Social network Graph Link Prediction - Facebook Challenge

Problem statement:

Given a directed social graph, have to predict missing links to recommend users (Link Prediction in graph)

Data Overview

Taken data from facebook's recruting challenge on kaggle https://www.kaggle.com/c/FacebookRecruiting

data contains two columns source and destination eac edge in graph

```
Data columns (total 2 columns):source_node int64destination node int64
```

Mapping the problem into supervised learning problem:

- Generated training samples of good and bad links from given directed graph and for each link got some features like no of followers, is he followed back, page rank, katz score, adar index, some svd fetures of adj matrix, some weight features etc. and trained ml model based on these features to predict link.
- Some reference papers and videos :
 - https://www.cs.cornell.edu/home/kleinber/link-pred.pdf
 - https://www3.nd.edu/~dial/publications/lichtenwalter2010new.pdf

- https://kaggle2.blob.core.windows.net/forum-messageattachments/2594/supervised link prediction.pdf
- https://www.youtube.com/watch?v=2M77Hgy17cg

Business objectives and constraints:

- No low-latency requirement.
- · Probability of prediction is useful to recommend highest probability links

Performance metric for supervised learning:

- Both precision and recall is important so F1 score is good choice
- Confusion matrix

```
In [1]: #Importing Libraries
        # please do go through this python notebook:
        import warnings
        warnings.filterwarnings("ignore")
        import csv
        import pandas as pd#pandas to create small dataframes
        import datetime #Convert to unix time
        import time #Convert to unix time
        # if numpy is not installed already : pip3 install numpy
        import numpy as np#Do aritmetic operations on arrays
        # matplotlib: used to plot graphs
        import matplotlib
        import matplotlib.pylab as plt
        import seaborn as sns#Plots
        from matplotlib import rcParams#Size of plots
        from sklearn.cluster import MiniBatchKMeans, KMeans#Clustering
        from sklearn.model selection import train test split
        import math
        import pickle
```

```
import os
        # to install xgboost: pip3 install xgboost
        import xgboost as xgb
        import warnings
        import networkx as nx
        import pdb
        import pickle
        from pandas import HDFStore,DataFrame
        from pandas import read hdf
        from scipy.sparse.linalg import svds, eigs
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import f1 score
        import tables
        from tqdm import tqdm
        from sklearn.model selection import RandomizedSearchCV
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import f1 score, auc, roc auc score, roc curve
        from xqboost import XGBClassifier
        pd.set option('max colwidth', 800)
In [2]: #reading graph
        if not os.path.isfile('data/after eda/train woheader.csv'):
            traincsv = pd.read csv('data/train.csv')
            print(traincsv[traincsv.isna().any(1)])
            print(traincsv.info())
            print("Number of diplicate entries: ",sum(traincsv.duplicated()))
            traincsv.to csv('data/after eda/train woheader.csv',header=False,in
        dex=False)
            print("saved the graph into file")
        g=nx.read edgelist('data/after eda/train woheader.csv',delimiter=',',cr
        eate using=nx.DiGraph(),nodetype=int)
        print(nx.info(g))
        Name:
        Type: DiGraph
        Number of nodes: 1862220
```

........

Number of edges: 9437519 Average in degree: 5.0679 Average out degree: 5.0679

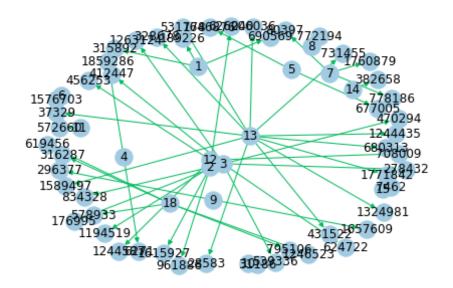
Displaying a sub graph

Name:

Type: DiGraph

Number of nodes: 66 Number of edges: 50

Average in degree: 0.7576 Average out degree: 0.7576



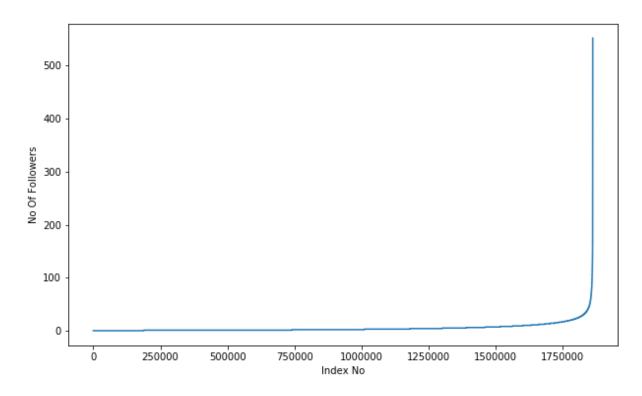
1. Exploratory Data Analysis

```
In [4]: # No of Unique persons
print("The number of unique persons",len(g.nodes()))
```

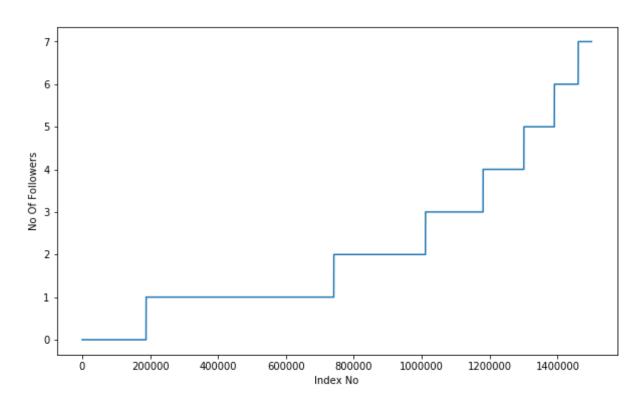
The number of unique persons 1862220

1.1 No of followers for each person

```
In [5]: indegree_dist = list(dict(g.in_degree()).values())
    indegree_dist.sort()
    plt.figure(figsize=(10,6))
    plt.plot(indegree_dist)
    plt.xlabel('Index No')
    plt.ylabel('No Of Followers')
    plt.show()
```



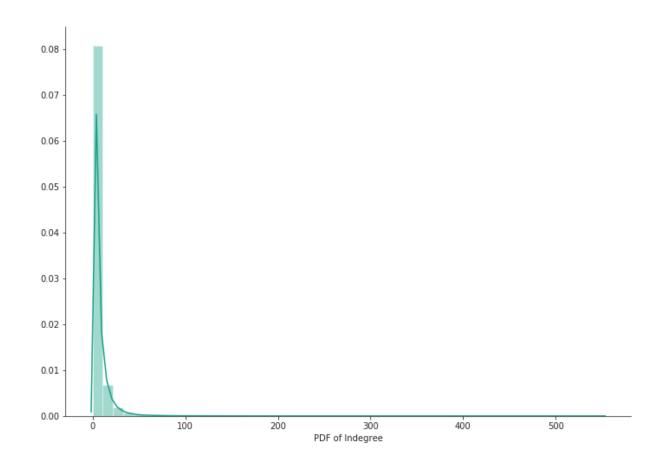
```
In [6]: indegree_dist = list(dict(g.in_degree()).values())
    indegree_dist.sort()
    plt.figure(figsize=(10,6))
    plt.plot(indegree_dist[0:1500000])
    plt.xlabel('Index No')
    plt.ylabel('No Of Followers')
    plt.show()
```



```
In [7]: plt.boxplot(indegree_dist)
  plt.ylabel('No Of Followers')
  plt.show()
```

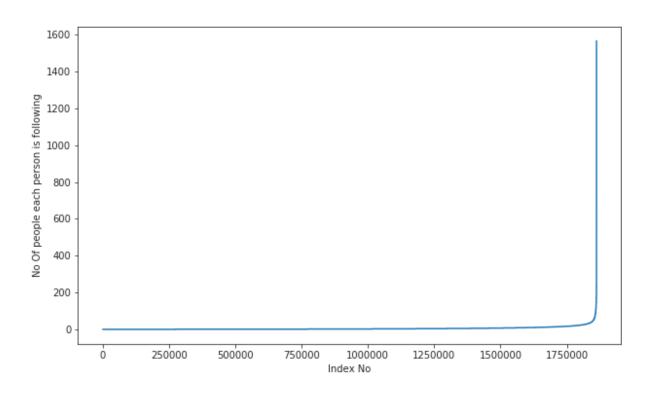
```
In [8]: ### 90-100 percentile
        for i in range(0,11):
             print(90+i, 'percentile value is', np.percentile(indegree dist, 90+i))
        90 percentile value is 12.0
        91 percentile value is 13.0
        92 percentile value is 14.0
        93 percentile value is 15.0
        94 percentile value is 17.0
        95 percentile value is 19.0
        96 percentile value is 21.0
        97 percentile value is 24.0
        98 percentile value is 29.0
        99 percentile value is 40.0
        100 percentile value is 552.0
        99% of data having followers of 40 only.
In [9]: ### 99-100 percentile
        for i in range(10,110,10):
             print(99+(i/100), 'percentile value is', np.percentile(indegree dist,
        99+(i/100))
```

```
99.1 percentile value is 42.0
         99.2 percentile value is 44.0
         99.3 percentile value is 47.0
         99.4 percentile value is 50.0
         99.5 percentile value is 55.0
         99.6 percentile value is 61.0
         99.7 percentile value is 70.0
         99.8 percentile value is 84.0
         99.9 percentile value is 112.0
         100.0 percentile value is 552.0
In [10]: %matplotlib inline
         sns.set style('ticks')
         fig, ax = plt.subplots()
         fig.set size inches(11.7, 8.27)
         sns.distplot(indegree dist, color='#16A085')
         plt.xlabel('PDF of Indegree')
         sns.despine()
         #plt.show()
```

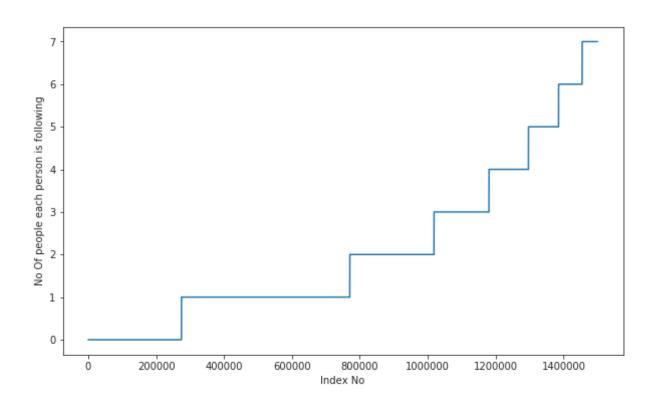


1.2 No of people each person is following

```
In [11]:    outdegree_dist = list(dict(g.out_degree()).values())
    outdegree_dist.sort()
    plt.figure(figsize=(10,6))
    plt.plot(outdegree_dist)
    plt.xlabel('Index No')
    plt.ylabel('No Of people each person is following')
    plt.show()
```



```
In [12]: indegree_dist = list(dict(g.in_degree()).values())
    indegree_dist.sort()
    plt.figure(figsize=(10,6))
    plt.plot(outdegree_dist[0:1500000])
    plt.xlabel('Index No')
    plt.ylabel('No Of people each person is following')
    plt.show()
```



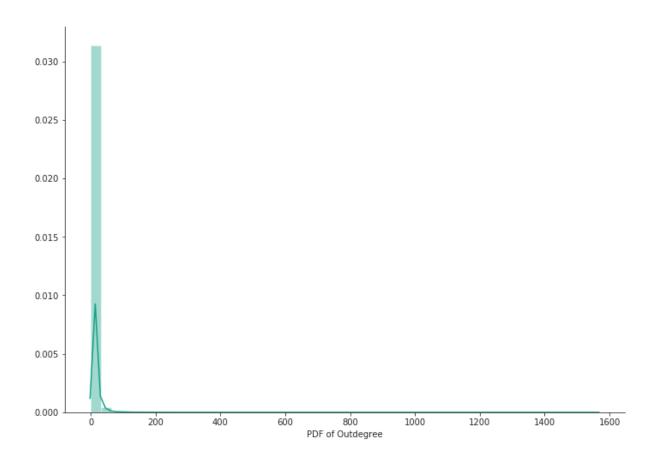
```
In [13]: plt.boxplot(indegree_dist)
  plt.ylabel('No Of people each person is following')
  plt.show()
```

```
No Of people each person is following and the search person of the searc
```

```
In [14]: ### 90-100 percentile
         for i in range(0,11):
             print(90+i, 'percentile value is', np.percentile(outdegree dist, 90+i
         90 percentile value is 12.0
         91 percentile value is 13.0
         92 percentile value is 14.0
         93 percentile value is 15.0
         94 percentile value is 17.0
         95 percentile value is 19.0
         96 percentile value is 21.0
         97 percentile value is 24.0
         98 percentile value is 29.0
         99 percentile value is 40.0
         100 percentile value is 1566.0
In [15]: ### 99-100 percentile
         for i in range(10,110,10):
             print(99+(i/100), 'percentile value is', np.percentile(outdegree dist
          ,99+(i/100)))
         99.1 percentile value is 42.0
         99.2 percentile value is 45.0
```

```
99.3 percentile value is 48.0
99.4 percentile value is 52.0
99.5 percentile value is 56.0
99.6 percentile value is 63.0
99.7 percentile value is 73.0
99.8 percentile value is 90.0
99.9 percentile value is 123.0
100.0 percentile value is 1566.0

In [16]: sns.set_style('ticks')
fig, ax = plt.subplots()
fig.set_size_inches(11.7, 8.27)
sns.distplot(outdegree_dist, color='#16A085')
plt.xlabel('PDF of Outdegree')
sns.despine()
```



No of persons those are not following anyone are 274512 and % is 14.741 115442858524

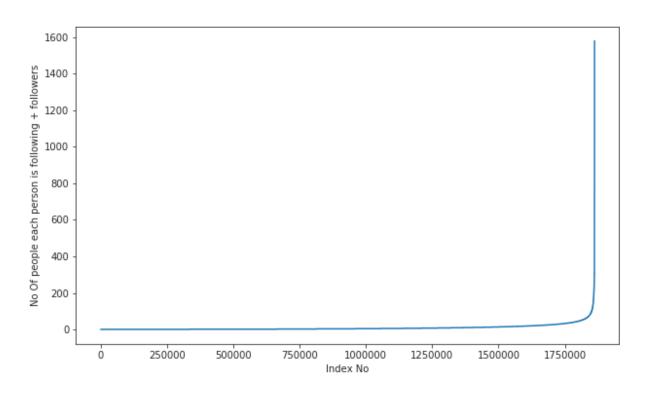
No of norcone baying zero followers are 1990/3 and & is 10 007796512971

No of persons those are not not following anyone and also not having an y followers are $\boldsymbol{\theta}$

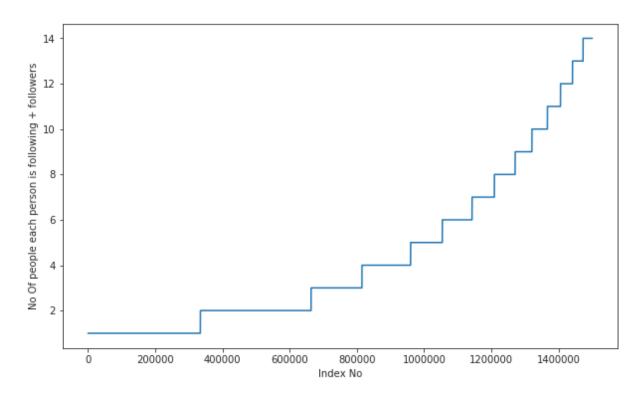
1.3 both followers + following

```
In [20]: from collections import Counter
dict_in = dict(g.in_degree())
dict_out = dict(g.out_degree())
d = Counter(dict_in) + Counter(dict_out)
in_out_degree = np.array(list(d.values()))
```

```
In [21]: in_out_degree_sort = sorted(in_out_degree)
    plt.figure(figsize=(10,6))
    plt.plot(in_out_degree_sort)
    plt.xlabel('Index No')
    plt.ylabel('No Of people each person is following + followers')
    plt.show()
```



```
In [22]: in_out_degree_sort = sorted(in_out_degree)
    plt.figure(figsize=(10,6))
    plt.plot(in_out_degree_sort[0:1500000])
    plt.xlabel('Index No')
    plt.ylabel('No Of people each person is following + followers')
    plt.show()
```



```
In [23]: ### 90-100 percentile
for i in range(0,11):
    print(90+i, 'percentile value is',np.percentile(in_out_degree_sort,9
    0+i))

90 percentile value is 24.0
91 percentile value is 26.0
92 percentile value is 28.0
93 percentile value is 31.0
94 percentile value is 33.0
95 percentile value is 37.0
96 percentile value is 41.0
97 percentile value is 48.0
98 percentile value is 58.0
99 percentile value is 79.0
100 percentile value is 1579.0
```

```
In [24]: ### 99-100 percentile
         for i in range(10,110,10):
             print(99+(i/100), 'percentile value is', np.percentile(in out degree
         sort,99+(i/100))
         99.1 percentile value is 83.0
         99.2 percentile value is 87.0
         99.3 percentile value is 93.0
         99.4 percentile value is 99.0
         99.5 percentile value is 108.0
         99.6 percentile value is 120.0
         99.7 percentile value is 138.0
         99.8 percentile value is 168.0
         99.9 percentile value is 221.0
         100.0 percentile value is 1579.0
In [25]: print('Min of no of followers + following is',in out degree.min())
         print(np.sum(in out degree==in out degree.min()),' persons having minim
         um no of followers + following')
         Min of no of followers + following is 1
         334291 persons having minimum no of followers + following
In [26]: print('Max of no of followers + following is',in out degree.max())
         print(np.sum(in out degree==in out degree.max()), ' persons having maxim
         um no of followers + following')
         Max of no of followers + following is 1579
         1 persons having maximum no of followers + following
In [27]: print('No of persons having followers + following less than 10 are', np.
         sum(in out degree<10))</pre>
         No of persons having followers + following less than 10 are 1320326
In [28]: print('No of weakly connected components', len(list(nx.weakly connected
         components(q))))
         count=0
```

```
for i in list(nx.weakly_connected_components(g)):
    if len(i)==2:
        count+=1
print('weakly connected components wit 2 nodes',count)
```

No of weakly connected components 45558 weakly connected components wit 2 nodes 32195

2. Posing a problem as classification problem

2.1 Generating some edges which are not present in graph for supervised learning

Generated Bad links from graph which are not in graph and whose shortest path is greater than 2.

```
In [29]: %time
         ###generating bad edges from given graph
         import random
         if not os.path.isfile('data/after eda/missing edges final.p'):
             #getting all set of edges
             r = csv.reader(open('data/after eda/train woheader.csv','r'))
             edges = dict()
             for edge in r:
                 edges[(edge[0], edge[1])] = 1
             missing edges = set([])
             while (len(missing edges)<9437519):</pre>
                 a=random.randint(1, 1862220)
                 b=random.randint(1, 1862220)
                 tmp = edges.get((a,b),-1)
                 if tmp == -1 and a!=b:
                      try:
```

```
if nx.shortest path length(g,source=a,target=b) > 2:
                             missing edges.add((a,b))
                          else:
                              continue
                     except:
                             missing edges.add((a,b))
                 else:
                     continue
             pickle.dump(missing edges,open('data/after eda/missing edges final.
         p','wb'))
         else:
             missing edges = pickle.load(open('data/after eda/missing edges fina
         l.p','rb'))
         CPU times: user 1.96 s, sys: 1.4 s, total: 3.36 s
         Wall time: 3.83 s
In [30]: len(missing edges)
Out[30]: 9437519
```

