):
          curr_arr = np.zeros((len(clusters),len(clusters)))
          for i in range(len(clusters)):
              for j in range(len(clusters)):
                  if(i==j):
                      curr_arr[i,j]=1e9
                      continue
                  min=1e9
                  for idx_i in clusters[i]:
                      for idx_j in clusters[j]:
                          if(min>dist_arr[idx_i,idx_j]):
                              min=dist_arr[idx_i,idx_j]
                  curr_arr[i,j]=min
          prev_clusters = clusters.copy()
          i, j = np.argwhere(curr_arr == np.min(curr_arr))[0]
          clusters.pop(i)
          if(i!=j):
              if(i>j):
                  clusters.pop(j)
              else:
                  clusters.pop(j-1)
          clusters.append(list(flatten([prev_clusters[i],prev_clusters[j]])))
          levels[str(len(clusters))] = clusters.copy()
     897it [07:49, 1.91it/s]
[16]: # choose a level to cut
      level_cut = 100
[17]: clusters = levels[str(level_cut)]
      labels list=[]
      for cluster in clusters:
          labels = np.array([Y.iloc[i,0] for i in cluster])
          label = stats.mode(labels)[0]
          labels list.append(label)
[18]: print('Train Accuracy: ',train_accuracy(X,Y,clusters,labels_list))
```

Train Accuracy: 0.48552338530066813

```
[19]: test_acc,pred_labels,label_map,cluster_centres =
       →test_accuracy(X_test,Y_test,X,Y,clusters,labels_list)
      print('Test Accuracy: ',test_acc)
     Test Accuracy: 0.36363636363636365
[20]: def plot_dendrogram(model, **kwargs):
          # Create linkage matrix and then plot the dendrogram
          # create the counts of samples under each node
          counts = np.zeros(model.children_.shape[0])
          n_samples = len(model.labels_)
          for i, merge in enumerate(model.children_):
              current_count = 0
              for child_idx in merge:
                  if child_idx < n_samples:</pre>
                      current_count += 1 # leaf node
                      current_count += counts[child_idx - n_samples]
              counts[i] = current_count
          linkage_matrix = np.column_stack(
              [model.children_, model.distances_, counts]
          ).astype(float)
          # Plot the corresponding dendrogram
          dendrogram(linkage_matrix, **kwargs)
      model = AgglomerativeClustering(distance_threshold=0, n_clusters=None,_
       →linkage='single')
      model = model.fit(X,Y)
[21]: # Visualising the Dendogram
      plt.title("Hierarchical Clustering Dendrogram")
```

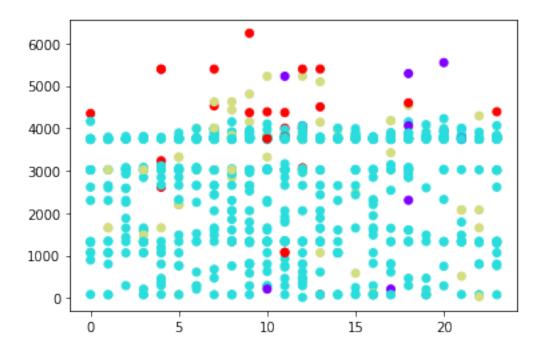
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[21]: # Visualising the Dendogram
plt.title("Hierarchical Clustering Dendrogram")
# plot the top three levels of the dendrogram
plot_dendrogram(model, truncate_mode="level", p=3)
plt.xlabel("Number of points in node (or index of point if no parenthesis).")
plt.show()
```



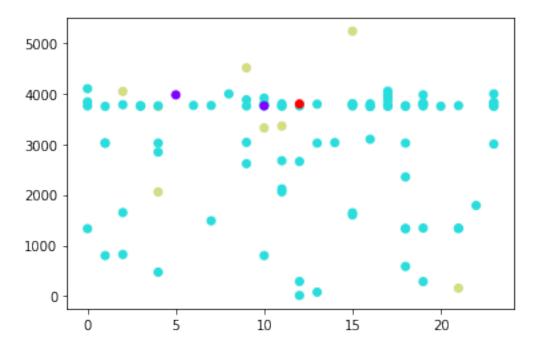
[22]: # We can see that our clusters are the exact same as the scipy package print(levels['3'])

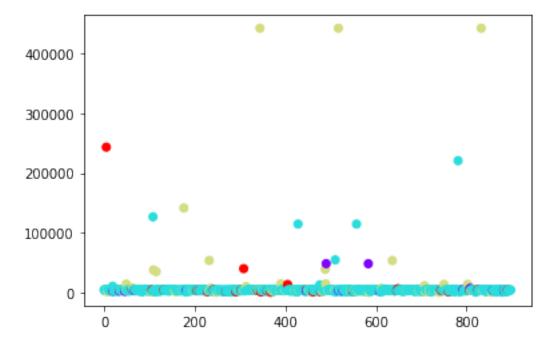
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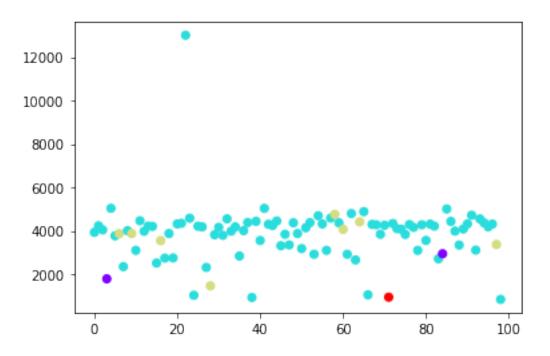
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536, 115, 308, 109, 488, 490, 583, 510, 232, 636]]
```



[24]: # Scatter plot of Oth and 7th attribute along with predicted labels for test set plt.scatter(X_test.iloc[:,0],X_test.iloc[:,7],c=pred_labels,cmap='rainbow') plt.show()







[]: