

## Import the needed packages

Entrée [1]:

```
import pandas as pd
import networkx as nt
import collections
import matplotlib.pyplot as plt
import numpy as np
from sklearn.cluster import KMeans
from sklearn.decomposition import PCA
from time import time
from sklearn import metrics
from sklearn.pipeline import make_pipeline
from sklearn.preprocessing import StandardScaler
```

## Import the data of the graph

Entrée [2]:

```
data_user_user = pd.read_csv("./data/Social_spammers_dataset/graphs/c_combined_edges.csv")
data_user_app = pd.read_csv("./data/Social_spammers_dataset/graphs/app_based_similarities.csv")
labels = pd.read_csv("./data/Social_spammers_dataset/users/coded_ids_labels_train.csv")
```

Let's see th overview of the graph data

Entrée [3]:

```
data_user_app.head()
```

Out[3]:

	user_id	app_id	weight
0	1	23	9
1	1	33	391
2	362	200	192
3	362	23	8
4	488	176	61

Entrée [4]:

```
data_user_user.head()
```

Out[4]:

	Source	Target	Weight	Sim
0	1	168	30	0.975
1	4	56	13	0.974
2	149	244	12	1.000
3	198	244	4	1.000
4	1	244	16	1.000

Get label in dictionary to facilitate the parsing

Entrée [5]:

```
labels.head()  
labels_dict=labels.to_dict()
```

## Creation the graph from dataframe

With networkx lib, we will build our graph. We will build two graph. The first graph is a graph on the relationship between user. The second one is about user and application used.

Entrée [6]:

```
graph_user_app = nt.from_pandas_edgelist(data_user_app,source="user_id",target="app"  
graph_user_user = nt.from_pandas_edgelist(data_user_user,source="Source",target="Ta
```

## The metric of graph

In this section we will do the graph exploration by computing some metrics

Entrée [7]:

```
print("Info graph user_app: ")
print(nt.info(graph_user_app))
print("Info graph user_app: ")
print(nt.info(graph_user_user))
```

Info graph user\_app:  
Name:  
Type: Graph  
Number of nodes: 767  
Number of edges: 2021  
Average degree: 5.2699  
Info graph user\_app:  
Name:  
Type: Graph  
Number of nodes: 295  
Number of edges: 1335  
Average degree: 9.0508

## Computing the diameter and radius of the graph

Entrée [8]:

```
print(f"radius of graph user app: {nt.radius(graph_user_app)}")
print(f"diameter of graph user app: {nt.diameter(graph_user_app)}")
```

radius of graph user app: 4  
diameter of graph user app: 6

## Computing of the density of graph

Entrée [9]:

```
density_user_user = nt.density(graph_user_user)
density_user_app = nt.density(graph_user_app)
print("The density of user_user_graph is {}".format(density_user_user))
print("The density of user_app_grap is {}".format(density_user_app))
```

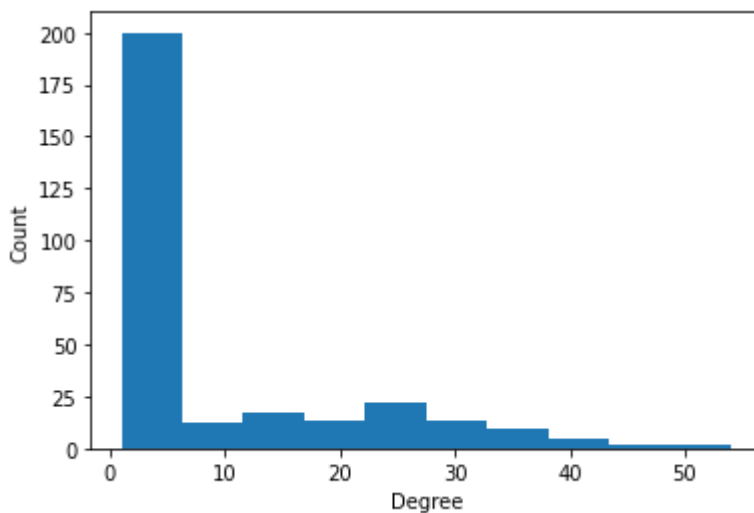
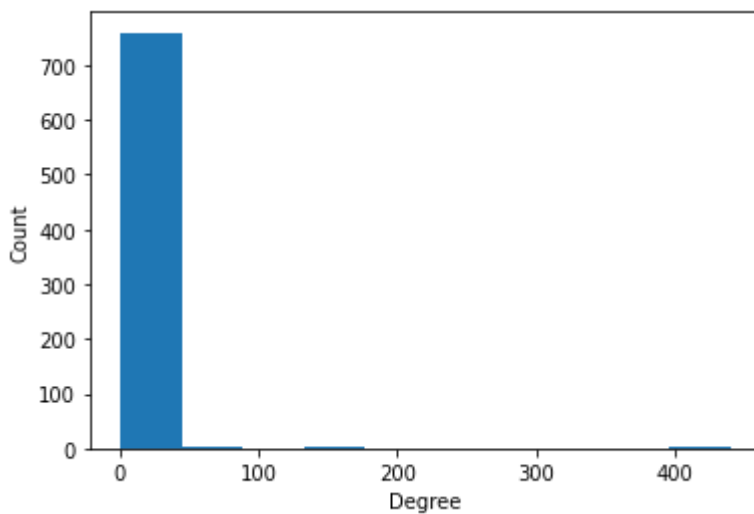
The density of user\_user\_graph is 0.030785195434105846  
The density of user\_app\_grap is 0.006879742375604658