2.2 Training and Test data split:

Removed edges from Graph and used as test data and after removing used that graph for creating features for Train and test data

```
In [31]: if (not os.path.isfile('data/after_eda/train_pos_after_eda.csv')) and (
    not os.path.isfile('data/after_eda/test_pos_after_eda.csv')):
        #reading total data df
        df_pos = pd.read_csv('data/train.csv')
        df_neg = pd.DataFrame(list(missing_edges), columns=['source_node',
        'destination_node'])

        print("Number of nodes in the graph with edges", df_pos.shape[0])
        print("Number of nodes in the graph without edges", df_neg.shape[0])
])
```

```
#Trian test split
             #Spiltted data into 80-20
             #positive links and negative links seperatly because we need positi
         ve training data only for creating graph
             #and for feature generation
             X_train_pos, X_test_pos, y_train_pos, y_test_pos = train_test_spli
         t(df pos,np.ones(len(df pos)),test size=0.2, random state=9)
             X train neg, X test neg, y train neg, y test neg = train test spli
         t(df neg,np.zeros(len(df neg)),test size=0.2, random state=9)
             print('='*60)
             print("Number of nodes in the train data graph with edges", X train
          _pos.shape[0],"=",y_train pos.shape[0])
             print("Number of nodes in the train data graph without edges", X tr
         ain neg.shape[0], "=", y train neg.shape[0])
             print('='*60)
             print("Number of nodes in the test data graph with edges", X test p
         os.shape[0], "=", y test pos.shape[0])
             print("Number of nodes in the test data graph without edges", X tes
         t neg.shape[0], "=", y test neg.shape[0])
             #removing header and saving
             X train pos.to csv('data/after eda/train pos after eda.csv',header=
         False, index=False)
             X test pos.to csv('data/after eda/test pos after eda.csv',header=Fa
         lse, index=False)
             X train neg.to csv('data/after eda/train neg after eda.csv',header=
         False. index=False)
             X test neg.to csv('data/after eda/test neg after eda.csv',header=Fa
         lse, index=False)
         else:
             #Graph from Traing data only
             del missing edges
In [32]: if (os.path.isfile('data/after eda/train pos after eda.csv')) and (os.p
         ath.isfile('data/after eda/test pos after eda.csv')):
             train graph=nx.read edgelist('data/after eda/train pos after eda.cs
         v',delimiter=',',create using=nx.DiGraph(),nodetype=int)
```

```
test graph=nx.read edgelist('data/after eda/test pos after eda.csv'
,delimiter=',',create using=nx.DiGraph(),nodetype=int)
    print(nx.info(train graph))
    print(nx.info(test graph))
    # finding the unique nodes in the both train and test graphs
    train nodes pos = set(train graph.nodes())
    test nodes pos = set(test graph.nodes())
    trY teY = len(train nodes pos.intersection(test nodes pos))
    trY teN = len(train nodes pos - test nodes pos)
    teY trN = len(test nodes pos - train nodes pos)
    print('no of people common in train and test -- ',trY teY)
    print('no of people present in train but not present in test -- ',t
rY teN)
    print('no of people present in test but not present in train -- ',t
eY trN)
    print(' % of people not there in Train but exist in Test in total T
est data are {} %'.format(teY trN/len(test nodes pos)*100))
Name:
Type: DiGraph
Number of nodes: 1780722
Number of edges: 7550015
Average in degree:
                    4.2399
Average out degree: 4.2399
Name:
Type: DiGraph
Number of nodes: 1144623
Number of edges: 1887504
Average in degree:
                     1.6490
Average out degree: 1.6490
no of people common in train and test -- 1063125
no of people present in train but not present in test -- 717597
no of people present in test but not present in train -- 81498
 % of people not there in Train but exist in Test in total Test data ar
e 7.1200735962845405 %
```

```
In [33]: #final train and test data sets
         if (not os.path.isfile('data/after eda/train after eda.csv')) and \
         (not os.path.isfile('data/after eda/test after eda.csv')) and \
         (not os.path.isfile('data/train y.csv')) and \
         (not os.path.isfile('data/test y.csv')) and \
         (os.path.isfile('data/after eda/train pos after eda.csv')) and \
         (os.path.isfile('data/after eda/test pos after eda.csv')) and \
         (os.path.isfile('data/after eda/train neg after eda.csv')) and \
         (os.path.isfile('data/after eda/test neg after eda.csv')):
             X train pos = pd.read csv('data/after eda/train pos after eda.csv',
          names=['source node', 'destination node'])
             X test pos = pd.read csv('data/after eda/test pos after eda.csv', n
         ames=['source node', 'destination node'])
             X train neg = pd.read csv('data/after eda/train neg after eda.csv',
          names=['source node', 'destination node'])
             X test neg = pd.read csv('data/after eda/test neg after eda.csv', n
         ames=['source node', 'destination node'])
             print('='*60)
             print("Number of nodes in the train data graph with edges", X train
          pos.shape[0])
             print("Number of nodes in the train data graph without edges", X tr
         ain neg.shape[0])
             print('='*60)
             print("Number of nodes in the test data graph with edges", X test p
         os.shape[0])
             print("Number of nodes in the test data graph without edges", X tes
         t neg.shape[0])
             X train = X train pos.append(X train neg,ignore index=True)
             y train = np.concatenate((y train pos,y train neg))
             X test = X test pos.append(X test neg,ignore index=True)
             y test = np.concatenate((y test pos,y test neg))
```

```
X_train.to_csv('data/after_eda/train_after_eda.csv',header=False,in
dex=False)
    X_test.to_csv('data/after_eda/test_after_eda.csv',header=False,inde
x=False)
    pd.DataFrame(y_train.astype(int)).to_csv('data/train_y.csv',header=
False,index=False)
    pd.DataFrame(y_test.astype(int)).to_csv('data/test_y.csv',header=False,index=False)
```

```
In [ ]: print("Data points in train data",X_train.shape)
    print("Data points in test data",X_test.shape)
    print("Shape of traget variable in train",y_train.shape)
    print("Shape of traget variable in test", y_test.shape)
```

1. Reading Data

Name:

Type: DiGraph

Number of nodes: 1780722 Number of edges: 7550015 Average in degree: 4.2399 Average out degree: 4.2399

2. Similarity measures

2.1 Jaccard Distance:

http://www.statisticshowto.com/jaccard-index/

$$j = \frac{|X \cap Y|}{|X \cup Y|}$$

```
In [3]: #for followees
        def jaccard for followees(a,b):
            try:
                if len(set(train graph.successors(a))) == 0 | len(set(train gr
        aph.successors(b))) == 0:
                    return 0
                sim = (len(set(train graph.successors(a)).intersection(set(trai))
        n graph.successors(b))))/\
                                             (len(set(train graph.successors(a))
         .union(set(train_graph.successors(b)))))
            except:
                return 0
            return sim
In [4]: #one test case
        print(jaccard for followees(273084,1505602))
        0.0
In [5]: #node 1635354 not in graph
        print(jaccard for followees(273084,1505602))
        0.0
In [6]: #for followers
        def jaccard for followers(a,b):
            try:
                if len(set(train graph.predecessors(a))) == 0 | len(set(g.pred
        ecessors(b)) == 0:
```

0

0

2.2 Cosine distance

$$CosineDistance = rac{|X \cap Y|}{|X| \cdot |Y|}$$

```
In [10]: print(cosine for followees(273084,1505602))
         0.0
In [11]: print(cosine for followees(273084,1635354))
         0
In [12]: def cosine for followers(a,b):
             try:
                 if len(set(train graph.predecessors(a))) == 0 | len(set(train
         graph.predecessors(b))) == 0:
                     return 0
                 sim = (len(set(train graph.predecessors(a)).intersection(set(tr
         ain graph.predecessors(b))))/\
                                               (math.sqrt(len(set(train_graph.pre
         decessors(a))))*(len(set(train graph.predecessors(b)))))
                 return sim
             except:
                 return 0
In [13]: print(cosine for followers(2,470294))
         0.02886751345948129
In [14]: print(cosine for followers(669354,1635354))
         0
```

2.3 Preferential Attachment

One well-known concept in social networks is that users with many friends tend to create more connections in the future. This is due to the fact that in some social networks, like in finance, the

rich get richer. We estimate how "rich" our two vertices are by calculating the multiplication between the number of friends (|N(x)|) or followers each vertex has. It may be noted that the similarity index does not require any node neighbor information; therefore, this similarity index has the lowest computational complexity.

$$PreferentialAttachment = |N(X)| \cdot |N(Y)|$$

For two nodes a,b, the preferential attachment= number of followers(a)* number of followers(b).

```
In [15]: def preferentialAttachment(a,b):
    try:
        if len(set(train_graph.predecessors(a))) == 0 | len(set(train_graph.predecessors(b))) == 0:
            return 0
            sim = (len(set(train_graph.successors(a)))*len((set(train_graph.successors(b)))))
            return sim
    except:
        return 0
```

```
In [16]: print(preferentialAttachment(273084,1505602))
```

120

```
In [17]: print(preferentialAttachment(273084,1635354))
```

0

3. Ranking Measures

https://networkx.github.io/documentation/networkx-

1.10/reference/generated/networkx.algorithms.link_analysis.pagerank_alg.pagerank.html

PageRank computes a ranking of the nodes in the graph G based on the structure of the incoming links.

