# **Data Reading and Cleaning**

### importing data

```
import matplotlib.pyplot as plt
In [11]:
         import numpy as np
         import pandas as pd
         import seaborn as sns
         import cv2 as cv
         import glob
         from sklearn import metrics
         from sklearn.preprocessing import LabelEncoder,MinMaxScaler,RobustScaler
         from sklearn.metrics.pairwise import rbf kernel
         from sklearn.preprocessing import normalize
         from sklearn.model selection import train test split
         from sklearn.metrics.cluster import contingency matrix
         from sklearn.cluster import KMeans
         import math
         from sklearn.metrics.pairwise import euclidean distances
         from scipy.spatial import distance matrix
         from scipy.optimize import linear sum assignment
```

```
In [12]: from google.colab import drive
drive.mount('/content/drive', force_remount=True)
```

Mounted at /content/drive

```
In [13]: | columns=[
             "duration"
             , "protocol_type",
             "service", "flag",
             "src bytes",
             "dst_bytes", "land", "wrong_fragment", "urgent",
             "hot", "num failed logins", "logged in",
             "num compromised", "root shell",
             "su attempted", "num root",
             "num file creations", "num shells",
             "num_access_files", "num_outbound_cmds",
             "is host login", "is guest login", "count",
             "srv count", "serror rate",
             "srv serror rate", "rerror rate",
             "srv_rerror_rate", "same_srv_rate",
             "diff srv rate", "srv diff host rate",
             "dst host count", "dst host srv count",
             "dst host same srv rate", "dst host diff srv rate",
             "dst host same src port rate",
             "dst host srv diff host rate",
             "dst host serror rate",
             "dst host srv serror rate", "dst host rerror rate", "dst host srv rerror rate", "target"]
In [14]: def import data(path, columns, categorical cols):
             data = pd.read csv(path,compression = "gzip", names = columns)
             data = data.drop duplicates(keep = 'first')
             data = data[~data['service'].isin(['red_i', 'urh_i', 'icmp'])]
             dummy data = pd.get dummies(data,columns=categorical cols)
             labels = dummy data['target']
             dummy data.drop('target',axis = 1, inplace=True)
             dummy data.drop('num outbound cmds',axis = 1, inplace=True)
             dummy data.drop('is host login',axis = 1, inplace=True)
             return [dummy data,labels]
```

data cleaning and encoding

```
In [15]: # One-Hot Encoding for categorical data.
         #data, labels = import data("kddcup.data_10_percent.gz", columns, ['service', 'flag', 'protocol_type'])
         path = "/content/drive/MyDrive/pattern/NetworkAnomaly/dataset/kddcup.data 10 percent.gz/kddcup.data 10 percent
         path of whole data = "/content/drive/MyDrive/pattern/NetworkAnomaly/dataset/kddcup.data.gz/kddcup.data.gz"
         path test="/content/drive/MyDrive/pattern/NetworkAnomaly/dataset/corrected.gz"
         # importing training set of kmeans
         data, labels = import data(path, columns, ['service', 'flag', 'protocol type'])
         # importing training set of spectral clustering
         whole data, whole labels = import data(path of whole data, columns, ['service', 'flag', 'protocol type'])
         #importing test set of kmeans
         test set, ground truth=import data(path test, columns, ['service', 'flag', 'protocol type'])
         # removing rows from test set taht have targets not in the training set
         test = pd.concat([test set, ground truth],axis = 1)
         test = test[test['target'].isin(np.unique(labels).tolist())]
         ground truth = test['target']
         test.drop('target', axis = 1, inplace=True)
         # label encoding for labels of training and testing set
         le = LabelEncoder()
         le.fit(labels)
         labels = le.transform(np.array(labels))
         ground truth = le.transform(np.array(ground truth))
```

In [ ]:

# **Evaluation Functions**

**Conditional Entropy** 

```
In [3]: # calculating TP, TN, FP, FN for precison, recall and F1-Score
        def confusion mat(contingency matrix):
            x=contingency matrix.shape[0]
            y=contingency matrix.shape[1]
            TP=0
            for i in range (x):
                 for j in range(y):
                    TP += math.comb(contingency matrix[i][j],2)
            FP=0
            for i in range (x):
                s=np.sum(contingency matrix[i])
                for j in range(y):
                    l=contingency matrix[i][j]*(s-contingency matrix[i][j])
                    FP+=1
            FP=FP//2
            FN=0
            for i in range(x):
                for j in range(y):
                    FN+=contingency matrix[i][j]*np.sum(contingency matrix[i+1:,j])
            total=math.comb(np.sum(contingency matrix),2)
            TN=total-(FN+TP+FP)
            conf=[TP,FP,FN,TN]
            return conf
```

#### Precision

```
In [4]: def precision(contingency_matrix):
    precision=[]
    for i in range(contingency_matrix.shape[0]):
        class_max=np.max(contingency_matrix[i])
        precision.append(class_max/np.sum(contingency_matrix[i]))
    return precision
```

### Recall

```
In [5]: def recall(contingency_matrix):
    recall=[]
    for i in range(contingency_matrix.shape[0]):
        class_max=np.max(contingency_matrix[i])
        index=np.where(contingency_matrix==class_max)[1]
        recall.append(class_max/np.sum(contingency_matrix[:,index]))
    return recall
```

# F1-Score

```
In [6]: def f1_score(precision,recall,clusters):
    f1=0
    for i in range(clusters):
        f1+=(2*precision[i]*recall[i])/(precision[i]+recall[i])
    return (1/clusters)*(f1)
```

# K-means

```
In []: k_list = [7, 15, 23, 31, 45]
In [7]: def euclidean(point, data):
    return np.sqrt(np.sum(np.subtract(point,data)**2, axis=1))
```

# Implementation

```
In [18]: def k means(data, k):
             new = data.sample(n=k).to numpy()
             epoch = 0
             while True:
                 points = [[] for i in range(k)]
                 previous=[]
                 cluster=[]
                 epoch += 1
                 #print(epoch)
                 for point in data.values.tolist():
                     distances=np.array(euclidean(np.array([np.array(point)]*k),np.array(new)))
                     #distances = np.array([np.linalq.norm(point-centroid) for centroid in new])
                     centroid idx = np.argmin(distances)
                     cluster.append(centroid idx)
                     points[centroid idx].append(point)
                 previous=new
                 new=[np.mean(cluster, axis=0) for cluster in points]
                 if(np.array equal(new, previous)):
                     break
             return cluster,new
```

```
In [9]: def predict(data, centroids):
    cluster=[]
    for point in test.values.tolist():
        distances=np.array(euclidean(np.array(point),centroids))
        centroid_idx = np.argmin(distances)
        cluster.append(centroid_idx)
    return cluster
```

### **Data Normalization**

```
In [10]: #normalizing data to improve k means performance
def standardization_robust(df, scaler):
    arr = np.array(df)
    arr=scaler.transform(arr)
    return arr

def standardization_minmax(arr, col, minmax_scaler):
    df=pd.DataFrame(minmax_scaler.transform(arr),columns=col)
    return df
```

```
In [16]:
    scaler = RobustScaler()
    minmax_scaler= MinMaxScaler()
    arr = np.array(data)
    scaler.fit(arr)
    arr = standardization_robust(data,scaler)
    minmax_scaler.fit(arr)
    data = standardization_minmax(arr, data.columns, minmax_scaler)

    arr_test = np.array(test)
    arr_test = standardization_robust(test,scaler)
    test = standardization_minmax(arr_test, test.columns, minmax_scaler)
```

### **Clustering Using K-Means**

```
In [ ]: cluster 7,centroids 7 = k means(data, 7)
In [ ]: clusters 7=predict(test, centroids 7)
 In [ ]: data
Out[54]: (145571, 114)
In [ ]: \#\#Evaluation \ when \ k = 7
        cont 7 = metrics.cluster.contingency matrix(ground truth,clusters 7)
        e 7=conditional entropy(cont 7.T)
        p 7=precision(cont 7.T)
        r 7=recall(cont 7.T)
        f1 7=f1 score(p 7,r 7,cont 7.shape[1])
        print(f"conditional entropy when k is 7 = \{e \ 7\}")
        print(f"precision when k is 7 = {p 7}")
        print(f"recall when k is 7 = \{r \}")
        print(f"F1 score when k is 7 = \{f1 7\}")
        conditional entropy when k is 7 = 0.35894129224731053
        precision when k is 7 = [0.9744268077601411, 0.9493218378719378, 0.768194611982308, 1.0, 0.7306805960977109,
        0.9490234104446599, 0.6322701688555347]
        926739162196573, 0.6739130434782609, 0.3918604651162791]
        F1 score when k is 7 = 0.4612489954277501
In [ ]: cluster 15,centroids 15 = k means(data, 15)
In [ ]: clusters 15=predict(test, centroids 15)
```

```
In [ ]: \##Evaluation when k = 15
       cont 15 = metrics.cluster.contingency_matrix(ground_truth,clusters_15)
       e 15=conditional entropy(cont 15.T)
       p 15=precision(cont 15.T)
       r 15=recall(cont 15.T)
       f1 15=f1 score(p 15,r 15,cont 15.shape[1])
       print(f"conditional entropy when k is 15= {e 15}")
       print(f"precision when k is 15 = {p 15}")
       print(f"recall when k is 15 = {r 15}")
       print(f"F1 score when k is 15 = \{f1 \ 15\}")
       conditional entropy when k is 15= 0.2853081783108416
       precision when k is 15 = [1.0, 1.0, 0.949210088691796, 0.9995949777237748, 0.9426778687652877, 0.99968622529]
       02416, 0.8085106382978723, 1.0, 0.49001248439450684, 0.6827133479212254, 0.9953488372093023, 0.9231448763250
       883, 0.9293119698397738, 0.7116883116883117, 0.9963518443453587]
       700194109912129, 0.06649829892926468, 0.95, 0.15310840055085578, 0.03276909269270105, 1.0, 0.013399845546951
       639, 0.05139681290576431, 0.02057982509235875, 0.3186046511627907, 0.05130345849596126]
       F1 score when k is 15 = 0.3347449968336772
In [ ]: cluster 23,centroids 23 = k means(data, 23)
```

In [ ]: clusters 23 = predict(test, centroids 23)

```
In [ ]: |cont 23 = metrics.cluster.contingency matrix(ground truth,clusters 23)
        e 23=conditional entropy(cont 23.T)
        p 23=precision(cont 23.T)
        r 23=recall(cont 23.T)
        f1 23=f1 score(p 23,r 23,cont 23.shape[1])
        print(f"conditional entropy when k is 15= {e 23}")
        print(f"precision when k is 15 = {p 23}")
        print(f"recall when k is 15 = \{r \ 23\}")
        print(f"F1 score when k is 15 = \{f1 \ 23\}")
        conditional entropy when k is 15= 0.2535268982825815
        precision when k is 15 = [0.9829706717123936, 0.4521072796934866, 0.49588607594936707, 0.9505833905284832,
        1.0, 1.0, 0.9953488372093023, 1.0, 0.8, 0.8631578947368421, 1.0, 0.9147003745318352, 0.964856455946825, 0.68
        27133479212254, 0.9824035192961408, 0.9676025917926566, 1.0, 0.9828473413379074, 0.9047619047619048, 0.99935
        58776167472, 0.9293119698397738, 0.7116883116883117, 1.0
        recall when k is 15 = [0.05110171158764509, 0.6781609195402298, 0.03270647659201436, 0.028907766483688507,
        0.043497317943687254, 0.26144309240049257, 0.013399845546951639, 0.12064725555774149, 0.0007513932082402788,
        0.09534883720930233, 0.06649829892926468, 0.20389889586942456, 0.6711095809561283, 1.0, 0.10254430089123584,
        0.009350671035879025, 0.15305921699783592, 0.035879025693473315, 0.95, 0.1295318402871992, 0.020579825092358
        75, 0.3186046511627907, 0.04257894846694914]
        F1 score when k is 15 = 0.26036820368872193
In [ ]: | cluster 31,centroids 31 = k means(data, 31)
```

In [ ]: clusters 31 = predict(test, centroids 31)

```
In [ ]: cont_31 = metrics.cluster.contingency_matrix(ground_truth,clusters_31)
    e_31=conditional_entropy(cont_31.T)
    p_31=precision(cont_31.T)
    r_31=recall(cont_31.T)
    f1_31=f1_score(p_31,r_31,cont_31.shape[1])
    print(f"conditional entropy when k is 45= {e_31}")
    print(f"precision when k is 45 = {p_31}")
    print(f"recall when k is 45 = {r_31}")
    print(f"F1 score when k is 45 = {f1_31}")
```

conditional entropy when k is 45 = 0.24736919864636803
precision when k is 45 = [0.7063711911357341, 0.9985974754558204, 1.0, 0.9987129987129987, 0.984591679506933
7, 1.0, 0.9504132231404959, 0.5778894472361809, 0.999618320610687, 0.8500196309383589, 0.9402618657937807,
0.9588268471517203, 1.0, 0.8954545454545455, 0.9182608695652174, 1.0, 1.0, 0.6428571428571429, 0.99019745133
73477, 0.3414154652686763, 1.0, 1.0, 1.0, 1.0, 0.8947368421052632, 0.6815245478036176, 0.9991652754590985,
0.9378125400692396, 0.7415730337078652, 1.0, 0.6827133479212254]
recall when k is 45 = [0.29651162790697677, 0.014860887896307738, 0.047167027346055476, 0.01619669804429045
6, 0.2199980326578792, 0.09285854810151485, 0.03360397403519025, 0.6609195402298851, 0.054663855899480286,
0.10648239228801888, 0.023981966563002232, 0.03548245705579094, 0.01903403501868975, 0.02906747983474326, 0.
022040867441714847, 0.06956648786291249, 0.21959466510822148, 0.010356731875719217, 0.34777690340350187, 0.5
199600798403193, 0.023397549623259795, 1.0, 0.07525083612040134, 0.11725908455260796, 0.0003485677964364069,
0.022019995408152616, 0.05887271296478458, 0.30531610694829997, 0.0032461144993114303, 0.022270459810899375,
1.0]
F1 score when k is 45 = 0.21402228442257912

```
In [ ]: cluster_45,centroids_45 = k_means(data, 45)
```

```
In [20]: clusters_45 = predict(test, centroids_45)
```

```
In [21]: cont_45 = metrics.cluster.contingency_matrix(ground_truth,clusters_45)
    e_45=conditional_entropy(cont_45.T)
    p_45=precision(cont_45.T)
    r_45=recall(cont_45.T)
    f1_45=f1_score(p_45,r_45,cont_45.shape[1])
    print(f"conditional entropy when k is 45= {e_45}")
    print(f"precision when k is 45 = {p_45}")
    print(f"recall when k is 45 = {r_45}")
    print(f"F1 score when k is 45 = {f1_45}")
```

conditional entropy when k is 45= 0.20687376379107345 precision when k is 45 = [1.0, 0.9796167957602935, 1.0, 0.6798444588464031, 1.0, 1.0, 1.0, 1.0, 0.9215909090]909091, 1.0, 0.6153846153846154, 0.9370932754880694, 0.9388609715242882, 1.0, 0.9926470588235294, 0.98365561 69429097, 0.9715846994535519, 1.0, 0.86, 1.0, 0.6827133479212254, 1.0, 1.0, 1.0, 0.9992896718283847, 0.34571 997345719974, 0.6370510396975425, 1.0, 0.9171240985900334, 0.995835646862854, 0.9382716049382716, 1.0, 0.870 9677419354839, 0.9831908831908832, 0.9984496124031008, 0.9708454810495627, 0.97135416666666666, 0.72093023255 81395, 0.9769784172661871, 1.0, 1.0, 0.9976958525345622, 0.9, 1.0] recall when k is 45 = [0.09236671257131615, 0.11818807790674798, 0.043664294212185095, 0.021894763206779237,0.011115482982490656, 0.0905428815929536, 0.01335810147982718, 0.017699484460771013, 0.016927219218968503, 0.023272317421886415, 0.14507772020725387, 0.018033436997766692, 0.023397549623259795, 0.01849262173613575, 0.47093023255813954, 0.21016132205390517, 0.01855523783682244, 0.07589022230965965, 0.00634467833956325, 0.0 3359236671257131, 1.0, 0.05689716349063889, 0.05593704994677631, 0.03531548078729311, 0.34595711194176665, 0.5199600798403193, 0.3918604651162791, 0.017052451420341883, 0.17785059798376157, 0.0748679843877189, 0.95, 0.0474621286641747, 0.0941860465116279, 0.07202938782325562, 0.0134415896140761, 0.034751935881112896, 0.007 785268518711778, 0.7126436781609196, 0.04251633236626245, 0.05813495966948652, 0.008703637995449896, 0.00903 7590532445575, 0.00036559358180156395, 0.06597649809020893] F1 score when k is 45 = 0.18001787411451572