## Degree ditribition analysis

Let'draw the histogram of degree distribution of our two graph

Entrée [10]:

```
def plot_degree_dist(G,name):  
    fig1 = plt.gcf()  
    degrees = [G.degree(n) for n in G.nodes()]  
    plt.hist(degrees)  
    plt.ylabel("Count")  
    plt.xlabel("Degree")  
    plt.show()  
    #plt.draw()  
    fig1.savefig(name, dpi=100)  
  
plot_degree_dist(graph_user_app, "graph_user_app_deg_dist.png")  
plot_degree_dist(graph_user_user, "graph_user_user_deg_dist.png")
```



## Compute the distribution of degree only on one class

First, we will separate the node in two sets: one for the spammers and another one for the non spammer And after that we will draw the degree distribution for each class of node.

In this section we also compute the mean of the degree on each class of node

Entrée [11]:

```
#In this section we look for the spammer node and the non spammer node
list_spam = []
list_non_spam = []
for node in graph_user_user.nodes():
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==1:
            list_spam.append(node)

for node in graph_user_user.nodes():
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==0:
            list_non_spam.append(node)
```

Entrée [12]:

```

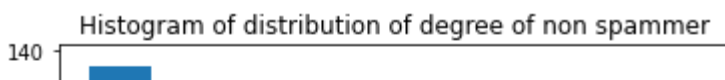
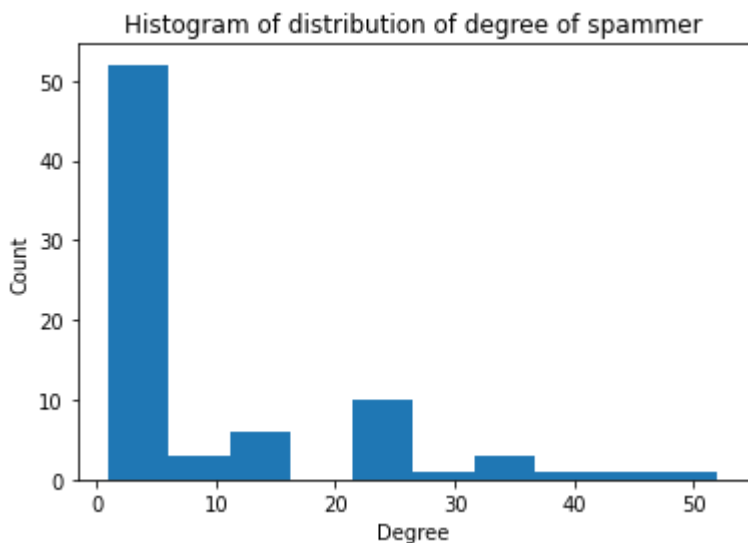
import numpy as np
def plot_degree_dist(G, list_node, title, name_file):
    fig1 = plt.gcf()
    degrees = [G.degree(n) for n in list_node]
    plt.hist(degrees)
    plt.ylabel("Count")
    plt.xlabel("Degree")
    plt.title(title)
    plt.show()
    fig1.savefig(name_file, dpi=100)

plot_degree_dist(graph_user_user, list_spam, "Histogram of distribution of degree of spammer", "spam.png")
plot_degree_dist(graph_user_user, list_non_spam, "Histogram of distribution of degree of non spammer", "non_spam.png")

def compute_mean_of_degree(G, list_node):
    degrees = [G.degree(n) for n in list_node]
    return np.mean(degrees)

print ("The mean of degree of the spammer: {}".format(compute_mean_of_degree(graph_user_user, list_spam)))
print ("The mean of degree of the non spammer: {}".format(compute_mean_of_degree(graph_user_user, list_non_spam)))

```



We can constate that the mean of the degree of group of spammer is greater than the non spammer mean of degree This observation allows us to say that the spammer share more edge than the non spammer.

Next, we will get the array of the degree for each class.

With these array, we will draw boxplot to see how it looks like.

Entrée [13]:

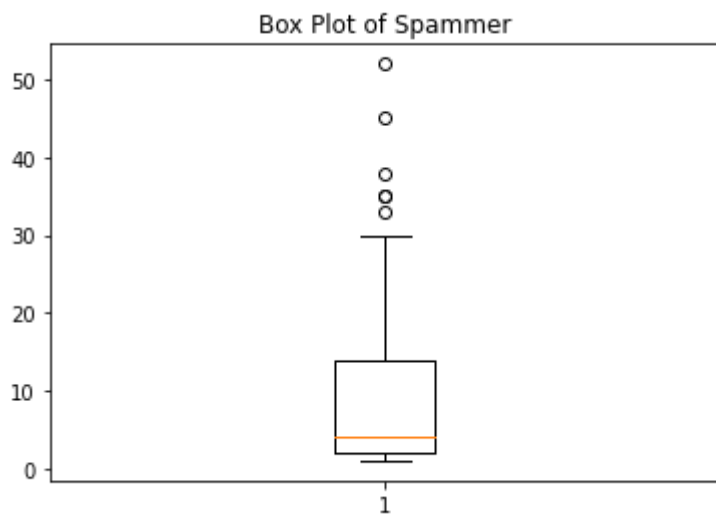
```

degrees_spammer = [graph_user_user.degree(n) for n in list_spam]
degrees_non_spammer = [graph_user_user.degree(n) for n in list_non_spam]

```

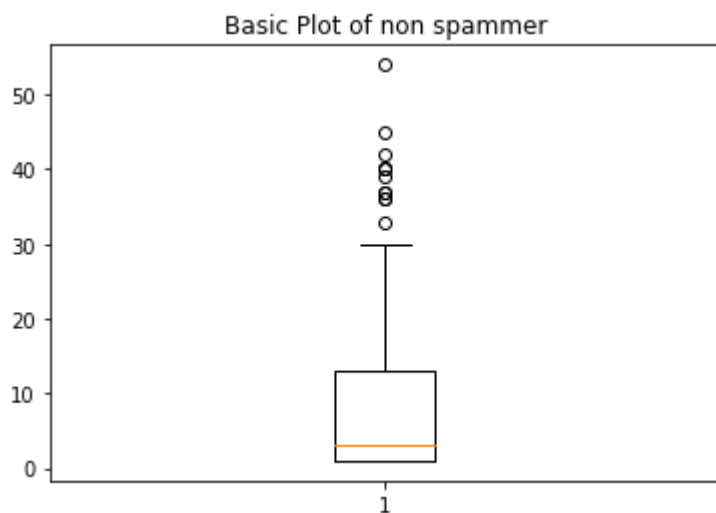
Entrée [14]:

```
fig1, ax1 = plt.subplots()
ax1.set_title('Box Plot of Spammer')
ax1.boxplot(degrees_spammer)
fig1.savefig("boxplot_spam.png", dpi=100)
```



Entrée [15]:

```
fig1, ax1 = plt.subplots()
ax1.set_title('Basic Plot of non spammer')
ax1.boxplot(degrees_non_spammer)
fig1.savefig("boxplot_non_spam.png", dpi=100)
```

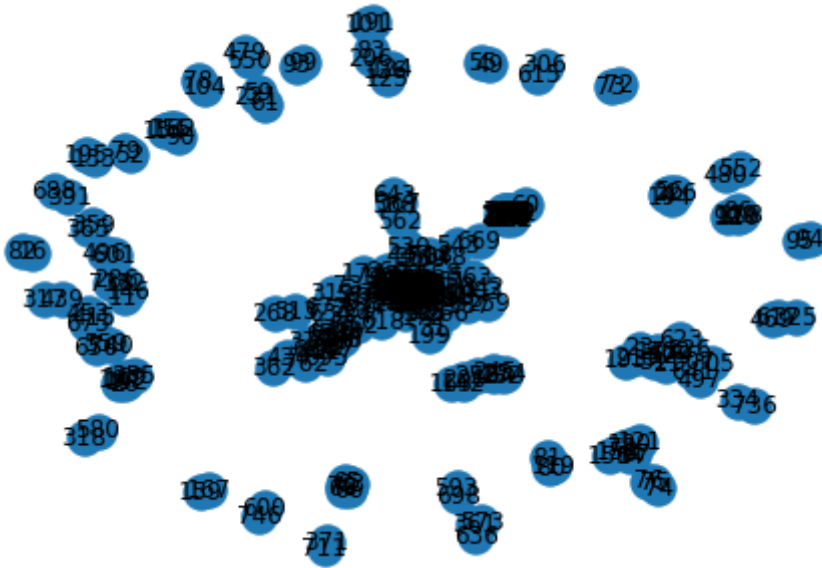


## Graph drawing

After computing some indicators, let dra the graphs of relationship between user

Entrée [16]:

```
nt.draw(graph_user_user, with_labels=True)
```

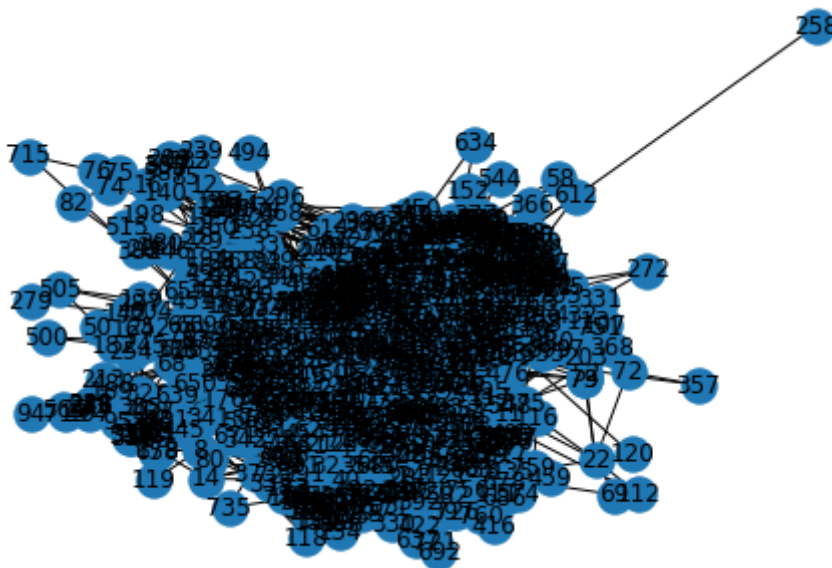


In the previous figure, we can notice that this graph are not connected.