Mathematical PageRanks for a simple network, expressed as percentages. (Google uses a logarithmic scale.) Page C has a higher PageRank than Page E, even though there are fewer links to C; the one link to C comes from an important page and hence is of high value. If web surfers who start on a random page have an 85% likelihood of choosing a random link from the page they are currently visiting, and a 15% likelihood of jumping to a page chosen at random from the entire web, they will reach Page E 8.1% of the time. (The 15% likelihood of jumping to an arbitrary page corresponds to a damping factor of 85%.) Without damping, all web surfers would eventually end up on Pages A, B, or C, and all other pages would have PageRank zero. In the presence of damping, Page A effectively links to all pages in the web, even though it has no outgoing links of its own.

3.1 Page Ranking

https://en.wikipedia.org/wiki/PageRank

```
In [18]: if not os.path.isfile('data/fea_sample/page_rank.p'):
    pr = nx.pagerank(train_graph, alpha=0.85)
    pickle.dump(pr,open('data/fea_sample/page_rank.p','wb'))
else:
    pr = pickle.load(open('data/fea_sample/page_rank.p','rb'))

In [19]: print('min',pr[min(pr, key=pr.get)])
    print('max',pr[max(pr, key=pr.get)])
    print('mean',float(sum(pr.values())) / len(pr))

min 1.6556497245737814e-07
    max 2.7098251341935817e-05
    mean 5.615699699365892e-07
In [20]: #for imputing to nodes which are not there in Train data
    mean_pr = float(sum(pr.values())) / len(pr)
    print(mean_pr)
```

4. Other Graph Features

4.1 Shortest path:

Getting Shortest path between twoo nodes, if nodes have direct path i.e directly connected then we are removing that edge and calculating path.

```
In [21]: #if has direct edge then deleting that edge and calculating shortest pa
         def compute shortest path length(a,b):
             p = -1
             try:
                 if train graph.has edge(a,b):
                     train graph.remove edge(a,b)
                     p= nx.shortest path length(train graph,source=a,target=b)
                     train graph.add edge(a,b)
                 else:
                      p= nx.shortest path length(train graph,source=a,target=b)
                 return p
             except:
                  return -1
In [22]: #testing
         compute shortest path length(77697, 826021)
Out[22]: 10
In [23]: #testing
         compute shortest path length(669354,1635354)
Out[23]: -1
```

4.2 Checking for same community

```
In [24]: #getting weekly connected edges from graph
         wcc=list(nx.weakly connected components(train graph))
         def belongs_to_same_wcc(a,b):
             index = []
             if train graph.has edge(b,a):
                  return 1
             if train graph.has edge(a,b):
                     for i in wcc:
                         if a in i:
                             index= i
                             break
                     if (b in index):
                         train graph.remove edge(a,b)
                         if compute_shortest_path_length(a,b)==-1:
                             train graph.add edge(a,b)
                              return 0
                          else:
                             train graph.add edge(a,b)
                              return 1
                     else:
                          return 0
             else:
                     for i in wcc:
                         if a in i:
                             index= i
                             break
                     if(b in index):
                          return 1
                     else:
                          return 0
In [25]: belongs_to_same_wcc(861, 1659750)
Out[25]: 0
```

```
In [26]: belongs_to_same_wcc(669354,1635354)
Out[26]: 0
```

4.3 Adamic/Adar Index:

Adamic/Adar measures is defined as inverted sum of degrees of common neighbours for given two vertices.

$$A(x,y) = \sum_{u \in N(x) \cap N(y)} rac{1}{log(|N(u)|)}$$

```
In [28]: calc_adar_in(1,189226)
Out[28]: 0
In [29]: calc_adar_in(669354,1635354)
Out[29]: 0
```

4.4 Is persion was following back:

```
In [30]: def follows_back(a,b):
    if train_graph.has_edge(b,a):
        return 1
    else:
        return 0

In [31]: follows_back(1,189226)

Out[31]: 1

In [32]: follows_back(669354,1635354)
Out[32]: 0
```

4.5 Katz Centrality:

https://en.wikipedia.org/wiki/Katz centrality

https://www.geeksforgeeks.org/katz-centrality-centrality-measure/ Katz centrality computes the centrality for a node based on the centrality of its neighbors. It is a generalization of the eigenvector centrality. The Katz centrality for node i is

$$x_i = lpha \sum_j A_{ij} x_j + eta,$$

where A is the adjacency matrix of the graph G with eigenvalues

 λ

The parameter

β

controls the initial centrality and

$$\alpha < rac{1}{\lambda_{max}}$$

4.6 Hits Score

The HITS algorithm computes two numbers for a node. Authorities estimates the node value based on the incoming links. Hubs estimates the node value based on outgoing links.

https://en.wikipedia.org/wiki/HITS algorithm

```
In [36]:
    if not os.path.isfile('data/fea_sample/hits.p'):
        hits = nx.hits(train_graph, max_iter=100, tol=1e-08, nstart=None, n
        ormalized=True)
        pickle.dump(hits,open('data/fea_sample/hits.p','wb'))
    else:
        hits = pickle.load(open('data/fea_sample/hits.p','rb'))
```

5. Featurization

5. 1 Reading a sample of Data from both train and test

```
In [38]: import random
         if os.path.isfile('data/after eda/train after eda.csv'):
             filename = "data/after eda/train after eda.csv"
             # you uncomment this line, if you dont know the lentah of the file
          name
             # here we have hardcoded the number of lines as 15100030
             # n train = sum(1 for line in open(filename)) #number of records in
          file (excludes header)
             n train = 15100028
             s = 100000 #desired sample size
             skip train = sorted(random.sample(range(1,n train+1),n train-s))
             #https://stackoverflow.com/a/22259008/4084039
In [39]: if os.path.isfile('data/after eda/train after eda.csv'):
             filename = "data/after eda/test after eda.csv"
             # you uncomment this line, if you don't know the lentgh of the file
          name
             # here we have hardcoded the number of lines as 3775008
             # n test = sum(1 for line in open(filename)) #number of records in
          file (excludes header)
             n test = 3775006
             s = 50000 #desired sample size
```

```
skip test = sorted(random.sample(range(1,n test+1),n test-s))
             #https://stackoverflow.com/a/22259008/4084039
In [40]:
         print("Number of rows in the train data file:", n train)
         print("Number of rows we are going to elimiate in train data are", len(s
         kip train))
         print("Number of rows in the test data file:", n test)
         print("Number of rows we are going to elimiate in test data are".len(sk
         ip test))
         Number of rows in the train data file: 15100028
         Number of rows we are going to elimiate in train data are 15000028
         Number of rows in the test data file: 3775006
         Number of rows we are going to elimiate in test data are 3725006
In [41]:
         df final train = pd.read csv('data/after eda/train after eda.csv', skip
         rows=skip train, names=['source node', 'destination node'])
         df final train['indicator link'] = pd.read csv('data/train y.csv', skip
         rows=skip train, names=['indicator link'])
         print("Our train matrix size ", df final train.shape)
         df final train.head(2)
         Our train matrix size (100002, 3)
Out[41]:
            source node destination node indicator link
                273084
          0
                             1505602
          1
               1757093
                              912379
In [42]: df final test = pd.read csv('data/after eda/test after eda.csv', skipro
         ws=skip test, names=['source node', 'destination node'])
         df final test['indicator link'] = pd.read csv('data/test y.csv', skipro
         ws=skip test, names=['indicator link'])
         print("Our test matrix size ",df final test.shape)
         df final test.head(2)
         Our test matrix size (50002, 3)
```

Out[42]: source_node destination_node indicator_link 0 848424 784690 1 1 1227220 827479 1

5.2 Adding a set of features

we will create these each of these features for both train and test data points

```
1. jaccard_followers
```