# **Spectral Clustering**

reading and encoding data

```
In [ ]: whole_data, whole_labels = import_data(path_of_whole_data, columns, ['service', 'flag', 'protocol_type'])
    whole_data['labels'] = whole_labels
    train_spectral, test_spectral = train_test_split(whole_data, train_size = 0.0025, random_state = 42, stratify
    train_spectral_labels = train_spectral['labels']
    le = LabelEncoder()
    train_spectral_labels_encoded = le.fit_transform(train_spectral_labels)
    train_spectral.drop(['labels'], axis = 1, inplace=True)
    k = len(np.unique(train_spectral_labels))
    train_spectral_labels_unique_alphabetically = sorted(train_spectral_labels.unique())
```

### **Implementation**

```
In []: def spectral_clustering(sim_matrix, k):
    degree_matrix = np.diag(np.sum(sim_matrix, axis=1))
    L = degree_matrix - sim_matrix
    La = np.dot(np.linalg.inv(degree_matrix), L)
    eigen_vals, eigen_vectors = np.linalg.eigh(La)
    idx = np.argsort(eigen_vals) #Index of arranged eigen values in ascending order
    eigen_vals = eigen_vals[idx]
    eigen_vectors = eigen_vectors[:,idx]
    U = eigen_vectors[:,:k]
    Y = normalize(U)
    #Y_df = pd.DataFrame(Y)
    #kms, centroids = k_means(Y_df, k)
    kms = KMeans(n_clusters=k)
    C = kms.fit_predict(Y)
    return C, Y
```

### Clustering using spectral clustering

```
In [ ]: sim_matrix = rbf_kernel(np.array(train_spectral), gamma=0.1)
C, Y = spectral_clustering(sim_matrix,k)
```

#### **Evaluation**

```
contg matrix spectral = contingency matrix(labels true = train spectral labels, labels pred = C).T
 In [ ]:
         pd.DataFrame(contg matrix spectral)
Out[112]:
            0 1
                  2 3
                         4 5 6 7 8 9 10
          0 2 9
                 20 4 2009 1 5 10 8 2 2
          1 0 0 143 0
                         0 0 1 0 0 0 0
          2 0 0 135 0
                        0 0 0 0 0 0
          3 0 0
                 63 0
                        1 0 1 1 0 0 0
          4 0 0
                 70 0
                        2 0 1 0 0 0 0
          5 0 0
                 36 0
                         6 0 0 0 0 0
          6 0 0
                 47 0
                         3 0 0 0 0 0 0
          7 0 0
                 28 0
                         8 0 1 1 0 0 0
          8 0 0
                 36 0
                        3 0 0 1 0 0 0
                 14 0
                         0 0 0 0 0 0
          10 0 0 13 0
                         0 0 0 0 0 0
 In [ ]: x=precision(contg matrix spectral)
         y=recall(contg matrix spectral)
         f1=f1 score(x,y,contg matrix spectral.shape[0])
         print("Precision for each Cluster: " + str(x))
         print("Recall for each Cluster: " +str(y))
         print("Average F1 score across all clusters:" + str(f1))
         Precision for each Cluster: [0.9695945945945946, 0.993055555555556, 1.0, 0.9545454545454546, 0.958904109589
         041, 0.8571428571428571, 0.94, 0.7368421052631579, 0.9, 1.0, 1.0]
         Recall for each Cluster: [0.9886811023622047, 0.236363636363636, 0.2231404958677686, 0.10413223140495868,
         23140495867768594, 0.021487603305785124]
         Average F1 score across all clusters:0.2320943551061556
         #num detected spectral = sum(contq matrix spectral[0])
         num detected spectral = sum(sum(contg_matrix_spectral[1:][:]))
```

```
In [ ]: | num_detected_spectral
 Out[99]: 615
  In [ ]: scaler 2 = RobustScaler()
          minmax scaler 2= MinMaxScaler()
          arr = np.array(train spectral)
          scaler 2.fit(arr)
          arr = standardization robust(train spectral, scaler 2)
          minmax scaler 2.fit(arr)
          train spectral means = standardization minmax(arr, train spectral.columns, minmax scaler 2)
          Kmeans on same K of spectral clustering
      1: C means, centroids means = k means(train spectral means, 11)
  In [ ]: contg_matrix_means = contingency_matrix(labels_true = train_spectral_labels, labels_pred = C_means).T
          pd.DataFrame(contg matrix means)
Out[113]:
              0 1
                          4 5 6 7 8 9 10
           0 0 0
                    0 0 91 0 0 0 0 0
                    0 0 494 0 0 0 0 0
           2 0 0 529 2
                          0 0 3 6 0 0 0
           3 0 0
                    0 0
                         98 0 0 0 0 0 0
           4 0 1 76 0 85 0 6 4 0 0 0
           5 0 1
                    0 0 211 1 0 1 1 0 2
           6 0 0
                    0 0 60 0 0 0 0 0
```