Let draw graph for app and user interaction

Entrée [17]:

```
nt.draw(graph_user_app, with_labels=True)
```



Let compute the degree of centrality of each node



Entrée [18]:

```
degree_dict_user_user = nt.degree centrality(graph_user_user)
degree_dict_user_app = nt.degree centrality(graph_user_app)
```

After computing the degree of centrality of each

Entrée [19]:

```
print("The mean of the degree of centrality for the user relation graph is {}".format(
print("The mean of the degree of centrality for the user interaction with app graph
```

The mean of the degree of centrality for the user relation graph is 0.030785195434105843

The mean of the degree of centrality for the user interaction with app graph is 0.006879742375604658

Now, we will compute this mean with this mean for only the spammer and only for the non-spammer

Entrée [20]:

```
list_to_compute_mean = []
for node in degree_dict_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==1:
            list_to_compute_mean.append(degree_dict_user_user[node])
print(" The mean of the centrality of spammer : {}".format(np.mean(list_to_compute_
```

```
list_to_compute_mean = []
for node in degree_dict_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==0:
            list_to_compute_mean.append(degree_dict_user_user[node])
print(" The mean of the centrality of non spammer : {}".format(np.mean(list_to_comp
```

The mean of the centrality of spammer : 0.03296703296703297

The mean of the centrality of non spammer : 0.0312184279652455

The mean of centrality of the spammer is pretty greater than the non spammer.

## Computing of the betweenness of each node

More easily, this metric will show the best\_connector

Entrée [21]:

```
betweenness_user_user = nt.betweenness centrality(graph_user_user)
```

Entrée [22]:

```
betweenness_user_user = dict(sorted(betweenness_user_user.items(), key=lambda item: i
```

Let compute the mean of betweenness by class of node.

Entrée [23]:

```
list_to_compute_mean = []
for node in betweenness_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==1:
            list_to_compute_mean.append(betweenness_user_user[node])
print(" The mean of the betweenness of spammer : {}".format(np.mean(list_to_compute_mean)))

list_to_compute_mean = []
for node in betweenness_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==0:
            list_to_compute_mean.append(betweenness_user_user[node])
print(" The mean of the betweenness of non spammer : {}".format(np.mean(list_to_compute_mean)))
```

The mean of the betweenness of spammer : 0.0016196661726249728

The mean of the betweenness of non spammer : 0.001884250561132695

## Computing of average of clustering coefficient

Entrée [24]:

```
average_clustering_user_user = nt.average_clustering(graph_user_user)
average_clustering_user_app = nt.average_clustering(graph_user_app)
print ("The average of clustering coefficient of graph user relation is {}".format(average_clustering_user_user))
print ("The average of clustering coefficient of graph user app is {}".format(average_clustering_user_app))
```

The average of clustering coefficient of graph user relation is 0.47837313796999104

The average of clustering coefficient of graph user app is 0.17656147836380048

From the previous indicators, the graph on user relationship is mre clustable than the other graph.

Entrée [25]:

```
closeness_user_user = nt.closeness centrality(graph_user_user)
```

Entrée [26]:

```
closeness_user_user = dict(sorted(closeness_user_user.items(), key=lambda item: ite
```

Entrée [27]:

```
list_to_compute_mean_spam=[]
for node in closeness_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==1:
            list_to_compute_mean_spam.append(closeness_user_user[node])
list_to_compute_mean_non_spam=[]
for node in closeness_user_user:
    if labels_dict['label'].get(node) is not None:
        if labels_dict['label'].get(node) ==0:
            list_to_compute_mean_non_spam.append(closeness_user_user[node])
print("the mean of clossness of spammer {}",np.mean(list_to_compute_mean_spam))
print("the mean of clossness of non spammer {}",np.mean(list_to_compute_mean_non_s
```

```
the mean of clossness of spammer {} 0.08221274876513192
the mean of clossness of non spammer {} 0.08526386823516259
```

## Link analysis

Analysis of the influence. The interpretation of this metric is influence of on node in the network. At the end, we will display orded in most influenced node

Entrée [28]:

```
most_influential_link = nt.degree centrality(graph_user_user)
```

Entrée [29]:

```
#Adding the label
for item in most_influential_link:
    most_influential_link[item]=(most_influential_link[item],labels_dict['label'].g
```

## Page rank

Entrée [30]:

```
page_rank=nt.pagerank(graph_user_user)
```

Entrée [31]:

```
# the max page_rank
import operator
max_page_rank =max(page_rank.items(), key=operator.itemgetter(1))[0]
```

Let show the most page\_rank node label

Entrée [32]:

```
labels_dict['label'].get(max_page_rank)
```

Out[32]:

1

Let compute the most important connection and its label. A node is high eigenvector centrality if it is connected to many other nodes who are themselves well connected.

Entrée [33]:

```
most_important_linked = nt.eigenvector_centrality(graph_user_user)
count = 0
for w in sorted(most_important_linked, key=most_important_linked.get, reverse=True):
    print(w, most_important_linked[w], labels_dict['label'].get(w))
    count = count + 1
    if count == 10:
        break
```

```
240 0.2053466952222802 0
137 0.20007116848212567 1
33 0.19154650519803795 0
136 0.1909179357115181 1
257 0.18913960441815067 0
223 0.1854685477854571 0
267 0.184522637831872 0
463 0.18053552510982565 0
241 0.18025828236593705 0
17 0.17975094125267083 0
```

Detection of community with girvan\_newman on user relationship graph

Entrée [34]:

```
from networkx.algorithms import community
communities_generator = community.girvan_newman(graph_user_user)
top_level_communities = next(communities_generator)
next_level_communities = next(communities_generator)
coun = 0
print("The number of community with girvan_newman is : ", len(sorted(map(sorted, ne
for arr in sorted(map(sorted, next_level_communities)):
    #print("In the community {} there are {} there are following node :".format(cou
    for node in arr:
        pass
        #print(node)
    coun = coun + 1
```