- 2. jaccard_followees
- 3. cosine_followers
- 4. cosine_followees
- 5. preferentialAttachment
- 6. num followers s
- 7. num_followees_s
- 8. num_followers_d
- 9. num followees d
- 10. inter_followers
- 11. inter followees

```
OW:
                                                      jaccard for followees(row[
         'source node'],row['destination node']),axis=1)
             df final test['jaccard followees'] = df final test.apply(lambda row
                                                      jaccard for followees(row[
          'source node'], row['destination node']), axis=1)
             #mapping cosine followers to train and test data
             df final train['cosine followers'] = df final train.apply(lambda ro
         W:
                                                      cosine for followers(row['s
         ource node'],row['destination node']),axis=1)
             df final test['cosine followers'] = df final test.apply(lambda row:
                                                      cosine for followers(row['s
         ource node'],row['destination node']),axis=1)
             #mapping cosine followees to train and test data
             df final train['cosine followees'] = df final train.apply(lambda ro
         w:
                                                      cosine for followees(row['s
         ource node'],row['destination node']),axis=1)
             df final test['cosine followees'] = df final test.apply(lambda row:
                                                      cosine for followees(row['s
         ource node'],row['destination_node']),axis=1)
             #mapping preferential Attachment to train and test data
             df final train['pref attachment'] = df final train.apply(lambda row
                                                      preferentialAttachment(row[
          'source node'],row['destination node']),axis=1)
             df final test['pref attachment'] = df_final_test.apply(lambda row:
                                                      preferentialAttachment(row[
          'source node'],row['destination node']),axis=1)
In [44]: def compute features stage1(df final):
             #calculating no of followers followees for source and destination
             #calculating intersection of followers and followees for source and
```

```
destination
             num followers s=[]
             num followees s=[]
             num followers d=[]
             num followees d=[]
             inter followers=[]
             inter followees=[]
             for i,row in df final.iterrows():
                 try:
                     s1=set(train graph.predecessors(row['source node']))
                     s2=set(train graph.successors(row['source node']))
                 except:
                      s1 = set()
                     s2 = set()
                 try:
                     dl=set(train graph.predecessors(row['destination node']))
                     d2=set(train graph.successors(row['destination node']))
                 except:
                     d1 = set()
                     d2 = set()
                 num followers s.append(len(s1))
                 num followees s.append(len(s2))
                 num followers d.append(len(d1))
                 num followees d.append(len(d2))
                 inter followers.append(len(s1.intersection(d1)))
                 inter followees.append(len(s2.intersection(d2)))
             return num followers s, num followers d, num followees s, num follo
         wees d, inter followers, inter followees
In [45]: if not os.path.isfile('data/fea sample/storage sample stage1.h5'):
             df final train['num followers s'], df final train['num followers d'
         ], \
             df final train['num followees s'], df final train['num followees d'
             df final train['inter followers'], df final train['inter followees'
         ]= compute features stage1(df final train)
```

```
df_final_test['num_followers_s'], df_final_test['num_followers_d'],
    df_final_test['num_followees_s'], df_final_test['num_followees_d'],
    df_final_test['inter_followers'], df_final_test['inter_followees']=
    compute_features_stagel(df_final_test)

    hdf = HDFStore('data/fea_sample/storage_sample_stagel.h5')
    hdf.put('train_df',df_final_train, format='table', data_columns=Tru
e)
    hdf.put('test_df',df_final_test, format='table', data_columns=True)
    hdf.close()
else:
    df_final_train = read_hdf('data/fea_sample/storage_sample_stagel.h
5', 'train_df',mode='r')
    df_final_test = read_hdf('data/fea_sample/storage_sample_stagel.h5',
    'test_df',mode='r')
```

5.3 Adding new set of features

we will create these each of these features for both train and test data points

- 1. adar index
- 2. is following back
- 3. belongs to same weakly connect components
- 4. shortest path between source and destination

```
In [46]: if not os.path.isfile('data/fea_sample/storage_sample_stage2.h5'):
    #mapping adar index on train
    df_final_train['adar_index'] = df_final_train.apply(lambda row: cal
    c_adar_in(row['source_node'],row['destination_node']),axis=1)
    #mapping adar index on test
    df_final_test['adar_index'] = df_final_test.apply(lambda row: calc_
    adar_in(row['source_node'],row['destination_node']),axis=1)
```

```
#mapping followback or not on train
    df final train['follows_back'] = df_final_train.apply(lambda row: f
ollows back(row['source node'], row['destination node']), axis=1)
    #mapping followback or not on test
   df final test['follows back'] = df final test.apply(lambda row: fol
lows back(row['source node'], row['destination node']), axis=1)
   #mapping same component of wcc or not on train
    df final train['same comp'] = df final train.apply(lambda row: belo
ngs to same wcc(row['source node'],row['destination node']),axis=1)
   ##mapping same component of wcc or not on train
    df final test['same comp'] = df final test.apply(lambda row: belong
s_to_same_wcc(row['source node'],row['destination node']),axis=1)
    #----
   #mapping shortest path on train
    df final train['shortest path'] = df final train.apply(lambda row:
compute shortest path length(row['source node'],row['destination node'
1).axis=1)
   #mapping shortest path on test
    df final test['shortest path'] = df final test.apply(lambda row: co
mpute shortest path length(row['source node'],row['destination node']),
axis=1)
   hdf = HDFStore('data/fea sample/storage sample stage2.h5')
   hdf.put('train df', df final train, format='table', data columns=Tru
e)
   hdf.put('test df', df final test, format='table', data columns=True)
   hdf.close()
else:
    df final train = read hdf('data/fea sample/storage sample stage2.h
5', 'train df', mode='r')
```

```
df_final_test = read_hdf('data/fea_sample/storage_sample_stage2.h5'
, 'test_df',mode='r')
```

5.4 Adding new set of features

we will create these each of these features for both train and test data points

- 1. Weight Features
 - weight of incoming edges
 - weight of outgoing edges
 - weight of incoming edges + weight of outgoing edges
 - weight of incoming edges * weight of outgoing edges
 - 2*weight of incoming edges + weight of outgoing edges
 - weight of incoming edges + 2*weight of outgoing edges
- 2. Page Ranking of source
- 3. Page Ranking of dest
- 4. katz of source
- 5. katz of dest
- 6. hubs of source
- 7. hubs of dest
- 8. authorities_s of source
- 9. authorities_s of dest

Weight Features

In order to determine the similarity of nodes, an edge weight value was calculated between nodes. Edge weight decreases as the neighbor count goes up. Intuitively, consider one million people following a celebrity on a social network then chances are most of them never met each other or the celebrity. On the other hand, if a user has 30 contacts in his/her social network, the chances are higher that many of them know each other. credit - Graph-based Features for Supervised Link Prediction William Cukierski, Benjamin Hamner, Bo Yang

$$W=rac{1}{\sqrt{1+|X|}}$$

it is directed graph so calculated Weighted in and Weighted out differently

```
In [64]: #weight for source and destination of each link
         Weight in = {}
         Weight out = {}
         for i in tqdm(train graph.nodes()):
             s1=set(train graph.predecessors(i))
             w in = 1.0/(np.sqrt(1+len(s1)))
             Weight in[i]=w in
             s2=set(train graph.successors(i))
             w_out = 1.0/(np.sqrt(1+len(s2)))
             Weight out[i]=w out
         #for imputing with mean
         mean weight in = np.mean(list(Weight in.values()))
         mean weight out = np.mean(list(Weight out.values()))
                        | 1780722/1780722 [00:16<00:00, 105780.58it/s]
In [65]: if not os.path.isfile('data/fea sample/storage sample stage3.h5'):
             #mapping to pandas train
             df final train['weight in'] = df final train.destination node.apply
         (lambda x: Weight in.get(x,mean weight in))
             df final train['weight out'] = df_final_train.source_node.apply(lam
         bda x: Weight out.get(x,mean weight out))
             #mapping to pandas test
             df final test['weight in'] = df final test.destination node.apply(l
         ambda x: Weight in.get(x,mean weight in))
             df final test['weight out'] = df final test.source node.apply(lambd
         a x: Weight out.get(x,mean weight out))
```

```
#some features engineerings on the in and out weights
             df final train['weight f1'] = df final train.weight in + df final t
         rain.weight out
             df_final_train['weight_f2'] = df final train.weight in * df final t
         rain.weight out
             df final train['weight f3'] = (2*df final train.weight in + 1*df fi
         nal train.weight out)
             df final train['weight f4'] = (1*df final train.weight in + 2*df fi
         nal train.weight out)
             #some features engineerings on the in and out weights
             df final test['weight f1'] = df final test.weight in + df final tes
         t.weiaht out
             df final test['weight f2'] = df final test.weight in * df final tes
         t.weight out
             df final test['weight f3'] = (2*df final test.weight in + 1*df fina
         l test.weight out)
             df final test['weight f4'] = (1*df final test.weight in + 2*df fina
         l test.weight out)
In [66]: if not os.path.isfile('data/fea sample/storage sample stage3.h5'):
             #page rank for source and destination in Train and Test
             #if anything not there in train graph then adding mean page rank
             df final train['page rank s'] = df final train.source node.applv(la
         mbda x:pr.get(x,mean pr))
             df final train['page rank d'] = df final train.destination node.app
         ly(lambda x:pr.get(x,mean pr))
             df final test['page rank s'] = df final test.source node.apply(lamb
         da x:pr.get(x,mean pr))
             df final test['page rank d'] = df final test.destination node.apply
         (lambda x:pr.get(x,mean pr))
             #Katz centrality score for source and destination in Train and test
             #if anything not there in train graph then adding mean katz score
             df final train['katz s'] = df final train.source node.apply(lambda
```

```
x: katz.get(x,mean katz))
    df_final_train['katz_d'] = df final train.destination node.apply(la
mbda x: katz.get(x,mean katz))
    df final test['katz s'] = df final test.source node.apply(lambda x:
katz.get(x,mean katz))
    df final test['katz d'] = df final test.destination node.apply(lamb
da x: katz.get(x,mean katz))
    #Hits algorithm score for source and destination in Train and test
    #if anything not there in train graph then adding 0
    df final train['hubs s'] = df final train.source node.apply(lambda
x: hits[0].get(x,0)
    df final train['hubs d'] = df final train.destination node.apply(la
mbda x: hits[0].get(x,0))
    df final test['hubs s'] = df final test.source node.apply(lambda x:
hits[0].get(x,0)
    df final test['hubs d'] = df final test.destination node.apply(lamb
da x: hits[0].get(x,0))
    #Hits algorithm score for source and destination in Train and Test
    #if anything not there in train graph then adding 0
    df final train['authorities s'] = df final train.source node.apply(
lambda x: hits[1].qet(x,0))
    df final train['authorities d'] = df final train.destination node.a
pply(lambda x: hits[1].get(x,0))
    df final test['authorities s'] = df final test.source node.apply(la
mbda x: hits[1].qet(x,0))
    df final test['authorities_d'] = df_final_test.destination_node.app
ly(lambda x: hits[1].get(x,0))
```

```
hdf = HDFStore('data/fea_sample/storage_sample_stage3.h5')
hdf.put('train_df',df_final_train, format='table', data_columns=Tru
e)
hdf.put('test_df',df_final_test, format='table', data_columns=True)
hdf.close()
else:
    df_final_train = read_hdf('data/fea_sample/storage_sample_stage3.h
5', 'train_df',mode='r')
    df_final_test = read_hdf('data/fea_sample/storage_sample_stage3.h5'
, 'test_df',mode='r')
```