**7** 1 0

**8** 0 7

**9** 1 0

**10** 0 0

0 0 467 0 0 0 0 0 0

0 2 126 0 0 2 7 2 0

0 0 268 0 0 0 0 0 0

0 0 132 0 0 0 0 0 0

```
In [ ]: | num detected means = sum(contg matrix means[2]) + sum(contg matrix means[4])
          num detected means
Out[114]: 712
  In [ ]: entropy spectral = conditional entropy(contg matrix spectral)
          p s = precision(contg matrix spectral)
          r s = recall(contg matrix spectral)
          f1 s = f1 score(p s, r s, 11)
  In [ ]: print("Conditional entropy = " + str(entropy spectral))
          print("Precision = " + str(p_s))
          print("Recall = " + str(r s))
          print("F1 score = " + str(f1 s))
          Conditional entropy = 0.2759862350568628
          Precision = [0.9695945945945946, 0.993055555555556, 1.0, 0.9545454545454546, 0.958904109589041, 0.857142857
          1428571, 0.94, 0.7368421052631579, 0.9, 1.0, 1.0]
          Recall = [0.9886811023622047, 0.23636363636363636, 0.2231404958677686, 0.10413223140495868, 0.11570247933884
          298, 0.02975206611570248, 0.07768595041322314, 0.04628099173553719, 0.02975206611570248, 0.02314049586776859
          4, 0.021487603305785124]
          F1 score = 0.2320943551061556
```

# **Agglomerative Clustering**

Implementation

```
In [ ]: def agglomerative(data, k):
            clusters = [[i] for i in range(data.shape[0])] # each point in a separate cluster
            means = data.to numpy()
            distances = distance matrix(means, means) # compute distance matrix
            distances[distances == 0] = np.Inf
            while(len(means) > k):
                print(len(means))
                print(len(clusters))
                # getting the minimum distance
                (index1,index2) = np.unravel_index(distances.argmin(), distances.shape)
                # compute the mean of the closest clusters
                mean = [means[index1], means[index2]]
                mean = np.mean(mean, axis = 0)
                # concatenating the clusters
                clusters[index1] = clusters[index1] + clusters[index2]
                #updating the mean vector
                means[index1] = mean
                # deleting the old clusters and their means
                clusters.pop(index2)
                means = np.delete(means, index2, axis = 0)
                #updating the distance matrix
                distances = distance matrix(means, means)
                distances[distances == 0] = np.Inf
                #print('----')
            return clusters, means
```

## **Clustering using Agglomerative Clustering**

```
In [ ]: clusters, means = agglomerative(train_spectral, 11)
```

```
In [ ]: def mapping(data,n):
              map = np.zeros(n)
             for i in range(len(data)):
                 for j in range(len(data[i])):
                     map[data[i][j]] = i
              return map
  In [ ]: | agg cluster = mapping(clusters, len(train spectral))
          np.unique(agg cluster, return counts = True)
Out[121]: (array([ 0., 1., 2., 3., 4., 5., 6., 7., 8., 9., 10.]),
                                           6,
           array([2575, 9, 78,
                                                       3,
                                                                              1]))
                                      3,
                                                 8,
                                                             2,
                                                                        1,
```