The number of community with girvan\_newman is : 47

Generation of graph bipartite

Entrée [35]:

```
from networkx.algorithms.community.kernighan_lin import kernighan_lin_bisection
print(kernighan_lin_bisection(graph_user_user))
```

```
{513, 8, 10, 11, 17, 19, 533, 24, 25, 26, 30, 31, 33, 34, 547, 36, 3
8, 41, 45, 49, 562, 53, 568, 57, 56, 572, 61, 573, 584, 73, 72, 75, 7
6, 77, 78, 593, 82, 84, 86, 87, 600, 89, 90, 93, 94, 607, 98, 99, 102,
615, 104, 106, 108, 109, 110, 623, 122, 634, 129, 643, 133, 648, 136,
138, 656, 146, 147, 660, 149, 662, 156, 669, 158, 672, 162, 163, 165,
168, 170, 687, 179, 180, 182, 698, 699, 188, 187, 190, 191, 192, 193,
194, 195, 197, 200, 201, 205, 220, 223, 736, 737, 227, 740, 231, 232,
233, 746, 236, 238, 239, 240, 241, 245, 247, 761, 762, 254, 257, 260,
262, 264, 265, 267, 286, 293, 305, 306, 307, 310, 317, 318, 319, 321,
325, 334, 338, 359, 361, 371, 377, 403, 407, 415, 424, 435, 480, 497},
{1, 3, 4, 516, 9, 16, 529, 18, 530, 532, 20, 21, 23, 27, 543, 35, 550,
552, 42, 43, 47, 48, 559, 51, 563, 52, 55, 570, 59, 60, 63, 64, 65, 6
6, 67, 68, 580, 70, 71, 74, 588, 79, 80, 81, 83, 601, 91, 92, 95, 101,
103, 105, 617, 618, 114, 119, 121, 636, 125, 126, 124, 641, 130, 132,
646, 137, 142, 654, 144, 145, 151, 154, 155, 668, 159, 160, 675, 164,
677, 166, 167, 169, 171, 172, 173, 174, 688, 183, 696, 185, 695, 196,
198, 199, 711, 204, 718, 206, 208, 207, 211, 214, 215, 218, 219, 738,
226, 228, 229, 237, 244, 250, 251, 252, 255, 259, 261, 266, 268, 269,
280, 282, 295, 309, 315, 326, 339, 342, 350, 362, 365, 367, 384, 387,
391, 411, 434, 439, 444, 447, 448, 453, 462, 463, 465, 469, 479, 496})
```

## Detection of community with k-clique

With k=3, we will detect the communities

Entrée [36]:

```
communities_generator = nt.community.k_clique_communities(graph_user_user, 3)
top_level_communities = next(communities_generator)
next_level_communities = next(communities_generator)
print(next_level_communities)
```

```
frozenset({56, 194, 226, 4})
```

## Clustering based on similarity

Now let's use K-means to do clustering. This part is inspired from [link \(https://scikit-learn.org/stable/auto\\_examples/cluster/plot\\_kmeans\\_digits.html\)](https://scikit-learn.org/stable/auto_examples/cluster/plot_kmeans_digits.html).

We will use k-means with different configurations after that, we will compute some metric to evaluate the cluster

Entrée [37]:

```
filepath = "./data/Social_spammers_dataset/graphs/app_based_similarity/sim_matrix.tx
matrix = open(filepath).read()
matrix = [item.split(",") for item in matrix.split('\n')[:-1]]
```

Entrée [38]:

```
matrix_np = np.array(matrix)
matrix_np = matrix_np.astype(np.float32)
```

Entrée [39]:

```
columns = [ index for index in range(1,768) ]
rows = [ index for index in range(1,768) ]
dataframe_from_sim_matrix = pd.DataFrame(data=matrix_np, index=rows, columns=columns)
```

Entrée [40]:

```
from sklearn.cluster import KMeans
```

## Setting our benchmarck

This code allows us to make comparasion beetween the variants of k-means, and adding PCA

Entrée [41]:

```
def bench_k_means(kmeans, name, data, labels):
    t0 = time()
    estimator = make_pipeline(StandardScaler(), kmeans).fit(data)
    fit_time = time() - t0
    results = [name, fit_time, estimator[-1].inertia_]

    # Define the metrics which require only the true labels and estimator
    # labels
    clustering_metrics = [
        metrics.homogeneity_score,
        metrics.completeness_score,
        metrics.v_measure_score,
        metrics.adjusted_rand_score,
        metrics.adjusted_mutual_info_score,
    ]
    results += [m(labels, estimator[-1].labels_) for m in clustering_metrics]

    # The silhouette score requires the full dataset
    results += [
        metrics.silhouette_score(data, estimator[-1].labels_,
                                metric="euclidean", sample_size=300,)
    ]

    # Show the results
    formatter_result = ("{:9s}\t{:.3f}s\t{:.0f}\t{:.3f}\t{:.3f}"
                        "\t{:.3f}\t{:.3f}\t{:.3f}\t{:.3f}")
    print(formatter_result.format(*results))
```

Entrée [42]:

```
def benchmark_k_mean(dataframe, labels):
    for i in range(2,10):
        print ("With K = {}".format(i))
        print(82 * '_')
        print('init\t\ttime\tinertia\thomo\tcompl\tv-meas\tARI\tAMI\tsilhouette')

        kmeans = KMeans(init="k-means++", n_clusters=i, n_init=4,
                        random_state=0)
        bench_k_means(kmeans=kmeans, name="k-means++", data=dataframe, labels=labels)

        kmeans = KMeans(init="random", n_clusters=i, n_init=4, random_state=0)
        bench_k_means(kmeans=kmeans, name="random", data=dataframe, labels=labels)

        pca = PCA(n_components=i).fit(dataframe)
        kmeans = KMeans(init=pca.components_, n_clusters=i, n_init=1)
        bench_k_means(kmeans=kmeans, name="PCA-based", data=dataframe, labels=labels)

    print(82 * '_')
```

## The matrix from sim\_matrix.txt

Entrée [43]:

```
dataframe_from_sim_matrix.shape
```

Out[43]:

```
(767, 767)
```

## Clustering of the data from sim matrix

In this section we will make cluster(community) with the file.

