# **ASSIGNMENT - 3**

# **Preparing the Data:**

- Importing the required libraries such as numpy, pandas, sklearn etc..
- Loading the dataset into a dataframe.

	Elevation	Aspect	Slope	Hillshade_9am	Hillshade_Noon	Horizontal_Distance_To_Hydrology	Vertical_Distance_To_Hydrology
0	elevation_medium	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	0	1
1	elevation_high	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	1	1
2	elevation_medium	aspect_low	slope_low	hillshade_9am_max	hillnoon_max	1	1
3	elevation_high	aspect_ultra	slope_medium	hillshade_9am_max	hillnoon_max	2	1
4	elevation_high	aspect_high	slope_low	hillshade_9am_max	hillnoon_max	2	1
406703	elevation_ultra	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	1	1
406704	elevation_medium	aspect_low	slope_medium	hillshade_9am_max	hillnoon_max	0	1
406705	elevation_medium	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	0	1
406706	elevation_high	aspect_high	slope_low	hillshade_9am_max	hillnoon_max	2	2
406707	elevation_medium	aspect_ultra	slope_low	hillshade_9am_max	hillnoon_max	0	1
406708 rows × 11 columns							

• Finding the data type, not null and count of each feature in the dataframe

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 406708 entries, 0 to 406707
Data columns (total 11 columns):

l	Data	columns (total 11 columns):		
	#	Column	Non-Null Count	Dtype
	0	Elevation	406708 non-null	object
	1	Aspect	406708 non-null	object
	2	Slope	406708 non-null	object
	3	Hillshade_9am	406708 non-null	object
	4	Hillshade_Noon	406708 non-null	object
	5	Horizontal_Distance_To_Hydrology	406708 non-null	int64
	6	Vertical_Distance_To_Hydrology	406708 non-null	int64
	7	Horizontal_Distance_To_Fire_Points	406708 non-null	object
	8	Soil_Type	406708 non-null	int64
	9	Wilderness	406708 non-null	int64
	10	target	406708 non-null	int64

dtypes: int64(5), object(6)
memory usage: 34.1+ MB

• Checking the count of each target label in the dataframe

1 148288 2 198310 3 25028 4 1923 5 6645 6 12157 7 14357 dtype: int64	target				
3 25028 4 1923 5 6645 6 12157 7 14357	1	148288			
4 1923 5 6645 6 12157 7 14357	2	198310			
5 6645 6 12157 7 14357	3	25028			
6 12157 7 14357	4	1923			
7 14357	5	6645			
. 1.55.	6	12157			
dtype: int64	7	14357			
	dtype	: int64			

• Checking if there are any null values in the dataframe.

```
Elevation
                                        0
Aspect
                                        0
Slope
                                        0
Hillshade_9am
                                        0
Hillshade Noon
                                        0
Horizontal_Distance_To_Hydrology
                                        0
Vertical_Distance_To_Hydrology
                                        0
Horizontal_Distance_To_Fire_Points
                                        0
Soil_Type
                                        0
Wilderness
                                        0
target
                                        0
dtype: int64
```

• Dropping the duplicate entries in the dataframe significantly reduced size of the dataframe from 406708 to 12495 rows.

	Elevation	Aspect	Slope	Hillshade_9am	Hillshade_Noon	Horizontal_Distance_To_Hydrology	Vertical_Distance_To_Hydrology
0	elevation_medium	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	0	1
1	elevation_high	aspect_medium	slope_low	hillshade_9am_max	hillnoon_max	1	1
2	elevation_medium	aspect_low	slope_low	hillshade_9am_max	hillnoon_max	1	1
3	elevation_high	aspect_ultra	slope_medium	hillshade_9am_max	hillnoon_max	2	1
4	elevation_high	aspect_high	slope_low	hillshade_9am_max	hillnoon_max	2	1
405630	elevation_high	aspect_high	slope_low	hillshade_9am_max	hillnoon_max	2	2
405688	elevation_medium	aspect_low	slope_medium	hillshade_9am_max	hillnoon_max	1	1
405828	elevation_high	aspect_ultra	slope_low	hillshade_9am_max	hillnoon_max	3	0
405883	elevation_low	aspect_ultra	slope_low	hillshade_9am_max	hillnoon_max	0	1
406025	elevation_high	aspect_low	slope_medium	hillshade_9am_max	hillnoon_max	0	1