### **Evaluation**

```
In [ ]: contg matrix agg = contingency matrix(labels true = train spectral labels, labels pred = agg cluster).T
        pd.DataFrame(contg matrix agg)
```

```
Out[127]:
            0 1
                  2 3
                         4 5 6 7 8 9 10
          0 0 9 605 4 1922 1 9 13 8 2 2
          1 0 0
                         9 0 0 0 0 0
          2 0 0
                  0 0
                        78 0 0 0 0 0 0
          3 2 0
                  0 0
                         1 0 0
                                0 0 0
          4 0 0
                  0 0
                         6 0 0
                                0 0 0 0
                                0 0 0 0
          5 0 0
                  0 0
                         8 0 0
          6 0 0
                  0 0
                         3 0 0
                               0 0 0
          7 0 0
                  0 0
                         2 0 0
                                0 0 0
          8 0 0
                  0 0
                         1 0 0
                               0 0 0
          9 0 0
                  0 0
                         1 0 0
                               0 0 0 0
          10 0 0
                  0 0
                         1 0 0 0 0 0 0
```

```
In []: entropy_agg = conditional_entropy(contg_matrix_agg)
    TP_agg, FP_agg, FN_agg, TN_agg = confusion_mat(contg_matrix_agg)
    p_agg = precision(contg_matrix_agg)
    r_agg = recall(contg_matrix_agg)
    f1_agg = f1_score(p_agg, r_agg, 11)

In []: print("Conditional entropy = " + str(entropy_agg))
    print("Precision = " + str(p_agg))
    print("Recall = " + str(r_agg))
    print("F1 score = " + str(f1_agg))

Conditional entropy = 0.9232042032831927
    Precision = [0.7464077669902912, 1.0, 1.0, 0.6666666666666666, 1.0, 1.0, 1.0, 1.0, 1.0, 1.0, 1.0]
    Recall = [0.9458661417322834, 0.004390243902439025, 0.038385826771653545, 0.0009813542688910696, 0.002952755
    905511811, 0.00392156862745098, 0.0014763779527559055, 0.0009813542688910696, 0.00012301636117603641, 0.0001
    2301636117603641, 0.00012301636117603641]
    F1 score = 0.08530598749559327
```

### **Describing agglomerative clustering**

Agglomerative clustering is a hierarchical clustering algorithm that begins with each data point as its own cluster and iteratively merges the two closest clusters until a stopping criterion is met. In this algorithm, the distance between clusters is defined as the distance between their closest points (i.e., single linkage), their furthest points (i.e., complete linkage), or the average distance between their points (i.e., average linkage).

At each iteration, the two closest clusters are merged into a single cluster, and the distance matrix between clusters is updated accordingly. This process continues until all points are in a single cluster or until a predefined number of clusters is reached.

One advantage of agglomerative clustering is that it does not require a predefined number of clusters, as the algorithm determines the number of clusters based on the merging process. Additionally, the hierarchical structure of the resulting clusters can provide insight into the relationships between data points.

### Disadvantages of agglomeartive clustering

One disadvantage of agglomerative clustering is that it can be computationally expensive for large datasets. Since the algorithm requires the computation of pairwise distances between all data points at each iteration, the time complexity can be O(n^3) or higher for large datasets, where n is the number of data points.

Another potential disadvantage of agglomerative clustering is that it is sensitive to the choice of linkage criterion. Different linkage criteria can result in different cluster structures, and the choice of linkage criterion may depend on the nature of the data and the goals of the analysis. Moreover, agglomerative clustering is a greedy algorithm and once a decision is made to merge two clusters, it cannot be undone, which can lead to suboptimal clustering in some cases.

Finally, agglomerative clustering can also suffer from the "chaining" effect, where small clusters may be merged into larger clusters too quickly, resulting in suboptimal clusters. This effect can be mitigated by using a stopping criterion that takes into account the size and density of the clusters, This is what happened with Neptune and Normal Classes

|--|