Entrée [44]:

```
benchmark_k_mean(dataframe_from_sim_matrix,[index for index in range(1,768)])
```

With K = 2

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.559	0.093s	378817	0.102	1.000	0.185	0.000	0.000
random 0.587	0.079s	378817	0.102	1.000	0.185	0.000	0.000
PCA-based 0.539	0.035s	378817	0.102	1.000	0.185	0.000	0.000

With K = 3

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.528	0.068s	291944	0.157	1.000	0.271	0.000	-0.000
random 0.563	0.051s	291944	0.157	1.000	0.271	0.000	-0.000
PCA-based 0.551	0.026s	291944	0.157	1.000	0.271	0.000	-0.000

With K = 4

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.533	0.070s	247854	0.189	1.000	0.318	0.000	0.000
random 0.543	0.044s	251112	0.188	1.000	0.317	0.000	0.000
PCA-based 0.551	0.025s	251112	0.188	1.000	0.317	0.000	0.000

With K = 5

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.505	0.080s	226393	0.208	1.000	0.345	0.000	0.000
random 0.573	0.055s	208387	0.219	1.000	0.359	0.000	0.000
PCA-based 0.578	0.028s	208386	0.218	1.000	0.359	0.000	-0.000

With K = 6

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
-----------------	------	---------	------	-------	--------	-----	-----



k-means++ 0.566	0.086s	192941	0.235	1.000	0.380	0.000	-0.000
random 0.582	0.055s	188108	0.237	1.000	0.383	0.000	0.000
PCA-based 0.574	0.026s	192941	0.235	1.000	0.380	0.000	-0.000

---

With K = 7

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.587	0.099s	173039	0.252	1.000	0.403	0.000	-0.000
random 0.546	0.069s	178700	0.259	1.000	0.412	0.000	-0.000
PCA-based 0.511	0.027s	178684	0.249	1.000	0.398	0.000	-0.000

---

With K = 8

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.581	0.116s	161308	0.263	1.000	0.417	0.000	-0.000
random 0.577	0.061s	162884	0.283	1.000	0.441	0.000	-0.000
PCA-based 0.567	0.026s	166754	0.260	1.000	0.412	0.000	-0.000

---

With K = 9

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.518	0.131s	155523	0.288	1.000	0.447	0.000	0.000
random 0.568	0.064s	153667	0.284	1.000	0.442	0.000	-0.000
PCA-based 0.630	0.027s	147732	0.276	1.000	0.433	0.000	-0.000

---

From the matrix file, we made clusters (communities). For the K=9, that is the best number of K according to the silhouette metric

## Now we will do the clustering with the matrix from the graph user interaction

We also apply a clustering algorithm ( k-means, hierarchical algorithms). we also generated the matrix with similarity from graph of user relationship.

Entrée [45]:

```
graph_user_user_matrix = nt.adjacency_matrix(graph_user_user,weight='Sim')
```

Entrée [46]:

```
print(graph_user_user_matrix.todense())
labels = graph_user_user.nodes()
graph_user_user_matrix = pd.DataFrame(graph_user_user_matrix.todense(), index=list(
[[0.    0.975 0.    ... 0.    0.    0.    ]
 [0.975 0.    0.    ... 0.    0.    0.    ]
 [0.    0.    0.    ... 0.    0.    0.    ]
 ...
 [0.    0.    0.    ... 0.    0.    0.    ]
 [0.    0.    0.    ... 0.    0.    0.    ]
 [0.    0.    0.    ... 0.    0.    0.    ]])
```

Let show the shape on second matrix

Entrée [47]:

```
graph_user_user_matrix.shape
```

Out[47]:

(295, 295)

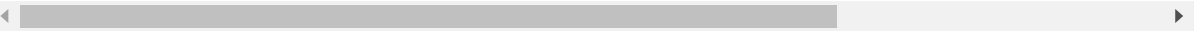
Entrée [48]:

```
graph_user_user_matrix
```

Out[48]:

	1	168	4	56	149	244	198	48	25	232	...	167	365	359	...
1	0.000	0.975	0.000	0.000	1.000	1.000	1.000	1.000	0.999	0.943	...	0.0	0.0	0.0	...
168	0.975	0.000	0.000	0.000	0.971	0.971	0.971	0.971	0.982	0.982	...	0.0	0.0	0.0	...
4	0.000	0.000	0.000	0.974	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
56	0.000	0.000	0.974	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
149	1.000	0.971	0.000	0.000	0.000	1.000	1.000	1.000	0.998	0.936	...	0.0	0.0	0.0	...
...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...
439	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
317	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
174	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
677	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...
193	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	...	0.0	0.0	0.0	...