12495 rows × 11 columns

separating target column to use clustering algorithms.

```
xx = df.drop(['target'], axis=1)
y = df['target']
```

- Encoding the categorical values into numeric values using one hot encoder.
- One hot encoder will encode categorical features to one-hot numeric array.
- It creates binary column for each category and will return a sparse matrix.

```
data = pd.get_dummies(xx)
data.head()
                                  Horizontal_Distance_To_Hydrology Vertical_Distance_To_Hydrology Soil_Type Wilderness Elevation_elevation_high Elevation_elevation_low Elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_elevation_el
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5 rows × 23 columns
```

- Dimensionality reduction using tsne into two components tsne1 and tsne2.
- TSNE stands for T-distributed Stochastic Neighbor Embedding. It is a tool to visualize high dimensional data.
- We used tsne over pca because it preserves the local stricture or cluster of the data and also it can handle outliers (pca gets highly affected by outliers).

```
tsne = TSNE(n_components=2)
x_emb = tsne.fit_transform(data)
```

#### Q1) (1) Representative object of each cluster

#### K-means Clustering:

```
km = KMeans(n clusters=7, random state=0)
km.fit_predict(data)
y_km = km.labels_
#attribute1: coordinate of cluster centres
print(km.cluster centers )
[[ 1.84917770e+00 1.63056558e+00 3.14384276e+01 1.80144404e+00
  1.66533454e-16 1.54432411e-01 1.31167268e-01 2.71159246e-01
  2.39069394e-01 2.76774970e-01 5.18652226e-01 4.02326514e-01
  3.61010830e-03 5.13437625e-02 9.62695548e-03 6.85118331e-01
  3.14480546e-011
 [ 1.25775194e+00 1.47093023e+00 1.03861434e+01 1.81637597e+00
  1.26937984e-01 5.24709302e-01 -1.80411242e-16 2.88759690e-01
  2.24321705e-01 2.91182171e-01 4.89341085e-01 3.90503876e-01
  3.87596899e-03 8.13953488e-02 1.55038760e-02 7.83914729e-01
  1.47286822e-01]
 [ 1.41123370e+00 1.26328987e+00 2.21850552e+01 1.09879639e+00
  1.52655666e-16 1.72016048e-01 1.70511535e-02 3.09929789e-01
  2.33701103e-01 2.77331996e-01 5.69709127e-01 3.87662989e-01
  4.01203611e-03 3.20962889e-02 3.51053159e-03 6.08324975e-01
  3.40020060e-01]
 [ 1.74327752e+00 1.52663623e+00 3.79766616e+01 9.75646880e-01
  1.24900090e-16 -4.99600361e-16 5.18011162e-01 3.06443430e-01
  2.63318113e-01 2.68899036e-01 5.35768645e-01 3.89649924e-01
  2.02942669e-03 3.09487570e-02 8.11770675e-03 6.31659056e-01
  3.51090817e-01]
 1.11022302e-16 3.02366864e-01 1.53846154e-02 2.89349112e-01
  2.62721893e-01 2.55621302e-01 4.19526627e-01 4.31952663e-01
  4.73372781e-03 6.50887574e-02 1.59763314e-02 5.29585799e-01
  3.15976331e-01]
 9.23988842e-01 1.46443515e+00 2.55299861e+00 2.60111576e+00
  3.89818689e-01 5.45327755e-01 1.24900090e-16 2.55927476e-01
  3.04044630e-01 1.91073919e-01 4.40027894e-01 4.30962343e-01
  6.97350070e-04 4.18410042e-02 3.48675035e-03 9.53974895e-01
  4.18410042e-021
 [ 8.22143698e-01 1.03062426e+00 1.73345112e+01 9.41107185e-01
  6.36042403e-02 3.75736160e-01 8.24499411e-03 2.93286219e-01
  2.54416961e-01 2.48527680e-01 7.53828033e-01 2.29681979e-01
  -4.33680869e-19 9.42285041e-03 -3.46944695e-18 5.34746761e-01
  2.76796231e-01]]
```

```
#attribute2: cluster label of each point
print(km.labels_)
[2 0 1 ... 4 5 3]
```

- Representative object: cluster centers represent coordinates of cluster centers.
- Labels represent the labels of each point.