5.5 Adding new set of features

we will create these each of these features for both train and test data points

1. SVD features for both source and destination

```
In [67]: def svd(x, S):
    try:
    z = sadj_dict[x]
    return S[z]
    except:
        return [0,0,0,0,0,0]
In [68]: #for svd features to get feature vector creating a dict node val and in edx in svd vector
    sadj_col = sorted(train_graph.nodes())
    sadj_dict = { val:idx for idx,val in enumerate(sadj_col)}

In [69]: Adj = nx.adjacency_matrix(train_graph,nodelist=sorted(train_graph.nodes ())).asfptype()

In [70]: U, s, V = svds(Adj, k = 6)
    print('Adjacency matrix Shape',Adj.shape)
    print('U Shape',U.shape)
```

```
print('V Shape', V.shape)
         print('s Shape',s.shape)
         Adjacency matrix Shape (1780722, 1780722)
         U Shape (1780722, 6)
         V Shape (6, 1780722)
         s Shape (6,)
In [71]: if not os.path.isfile('data/fea sample/storage sample stage4.h5'):
             df final train[['svd u s 1', 'svd u s 2', 'svd u s 3', 'svd u s 4',
         'svd u s 5', 'svd u s 6']] = \
             df final train.source node.apply(lambda x: svd(x, U)).apply(pd.Seri
         es)
             df final train[['svd u d 1', 'svd u d 2', 'svd u d 3', 'svd u d 4',
          'svd u d 5','svd u d 6']] = \
             df final train.destination node.apply(lambda x: svd(x, U)).apply(pd
          .Series)
             df final train[['svd v s 1','svd v s 2', 'svd v s 3', 'svd v s 4',
         'svd v s 5', 'svd v s 6',]] = \
             df final train.source node.apply(lambda x: svd(x, V.T)).apply(pd.Se
         ries)
             df final train[['svd v d 1', 'svd v d 2', 'svd v d 3', 'svd v d 4',
          'svd v d 5', 'svd v d 6']] = \
             df final train.destination node.apply(lambda x: svd(x, V.T)).apply(
         pd.Series)
             df final test[['svd u s 1', 'svd u s 2','svd u s 3', 'svd u s 4',
         'svd u s 5', 'svd u s 6']] = \
             df final test.source node.apply(lambda x: svd(x, U)).apply(pd.Serie
```

```
s)
    df final test[['svd u d 1', 'svd u d 2', 'svd u d 3', 'svd u d 4',
'svd u d 5', 'svd u d 6'11 = \
    df final test.destination node.apply(lambda x: svd(x, U)).apply(pd.
Series)
    df final test[['svd v s 1','svd v s 2', 'svd v s 3', 'svd v s 4',
'svd v s 5', 'svd v s 6',]] = \
    df final test.source node.apply(lambda x: svd(x, V.T)).apply(pd.Ser
ies)
    df final test[['svd v d 1', 'svd v d 2', 'svd v d 3', 'svd v d 4',
'svd v d 5', 'svd v d 6'11 = \
    df final test.destination node.apply(lambda x: svd(x, V.T)).apply(p
d.Series)
    ###### SVD dot is the product between source node svd and destinati
on node syd
    #Train Dataset
    s1,s2,s3,s4,s5,s6=df final train['svd u s 1'],df final train['svd u
s 2'], df final train['svd u s 3'], df final train['svd u s 4'], df final
train['svd u s 5'], df final train['svd u s 6']
    s7,s8,s9,s10,s11,s12=df final train['svd v s 1'],df final train['sv
d v s 2'], df final train['svd v s 3'], df final train['svd v s 4'], df fi
nal train['svd v s 5'], df final train['svd v s 6']
    d1,d2,d3,d4,d5,d6=df final train['svd u d 1'],df final train['svd u
d 2'], df final train['svd u d 3'], df final train['svd u d 4'], df final
train['svd u d 5'], df final train['svd u d 6']
    d7,d8,d9,d10,d11,d12=df final train['svd v d 1'],df final train['sv
d v d 2'], df final train['svd v d 3'], df final train['svd v d 4'], df fi
nal train['svd v d 5'],df final train['svd v d 6']
```

```
svd dot=[]
    for i in range(len(np.array(s1))):
        a=[]
        b=[]
        a.append(np.array(s1[i]))
        a.append(np.array(s2[i]))
        a.append(np.arrav(s3[i]))
        a.append(np.array(s4[i]))
        a.append(np.array(s5[i]))
        a.append(np.array(s6[i]))
        a.append(np.array(s7[i]))
        a.append(np.array(s8[i]))
        a.append(np.array(s9[i]))
        a.append(np.array(s10[i]))
        a.append(np.array(s11[i]))
        a.append(np.array(s12[i]))
        b.append(np.array(d1[i]))
        b.append(np.array(d2[i]))
        b.append(np.array(d3[i]))
        b.append(np.array(d4[i]))
        b.append(np.array(d5[i]))
        b.append(np.array(d6[i]))
        b.append(np.array(d7[i]))
        b.append(np.array(d8[i]))
        b.append(np.array(d9[i]))
        b.append(np.array(d10[i]))
        b.append(np.array(d11[i]))
        b.append(np.array(d12[i]))
        svd dot.append(np.dot(a,b))
    df final train['svd dot']=svd dot
    # Test dataset
    s1,s2,s3,s4,s5,s6=df final test['svd u s 1'],df final test['svd u s
2'], df final test['svd u s 3'], df final test['svd u s 4'], df final tes
t['svd u s 5'], df final test['svd u s 6']
    s7,s8,s9,s10,s11,s12=df final test['svd v s 1'],df final test['svd
v s 2'], df final test['svd v s 3'], df final test['svd v s 4'], df final
test['svd v s 5'], df final test['svd v s 6']
```

```
d1,d2,d3,d4,d5,d6=df final test['svd u d 1'],df final test['svd u d
2'], df final test['svd u d 3'], df final test['svd u d 4'], df final tes
t['svd u d 5'], df final test['svd u d 6']
    d7,d8,d9,d10,d11,d12=df final test['svd v d 1'],df final test['svd
v d 2'], df final test['svd v d 3'], df final test['svd v d 4'], df final
test['svd v d 5'], df final test['svd v d 6']
    svd dot=[]
    for i in range(len(np.array(s1))):
        a=[]
        b=[1]
        a.append(np.array(s1[i]))
        a.append(np.array(s2[i]))
        a.append(np.array(s3[i]))
        a.append(np.array(s4[i]))
        a.append(np.array(s5[i]))
        a.append(np.array(s6[i]))
        a.append(np.array(s7[i]))
        a.append(np.array(s8[i]))
        a.append(np.array(s9[i]))
        a.append(np.array(s10[i]))
        a.append(np.array(s11[i]))
        a.append(np.array(s12[i]))
        b.append(np.array(d1[i]))
        b.append(np.array(d2[i]))
        b.append(np.array(d3[i]))
        b.append(np.array(d4[i]))
        b.append(np.array(d5[i]))
        b.append(np.array(d6[i]))
        b.append(np.array(d7[i]))
        b.append(np.array(d8[i]))
        b.append(np.array(d9[i]))
        b.append(np.array(d10[i]))
        b.append(np.array(d11[i]))
        b.append(np.array(d12[i]))
        svd dot.append(np.dot(a,b))
    df final test['svd dot']=svd dot
```

```
hdf = HDFStore('data/fea_sample/storage_sample_stage4.h5')
               hdf.put('train df',df_final_train, format='table', data_columns=Tru
          e)
               hdf.put('test df', df final test, format='table', data columns=True)
               hdf.close()
In [72]: df final train.head(2)
Out[72]:
             source_node destination_node indicator_link jaccard_followers jaccard_followees cosine_foll-
           0
                  273084
                                1505602
                                                  1
                                                                               0.0
                                                                                          0.0
                                 912379
                                                                 0
                                                                               0.0
           1
                 1757093
                                                                                          0.0
          2 rows × 57 columns
In [73]: df final test.head(2)
Out[73]:
              source_node destination_node indicator_link jaccard_followers jaccard_followees cosine_followers
           0
                  848424
                                 784690
                                                                               0.0
                                                                                          0.0
                                                                               0.0
           1
                 1227220
                                 827479
                                                  1
                                                                 0
                                                                                          0.0
          2 rows × 57 columns
          Modelling
 In [2]: from sklearn.metrics import confusion matrix
          def plot confusion matrix(test y, predict y):
```

```
C = confusion matrix(test y, predict y)
            A = (((C.T)/(C.sum(axis=1))).T)
            B = (C/C.sum(axis=0))
            plt.figure(figsize=(20,4))
            labels = [0.1]
            # representing A in heatmap format
            cmap=sns.light palette("blue")
            plt.subplot(1, 3, 1)
            sns.heatmap(C, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels
         , yticklabels=labels)
            plt.xlabel('Predicted Class')
            plt.ylabel('Original Class')
            plt.title("Confusion matrix")
            plt.subplot(1, 3, 2)
            sns.heatmap(B, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels
         , vticklabels=labels)
            plt.xlabel('Predicted Class')
            plt.ylabel('Original Class')
            plt.title("Precision matrix")
            plt.subplot(1, 3, 3)
            # representing B in heatmap format
            sns.heatmap(A, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels
         , yticklabels=labels)
            plt.xlabel('Predicted Class')
            plt.ylabel('Original Class')
            plt.title("Recall matrix")
            plt.show()
In [3]: global result = pd.DataFrame(columns=['Model', 'Hyperparameters', 'Trai
        n-F1-Score', 'Test-F1-Score', 'Train-AUC', 'Test-AUC'])
In [4]: #reading
```