295 rows × 295 columns



## Entrée [49]:

```
benchmark_k_mean(graph_user_user_matrix,graph_user_user.nodes())
```

With K = 2

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.493	0.034s	79735	0.059	1.000	0.112	0.000	-0.000
random 0.493	0.022s	79735	0.059	1.000	0.112	0.000	-0.000
PCA-based 0.493	0.021s	79735	0.059	1.000	0.112	0.000	-0.000

With K = 3

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.481	0.033s	73186	0.106	1.000	0.191	0.000	-0.000
random 0.095	0.024s	78451	0.076	1.000	0.142	0.000	-0.000
PCA-based 0.481	0.012s	73171	0.107	1.000	0.193	0.000	-0.000

With K = 4

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.447	0.026s	71016	0.154	1.000	0.267	0.000	-0.000
random 0.138	0.021s	76170	0.129	1.000	0.229	0.000	-0.000
PCA-based 0.424	0.011s	71009	0.161	1.000	0.277	0.000	-0.000

With K = 5

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.127	0.028s	70059	0.174	1.000	0.297	0.000	-0.000
random 0.051	0.019s	76395	0.118	1.000	0.210	0.000	-0.000
PCA-based 0.367	0.010s	70086	0.191	1.000	0.320	0.000	-0.000

With K = 6

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
-----------------	------	---------	------	-------	--------	-----	-----

k-means++ 0.136	0.027s	68877	0.188	1.000	0.317	0.000	-0.000
random -0.103	0.025s	75790	0.127	1.000	0.226	0.000	-0.000
PCA-based 0.217	0.014s	68864	0.203	1.000	0.337	0.000	-0.000

---

With K = 7

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.104	0.029s	68324	0.198	1.000	0.330	0.000	-0.000
random -0.063	0.023s	73679	0.177	1.000	0.301	0.000	-0.000
PCA-based 0.218	0.012s	68440	0.219	1.000	0.359	0.000	-0.000

---

With K = 8

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ -0.020	0.034s	68672	0.151	1.000	0.262	0.000	-0.000
random -0.078	0.021s	73223	0.163	1.000	0.280	0.000	-0.000
PCA-based 0.217	0.011s	67831	0.227	1.000	0.370	0.000	-0.000

---

With K = 9

---

init silhouette	time	inertia	homo	compl	v-meas	ARI	AMI
k-means++ 0.127	0.031s	67201	0.207	1.000	0.342	0.000	-0.000
random -0.157	0.022s	72923	0.167	1.000	0.286	0.000	-0.000
PCA-based 0.214	0.012s	67339	0.254	1.000	0.405	0.000	-0.000

---

The best silhouette is obtained when we uses k as 2 . Let draw to get more insight about these clusters.

Entrée [50]:

```

import matplotlib.pyplot as plt

reduced_data = PCA(n_components=2).fit_transform(graph_user_user_matrix)
kmeans = KMeans(init="k-means++", n_clusters=2, n_init=4)
kmeans.fit(reduced_data)

h = .02
x_min, x_max = reduced_data[:, 0].min() - 1, reduced_data[:, 0].max() + 1
y_min, y_max = reduced_data[:, 1].min() - 1, reduced_data[:, 1].max() + 1
xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))

# Obtain labels for each point in mesh. Use last trained model.
Z = kmeans.predict(np.c_[xx.ravel(), yy.ravel()])

# Put the result into a color plot
Z = Z.reshape(xx.shape)
plt.figure(1)
#plt.figure(figsize=(200,100))
plt.clf()
plt.imshow(Z, interpolation="nearest",
           extent=(xx.min(), xx.max(), yy.min(), yy.max()),
           cmap=plt.cm.Paired, aspect="auto", origin="lower")

plt.plot(reduced_data[:, 0], reduced_data[:, 1], 'k.', markersize=2)
# Plot the centroids as a white X
centroids = kmeans.cluster_centers_
plt.scatter(centroids[:, 0], centroids[:, 1], marker="x", s=169, linewidths=3,
           color="w", zorder=10)
plt.title("K-means clustering user (PCA-reduced data)\n"
          "Centroids are marked with white cross")
plt.xlim(x_min, x_max)
plt.ylim(y_min, y_max)
plt.xticks(())
plt.yticks(())

plt.show()

```

K-means clustering user (PCA-reduced data)  
Centroids are marked with white cross