#### **Hierarchical Clustering: Agglomerative Clustering:**

```
#compute agglomerative clustering
 am = AgglomerativeClustering(n clusters=7,compute distances=True)
 am.fit predict(data)
y am = am.labels
 #attribute1: cluster label of each point
print(am.labels_)
 [6 2 0 ... 5 1 3]
 #attribute2: distance between nodes in corresponding place in children(of non-leaf node)
print(am.distances )
 0.
                                          ... 416.02339628 736.13165651
 1543.22871934]
#attribute3: number of leaves in hierarchical tree
print(am.n leaves )
#children of each non-leaf node
am.children
12495
array([[
            0, 2643],
            1, 1045],
            2, 1381],
       [24977, 24984],
       [24983, 24986],
       [24985, 24987]], dtype=int64)
```

- Representative object: Labels represent the labels of each point.
- N\_leaves represents the numbr of leaves in hierarchical tree.
- Distances shows distance between nodes in corresponding place in children(of non-leaf node)

#### **BIRCH Clustering:**

```
#compute BIRCH clustering
br = Birch(n clusters=7)
br.fit_predict(data)
y_br = br.labels_
##attribute1: cluster label of each point
print(br.labels_)
[6 0 1 ... 2 4 3]
#attribute2: centroid of all subclusters
print(br.subcluster_centers_)
                              0.]
[[ 1.6 2.
           26.
                     0.
                         1.
           25. ... 0.
25. ... 0.
[ 3.
                              0.]
       1.
                         1.
 [ 2.
                              0.]
       2.
                         1.
 . . .
                              0.]
 [ 1.
          3. ... 0.
                         1.
      1.
                              0. ]
 [ 2. 0. 2. ... 0.
                         1.
 [ 2. 0. 2. ... 0. 1.
                              0.]]
```

- Representative object: Labels represent the labels of each point.
- Subcluster centers shows centroid of all subclusters

#### **Gaussian Mixture Clustering:**

```
gmm = GaussianMixture(n components = 7, random state=0)
y_gmm = gmm.fit_predict(data)
#attribute1: mean of each mixture component
print(gmm.means )
[[1.84182652e+00 1.49928428e+00 2.80377904e+01 1.18880615e+00
  0.00000000e+00 0.00000000e+00 1.88233618e-01 2.80131694e-01
  2.62381908e-01 2.51503007e-01 5.87317504e-01 4.12682496e-01
  0.00000000e+00 0.00000000e+00 0.00000000e+00 6.39278543e-01
  3.60721457e-01]
 [3.88276291e-01 1.26712462e+00 1.50317747e+01 1.47390958e+00
  2.80019109e-01 2.94761598e-01 0.00000000e+00 4.41915254e-01
  4.42242598e-02 4.65987177e-01 0.00000000e+00 1.32432888e-01
  0.00000000e+00 5.44185974e-01 5.21238257e-01 5.21238257e-01
  2.58060400e-011
 [1.00720739e+00 1.30853136e+00 1.60282591e+01 1.41233770e+00
  0.0000000e+00 8.65051057e-01 0.0000000e+00 3.15744887e-01
  2.67301944e-01 2.32121752e-01 5.91121534e-01 4.08878466e-01
  0.00000000e+00 0.00000000e+00 0.00000000e+00 6.49940290e-01
  1.58017093e-01]
 [8.19414908e-01 1.51769984e+00 1.12632025e+01 2.53026812e+00
  5.84171049e-01 2.14311883e-01 0.00000000e+00 2.38789079e-01
  1.75868663e-01 3.53880904e-01 2.15653893e-01 3.48566194e-01
  0.0000000e+00 2.67246869e-01 0.00000000e+00 1.00000000e+00
  0.00000000e+001
 [9.72130742e-01 1.46933759e+00 3.56607355e+01 7.80095752e-01
  0.00000000e+00 0.00000000e+00 4.47529731e-01 2.68014032e-01
  1.98191928e-01 4.01809108e-01 2.95074622e-02 2.16952715e-01
  0.00000000e+00 4.20276782e-01 0.00000000e+00 4.02133568e-01
  5.48687328e-011
 [8.61612792e-01 1.86371274e+00 2.88523165e+01 1.72926548e+00
  0.00000000e+00 2.84269052e-01 1.21829486e-01 4.55479717e-01
  1.89502610e-01 2.88789285e-01 0.00000000e+00 3.01706651e-02
  4.87316681e-01 4.09164003e-01 7.39756341e-01 8.07439775e-01
  1.92560225e-01]
 [9.39374722e-01 1.36879716e+00 2.84458268e+01 0.00000000e+00
  0.00000000e+00 2.94773058e-01 0.00000000e+00 3.42403729e-01
  2.81824780e-01 3.28665464e-01 0.00000000e+00 1.12413416e-01
  8.67182498e-03 2.37553106e-01 9.09846326e-02 6.02307397e-01
  3.28209402e-01]]
```