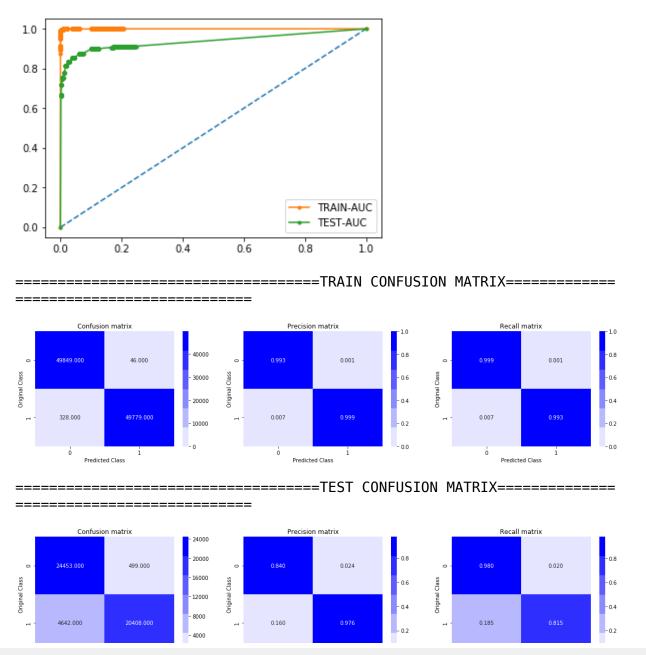
```
from pandas import read hdf
        df final train = read h\overline{d}f('data/fea sample/storage sample stage4.h5')
        'train df',mode='r')
        df final test = read hdf('data/fea sample/storage sample stage4.h5', 't
        est df', mode='r')
In [5]: df final train.columns
Out[5]: Index(['source node', 'destination node', 'indicator link',
                'jaccard followers', 'jaccard followees', 'cosine followers',
                'cosine followees', 'pref attachment', 'num followers s',
                'num followers d', 'num followees s', 'num followees d',
               'inter followers', 'inter followees', 'adar index', 'follows bac
        k',
                'same comp', 'shortest path', 'weight in', 'weight out', 'weight
        _f1',
                'weight f2', 'weight f3', 'weight_f4', 'page_rank_s', 'page_rank
        _d',
                'katz_s', 'katz_d', 'hubs_s', 'hubs_d', 'authorities s',
               'authorities d', 'svd u s 1', 'svd u s 2', 'svd u s 3', 'svd u s
        _4',
                'svd u s 5', 'svd u s 6', 'svd u d 1', 'svd u d 2', 'svd u d 3',
               'svd u d 4', 'svd u d 5', 'svd u d 6', 'svd v s 1', 'svd v s 2',
               'svd_v_s_3', 'svd_v_s_4', 'svd_v_s_5', 'svd_v_s_6', 'svd_v_d_1',
               'svd v d 2', 'svd v d 3', 'svd v d 4', 'svd_v_d_5', 'svd_v_d_6',
               'svd dot'],
              dtvpe='object')
In [6]: y train = df final train.indicator link
        y test = df final test.indicator link
In [7]: df final train.drop(['source node', 'destination node', 'indicator link'
        l,axis=1,inplace=True)
        df final test.drop(['source node', 'destination node', 'indicator link'
        l,axis=1,inplace=True)
In [8]: print("The shape of Train data :",df_final_train.shape)
        print("The shape of Train data :",df final test.shape)
```

```
The shape of Train data : (100002, 54)
The shape of Train data : (50002, 54)
```

RandomForest Classifier with Default Parameters

```
In [9]: clf = RandomForestClassifier(n jobs=-1, random state=42)
        clf.fit(df final train, y train)
        y pred train = clf.predict(df final train)
        y pred test = clf.predict(df final test)
        y pred train proba = clf.predict proba(df final train)[:,1]
        y pred test proba = clf.predict proba(df final test)[:,1]
        f1 score val train = f1 score(y train, y pred train)
        roc auc val train = roc auc score(y train, y pred train)
        f1 score val test = f1 score(y test, y pred test)
        roc auc val test = roc auc score(y test, y pred test)
        global result = global result.append({'Model': "RandomForest",
                               'Hyperparameters': "Default Parameters",
                               'Train-F1-Score': '{0:.4}'.format(f1 score val tr
        ain),
                               'Test-F1-Score': '{0:.4}'.format(f1 score val tes
        t),
                               'Train-AUC': '{0:.4}'.format(roc auc val train),
                               'Test-AUC': '{0:.4}'.format(roc auc val test)
                             }, ignore index=True)
        print("TRAIN Scores","*"*20)
        print("\tF1-Score: ", f1 score val train)
        print("\tROC-AUC : ", roc auc val train)
        print("TEST Scores","*"*20)
        print("\tF1-Score: ", f1_score_val_test)
        print("\tROC-AUC : ", roc auc val test)
```

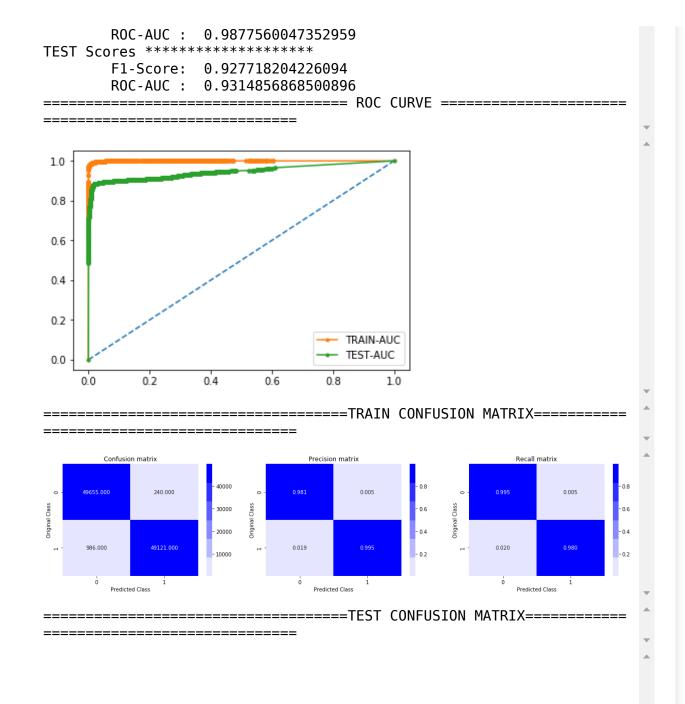
```
# calculate roc curve
fpr_train, tpr_train, thresholds = roc_curve(y_train, y_pred_train_prob
a)
plt.plot([0, 1], [0, 1], linestyle='--')
plt.plot(fpr_train, tpr_train, marker='.', label='TRAIN-AUC')
fpr test, tpr test, thresholds = roc curve(y test, y pred test proba)
plt.plot(fpr test, tpr test, marker='.', label='TEST-AUC')
# show the plot
plt.legend()
plt.show()
print("=======TRAIN CONFUSION MATRIX=====
plot confusion matrix(y train, y pred train)
plot confusion matrix(y test, y pred test)
TRAIN Scores *************
     F1-Score: 0.9962574550694472
     ROC-AUC: 0.9962660361781194
TEST Scores **************
     F1-Score: 0.8881345605674871
     ROC-AUC: 0.8973461109201925
```

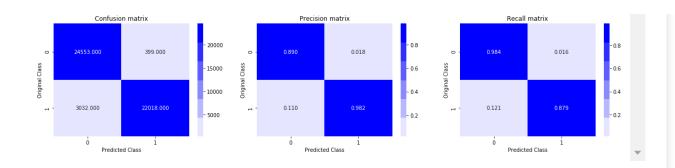


RandomForest Classifier with RandomSearchCV

```
In [10]: parameters = {
                          'n estimators': [10, 50, 100, 125, 150, 200, 300, 500],
                          'min samples leaf': [1, 2, 3, 5, 7, 9],
                          'min samples split': [2, 5, 6, 7, 8, 9]
         clf = RandomForestClassifier(n jobs=-1, random state=42, oob score=True
         r clf = RandomizedSearchCV(clf, param distributions=parameters, scoring
         ='f1', cv=5, random state=42, n jobs=-1, return train score=True)
         r clf.fit(df final train, y train)
         y pred train = r clf.predict(df final train)
         y pred test = r clf.predict(df final test)
         y pred train proba = r clf.predict proba(df final train)[:,1]
         y pred test proba = r clf.predict proba(df final test)[:,1]
         fl score val train = fl score(y train, y pred train)
         roc auc val train = roc auc score(y train, y pred train)
         f1 score val test = f1 score(y test, y pred test)
         roc auc val test = roc auc score(y test, y pred test)
         global result = global result.append({'Model': "RandomForest",
                                'Hyperparameters': r clf.best params ,
                                'Train-F1-Score': '{0:.4}'.format(f1 score val tr
         ain),
                                'Test-F1-Score': '{0:.4}'.format(f1 score val tes
         t),
                                'Train-AUC': '{0:.4}'.format(roc auc val train),
                                'Test-AUC': '{0:.4}'.format(roc auc val test)
                               }, ignore index=True)
```

```
print("*"*100)
print("Best Parameters: ", r clf.best params )
print("*"*100)
print("TRAIN Scores","*"*20)
print("\tF1-Score: ", f1_score_val_train)
print("\tROC-AUC : ", roc_auc_val_train)
print("TEST Scores","*"*20)
print("\tF1-Score: ", f1 score val test)
print("\tROC-AUC : ", roc auc val test)
# calculate roc curve
fpr train, tpr train, thresholds = roc curve(y train, y pred train prob
a)
plt.plot([0, 1], [0, 1], linestyle='--')
plt.plot(fpr train, tpr train, marker='.', label='TRAIN-AUC')
fpr test, tpr test, thresholds = roc curve(y test, y pred test proba)
plt.plot(fpr test, tpr test, marker='.', label='TEST-AUC')
# show the plot
plt.legend()
plt.show()
plot confusion matrix(y train, y pred train)
plot confusion matrix(y test, y pred test)
**************************
**********
Best Parameters: {'n estimators': 200, 'min samples split': 9, 'min
samples leaf': 1}
   *************************
***********
TRAIN Scores *************
     F1-Score: 0.9876744279567297
```