## Hierarchical Clustering

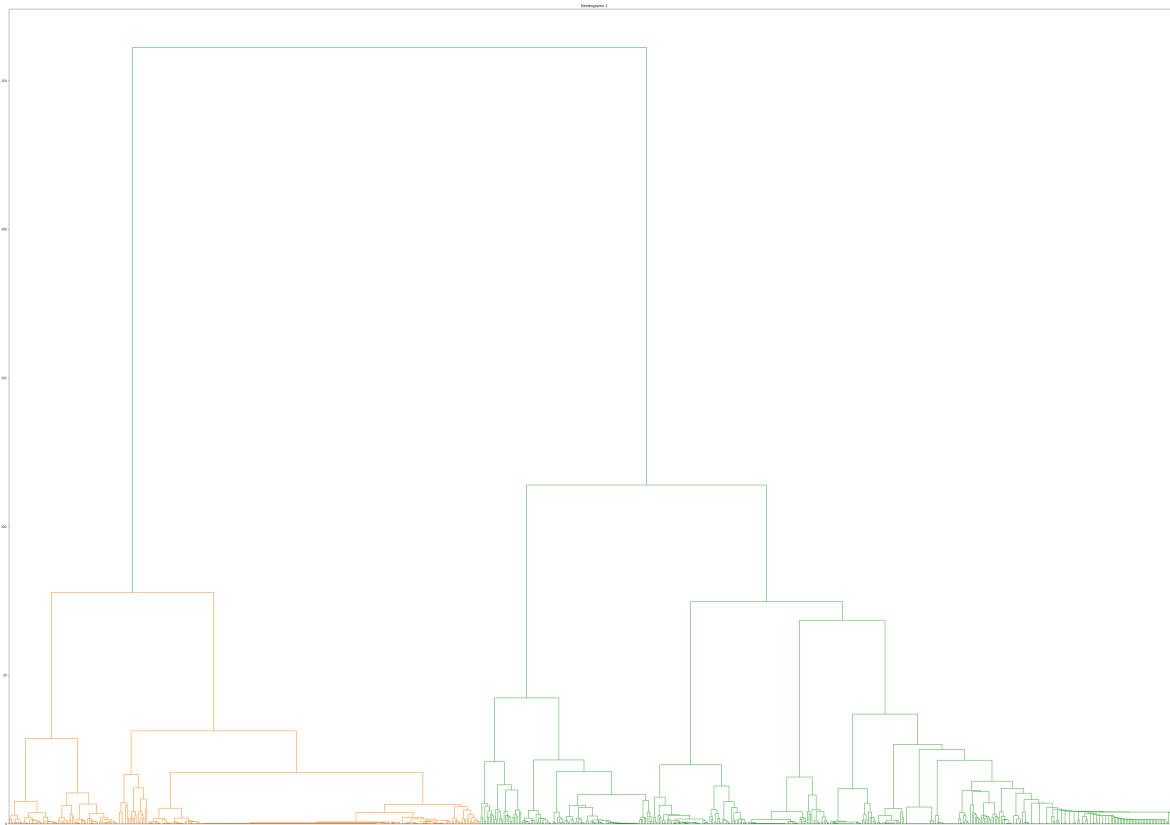
To end this study let uild the dendrogram from Hierarchical Clustering

The following section will compute the cluster from the graph matrix of similarity. From the previous section the right number of cluster is 2 for this matrix

Entrée [51]:

```
import scipy.cluster.hierarchy as shc

plt.figure(figsize=(70, 50))
plt.title("Dendograms 1")
dend = shc.dendrogram(shc.linkage(dataframe_from_sim_matrix, method='ward'))
```



This section will compute the cluter from the sim\_matrix.txt file.

Let's build the dendrogram before make cluster.

According to the dendrogram, the value 2 is right for the number of cluster.

Entrée [52]:

```
from sklearn.cluster import AgglomerativeClustering

cluster_AgglomerativeClustering = AgglomerativeClustering(n_clusters=2, affinity='e
cluster_AgglomerativeClustering.fit_predict(dataframe_from_sim_matrix)
```

Out[52]:

```
array([0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 0,
0,
      0, 1, 0, 0, 1, 0, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 0, 1, 1, 1,
0,
      1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1, 0, 0, 0,
0,
      0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1,
1,
      1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 1, 1, 0, 0, 0,
0,
      0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0,
0,
      0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0,
0,
      1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 1, 0, 1, 0,
0,
      0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0,
      1, 0, 1, 1, 1, 1, 1, 0, 1, 1, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1,
1,
      0, 1, 1, 0, 0, 0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1,
1,
      0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 1, 1, 0, 1, 1, 0,
1,
      1, 1, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 0, 1,
0,
      0, 1, 0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 1, 0,
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      1, 1, 0, 1, 1, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0,
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      0, 1, 0, 0, 0, 0, 1, 0, 0, 1, 1, 1, 0, 0, 1, 1, 1, 1, 1, 0, 0,
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      0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 1, 0, 0, 0,
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      1, 0, 0, 1, 0, 1, 1, 1, 0, 1, 0, 0, 1, 0, 1, 1, 0, 0, 0, 1, 1,
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      1, 1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1, 1, 0, 1, 1, 0, 0,
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      0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 0,
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      1, 0, 1, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1, 0,
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      0, 1, 0, 1, 1, 0, 0, 1, 1, 1, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0,
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      0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 1, 1, 0, 0, 1, 1, 0, 1, 0,
1,
```

```
1, 0, 1, 1, 1, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1, 0, 1, 0, 1, 1,
1,
0, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0, 1, 1, 1,
0,
1, 1, 1, 0, 0, 0, 1, 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 1, 1, 1, 0,
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0, 1, 0, 0, 1, 1, 0, 1, 0, 0, 0, 0, 1, 1, 1, 1, 1, 0, 1, 0, 1,
0,
0, 1, 0, 0, 1, 1, 1, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 1,
0,
0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0,
1,
0, 0, 0, 0, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 1, 1, 0, 0])
```

We call add this group to the dataset for the prediction task for the spammer. We think that can improve the accuracy of model.

Entrée [ ]:

Let build the dendrogram of this two dataset



Entrée [53]:

```
import scipy.cluster.hierarchy as shc

plt.figure(figsize=(70, 50))
plt.title("Dendrogram 2 ")
dend = shc.dendrogram(shc.linkage(graph_user_user_matrix, method='ward'))
```

<ipython-input-53-7f43d5beea76>:5: ClusterWarning: scipy.cluster: The symmetric non-negative hollow observation matrix looks suspiciously like an uncondensed distance matrix

```
dend = shc.dendrogram(shc.linkage(graph_user_user_matrix, method='ward'))
```