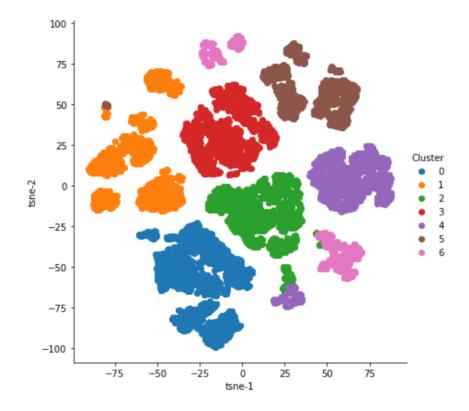
# #attribute2: weights of each mixture component print(gmm.weights\_)

[0.55910362 0.00543037 0.27754936 0.11727285 0.01627358 0.00591228 0.01845794]

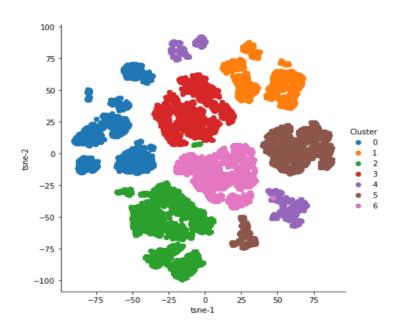
- Representative object: weights shows the weight of each mixture component
- Means shows the mean of each mixture component.

# Q1 (2) Visualization of clusters

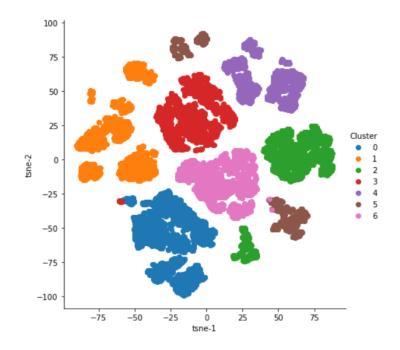
#### K-means Clustering:



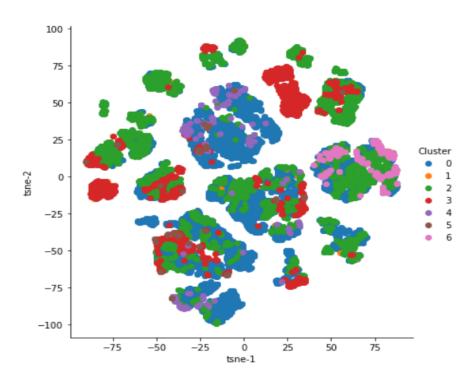
# Hierarchical Clustering: Agglomerative Clustering:



### **BIRCH Clustering:**



#### **Gaussian Mixture Clustering:**



# Q1 (3) Cluster distribution with true label count

• Finding the true label count of target column

```
#value count of each label present in target column
y.value_counts()
2     4711
```