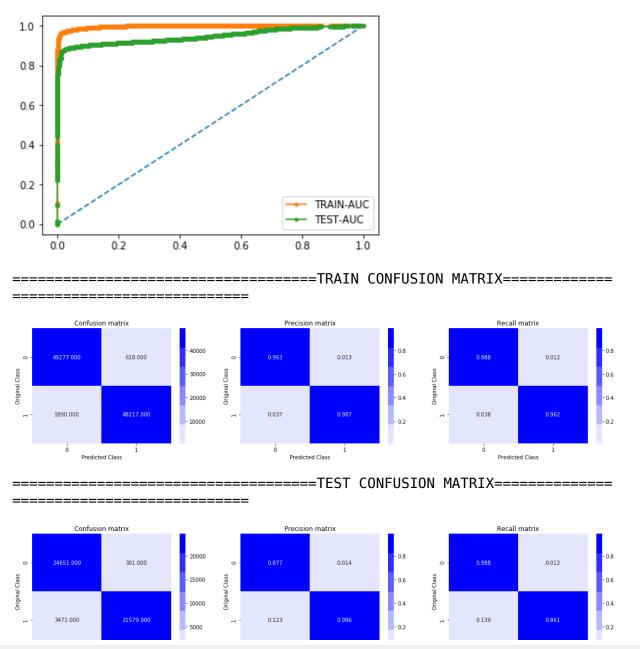




XGBoost Classifier with Default parameters

```
In [11]: clf = XGBClassifier(n jobs=-1, random state=42)
         clf.fit(df final train, y train)
         y pred train = clf.predict(df_final_train)
         y pred test = clf.predict(df final test)
         y pred train proba = clf.predict proba(df final train)[:,1]
         y pred test proba = clf.predict proba(df final test)[:,1]
         f1 score val train = f1 score(y train, y pred train)
         roc auc val train = roc auc score(y train, y pred train)
         fl score val test = fl score(y test, y pred test)
         roc auc val test = roc auc score(y test, y pred test)
         global result = global result.append({'Model': "XGBoost",
                               'Hyperparameters': "Default Parameters",
                               'Train-F1-Score': '{0:.4}'.format(f1 score val tr
         ain),
                               'Test-F1-Score': '{0:.4}'.format(f1 score val tes
         t),
                               'Train-AUC': '{0:.4}'.format(roc auc val train),
                               'Test-AUC': '{0:.4}'.format(roc auc val test)
```

```
}, ignore index=True)
print("TRAIN Scores","*"*20)
print("\tF1-Score: ", f1_score_val_train)
print("\tROC-AUC : ", roc_auc val train)
print("TEST Scores","*"*20)
print("\tF1-Score: ", f1 score val test)
print("\tROC-AUC : ", roc auc val test)
# calculate roc curve
fpr train, tpr train, thresholds = roc curve(y train, y pred train prob
a)
plt.plot([0, 1], [0, 1], linestyle='--')
plt.plot(fpr train, tpr train, marker='.', label='TRAIN-AUC')
fpr test, tpr test, thresholds = roc curve(y test, y pred test proba)
plt.plot(fpr_test, tpr test, marker='.', label='TEST-AUC')
# show the plot
plt.legend()
plt.show()
plot confusion matrix(y train, y pred train)
plot confusion matrix(y_test, y_pred_test)
TRAIN Scores *************
     F1-Score: 0.9746518162155606
     ROC-AUC: 0.9749473543192375
TEST Scores **************
     F1-Score: 0.9196249733645856
     ROC-AUC: 0.9246869822394327
```



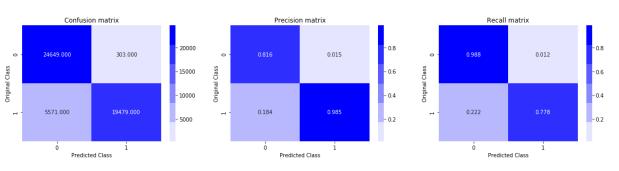
XGBoost Classifier with RandomSearchCV

```
In [12]: parameters = {
                          'n estimators': [10, 50, 100, 125, 150, 175, 300],
                          'learning rate': [0.1, 0.001, 0.5, 1, 0.005],
                          'max depth': [2, 3, 4, 5, 7, 9],
                          'gamma': [0, 0.01, 0.005, 0.5, 0.001],
                          'min child weight':range(1,6,2)
         clf = XGBClassifier(n jobs=-1, random state=42, oob score=True)
         r clf = RandomizedSearchCV(clf, param distributions=parameters, scoring
         ='f1', cv=5, random state=42, n jobs=-1, return train score=True)
         r clf.fit(df final train, v train)
         y pred train = r clf.predict(df final train)
         y pred test = r clf.predict(df final test)
         y pred train proba = r clf.predict proba(df final train)[:,1]
         y pred test proba = r clf.predict proba(df final test)[:,1]
         f1 score val train = f1 score(y train, y pred train)
         roc auc val train = roc auc score(y train, y pred train)
         f1 score val test = f1 score(y test, y pred test)
         roc auc val test = roc auc score(y test, y pred test)
         global result = global result.append({'Model': "XGBoost",
                                'Hyperparameters': r clf.best params ,
                                'Train-F1-Score': '{0:.4}'.format(f1 score val tr
         ain),
```

```
'Test-F1-Score': '{0:.4}'.format(f1 score val tes
t),
                 'Train-AUC': '{0:.4}'.format(roc auc val train),
                 'Test-AUC': '{0:.4}'.format(roc auc val test)
                }, ignore index=True)
print("*"*100)
print("Best Parameters: ", r clf.best params )
print("*"*100)
print("TRAIN Scores","*"*20)
print("\tF1-Score: ", f1 score val train)
print("\tROC-AUC : ", roc auc val train)
print("TEST Scores","*"*20)
print("\tF1-Score: ", f1 score val test)
print("\tROC-AUC : ", roc auc val test)
print("========= ROC CURVF =========
# calculate roc curve
fpr train, tpr train, thresholds = roc curve(y train, y pred train prob
a)
plt.plot([0, 1], [0, 1], linestyle='--')
plt.plot(fpr train, tpr train, marker='.', label='TRAIN-AUC')
fpr test, tpr test, thresholds = roc curve(y test, y pred test proba)
plt.plot(fpr test, tpr test, marker='.', label='TEST-AUC')
# show the plot
plt.legend()
plt.show()
_____")
plot confusion matrix(y train, y pred train)
plot confusion matrix(y test, y pred test)
**********
```

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```
Best Parameters: {'n_estimators': 300, 'learning_rate': 0.5, 'min_chil
d_weight': 1, 'gamma': 0.5, 'max_depth': 3}
TRAIN Scores *************
       F1-Score: 0.9967141730002895
       ROC-AUC: 0.9967118953474439
TEST Scores **************
       F1-Score: 0.8689775160599571
       ROC-AUC: 0.8827307376270224
           1.0
0.8
0.6
0.4
0.2
                                TRAIN-AUC
                              TEST-AUC
0.0
                 0.4
                       0.6
    0.0
          0.2
                              0.8
                                    1.0
                      ======TRAIN CONFUSION MATRIX=======
      Confusion matrix
                            Precision matrix
                                                   Recall matrix
                   40000
    49774.000
                   30000
                  20000
    208.000
                             Predicted Class
                                                   Predicted Class
```

Conclusion

In [13]: global_result

Out[13]:

	Model	Hyperparameters	Train- F1- Score	Test- F1- Score	Train- AUC	Test- AUC
0	RandomForest	Default Parameters	0.9963	0.8881	0.9963	0.8973
1	RandomForest	{'n_estimators': 200, 'min_samples_split': 9, 'min_samples_leaf': 1}	0.9877	0.9277	0.9878	0.9315
2	XGBoost	Default Parameters	0.9747	0.9196	0.9749	0.9247
3	XGBoost	{'n_estimators': 300, 'learning_rate': 0.5, 'min_child_weight': 1, 'gamma': 0.5, 'max_depth': 3}	0.9967	0.869	0.9967	0.8827

Summary

Given a directed social graph, we had to predict missing links to recommend users.

The Dataset contains 2 columns - Source Node and Destination Node.

The graph based dataset was converted to supervised classification learning problem. I generated training samples of good and bad links from given directed graph and for each link got some features like no of followers, has he followed back, page rank, katz score, adar index, some svd fetures of adj matrix, some weight features, preferential attachment, svd dot etc. and trained ml model based on these features to predict link.

I deployed two Models - RandomForest and XGBoost with hyperparameter tuning using RandomizedSearchCV. Each model's performance was assessed using F1-score and AUC score.

XGBoost with extreme hypertuning was overfitting to a little extent but RandomForest with hyperparameter tuning was performing the best with test f1 score of 0.92 and auc score of 0.93.