Name: target, dtype: int64

• True label counts for K-means clustering

```
#K-means clustering distribution with true label count tl_count(y_km,y)
```

```
Cluster:0
2 1125
    1018
1
7
     180
5
    113
6
     43
3
     14
Name: target, dtype: int64
Cluster:1
    973
3
    329
1
    318
6
    250
5
    156
   35
4
     3
Name: target, dtype: int64
Cluster:2
2
    372
1
    304
5
    64
3
    48
6
    46
4
    13
7
     2
Name: target, dtype: int64
Cluster:3
3
    665
6
    327
2
    237
    127
5
    62
1
    11
7
     5
Name: target, dtype: int64
Cluster:4
1
    1002
7
     814
2
     155
Name: target, dtype: int64
Cluster:5
2
   959
1
    881
7
    81
5
     53
     20
Name: target, dtype: int64
Cluster:6
1
    578
5
    149
7
    73
Name: target, dtype: int64
```

Datapoints present in each cluster:

```
#check the count of datapoints present in each cluster: K-means
dataframe1 = pd.DataFrame()
dataframe1['kmean'] = km.labels_
dataframe1['kmean'].value_counts()
0
     2493
1
    2064
5
    1994
4
    1971
6
    1690
3
    1434
2
     849
Name: kmean, dtype: int64
```

• True label counts for Hierarchical clustering (agglomerative clustering)

```
#Agglomerative clustering distribution with true label count tl_count(y_am,y)
```

```
Cluster:0
   984
3
   329
1
   318
   251
6
5
  156
    35
7
     3
Name: target, dtype: int64
Cluster:1
3 665
6 327
2
  226
   127
4
5
   62
1
    11
     - 5
Name: target, dtype: int64
Cluster:2
2 1133
1
  1023
7
    180
5
    113
6
     43
3
     14
Name: target, dtype: int64
Cluster:3
1 997
    814
2
    147
Name: target, dtype: int64
Cluster:4
2 371
1
   304
5
   64
3
    48
    45
6
4
    13
7
     2
Name: target, dtype: int64
Cluster:5
2 973
1
   592
5
  161
    73
Name: target, dtype: int64
Cluster:6
   877
1
    867
7
    81
5
    41
     20
Name: target, dtype: int64
```

#### Datapoints in each clusters:

• True label counts for BIRCH clustering

# #BIRCH clustering distribution with true label count tl\_count(y\_br,y)

```
Cluster:0
   984
3
   329
1
    318
    251
6
5
   156
   35
7
    3
Name: target, dtype: int64
Cluster:1
1 1006
  820
2 147
Name: target, dtype: int64
Cluster:2
   1133
2
1
   1014
7
    174
    113
5
6
     43
    14
3
Name: target, dtype: int64
Cluster:3
   376
2
   316
1
    64
5
    48
3
6
   45
4
   13
Name: target, dtype: int64
Cluster:4
3 665
   327
6
2
   226
   127
4
5
    62
   11
1
7
    5
Name: target, dtype: int64
Cluster:5
2 979
1
   593
5 161
   73
Name: target, dtype: int64
Cluster:6
2 866
   854
1
7
    79
5
     41
6
     20
Name: target, dtype: int64
```

#### Datapoints in each cluster:

```
#check the count of datapoints present in each cluster: Birch
dataframe3 = pd.DataFrame()
dataframe3['BIRCH'] = br.labels_
dataframe3['BIRCH'].value_counts()
     2491
0
    2076
1
    1973
    1860
6
5
     1806
    1423
4
3
     866
Name: BIRCH, dtype: int64
```

#### True label counts for Gaussian mixture model clustering

```
#Gaussian mixture clustering distribution with true label count
tl_count(y_gmm,y)
Cluster:0
2
   27
3
    18
1
    12
Name: target, dtype: int64
Cluster:1
   54
1
     48
    20
     3
Name: target, dtype: int64
Cluster:2
    24
1
    21
Name: target, dtype: int64
Cluster:3
3
     373
    276
6
4
    143
1
    118
     17
5
      8
Name: target, dtype: int64
Cluster:4
2
    4042
    3077
1
5
     560
3
     531
7
     492
     405
6
      32
Name: target, dtype: int64
Cluster:5
   750
1
    623
2 101
Name: target, dtype: int64
Cluster:6
    77
1
5
    26
     6
Name: target, dtype: int64
```

#### Datapoints in each cluster:

```
#check the count of datapoints present in each cluster: Gaussian Mixture
dataframe4 = pd.DataFrame()
dataframe4['GMM'] = y_gmm
dataframe4['GMM'].value_counts()

4 9139
5 1474
3 1442
6 208
1 125
0 62
2 45
Name: GMM, dtype: int64
```

# Q1 (4) Comparison of cluster formation of gaussian based model

- We can see in the clusters formed above that the Gaussian model is considering a
  certain number of distributions as clusters. So, with the plot, we can see that the
  grouping is done for data points based on the distribution of clusters. Visualizing the
  Gaussian mixture model in 3D will give us show us better visualization of distributions.
- Gaussian Mixture Models are probabilistic models and use the soft clustering approach
  for distributing the points in different clusters. Uses mean as well as the variance of the
  data to update the centroid.
- With K-means clustering: In the above-visualized plots, we can see that the clusters formed are almost circular in shape. This is because the centroids of the clusters are updated iteratively using the mean value. But as the distribution of points is not circular so it fails to identify the right clusters. Hence, instead of distance-based, a distribution-based model like Gaussian gives better clusters.
- With agglomerative clustering: We can see that agglomerative clustering doesn't
  handle the outliers well. Also, as it is creating equal size clusters so it didn't predict the
  data points in the right clusters as well.
- With BIRCH (Balanced Iterative Reducing and Clustering using Hierarchies): Birch performs better than the above two clustering algorithms in terms of scaling the data but the Gaussian mixture model performs better in terms of distribution.
- The Rand index computes the similarity measure between two clustering by considering all pairs of samples and counting pairs assigned in the same or different clusters in true and predicted clustering.
- The raw RI score is:

RI = (number of agreeing pairs) / (number of pairs)

#rand score for K-means
rand\_score(y,km.labels\_)

0.6901680089356056

#rand score for agglomerative
rand\_score(y,am.labels\_)

0.6906230652978336

#rand score for BIRCH
rand\_score(y,br.labels\_)

0.6908459045535935

#rand score for gaussian mixture
rand\_score(y,y\_gmm)

0.6016009989717033

# **Q2 Prediction of clusters**

```
#compute Gaussian mixture clustering (Gaussian Model)
gmm = GaussianMixture(n components = 7)
gmm.fit(data)
y gmm = gmm.predict(data)
def map cl(y target,y label):
    cluster map={}
    for i in range(7):
        add=np.where(y_label==i)[0].tolist()
        y=y target.iloc[add]
        check y=check label(y)
        cluster map[i]=check y
    return cluster map
#checking after mapping with GMM
cluster map=map cl(y,y gmm)
cluster map
{0: 1, 1: 7, 2: 6, 3: 2, 4: 7, 5: 5, 6: 3}
#function to assign clusters
def assign(cluster map,y label):
    yp=np.zeros(y label.shape)
    for i in cluster map:
        add=np.where(y label==i)[0].tolist()
        yp[add]=cluster map[i]
    return yp
yp=assign(cluster map,y gmm)
f1 score(y,yp,average='weighted')
0.3466280015913269
gmm.cluster map=cluster map
```

**Conclusion:** So, we are able to get 6 out of 7 clusters mapped and predicting based on the GMM which is giving best score.

#### For inference:

```
def predict(test_set) :
    prediction=[]
    model= pickle.load(open('final_model','rb'))
    testpd=pd.read_csv(test_set)
    testpd=pd.get_dummies(testpd)
    prediction=model.predict(testpd)
    yp=assign(model.cluster_map,np.array(prediction))
    return yp
```

- In the above predict function, we pass the .csv file.
- We do the pre processing steps inside the function itself like encoding.
- Then we load the dumped model into model and do the prediction on it.
- Finally, we return list of output labels.

#### **References:**

- <a href="https://scikit-learn.org/stable/modules/clustering.html">https://scikit-learn.org/stable/modules/clustering.html</a>
- <a href="https://www.analyticsvidhya.com/blog/2016/11/an-introduction-to-clustering-and-different-methods-of-clustering/">https://www.analyticsvidhya.com/blog/2016/11/an-introduction-to-clustering-and-different-methods-of-clustering/</a>
- https://www.geeksforgeeks.org/clustering-in-machine-learning/
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