# Facebook Friend Recommendation - Social network Graph Link Prediction

#### **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 recruiting challenge on kaggle <a href="https://www.kaggle.com/c/FacebookRecruiting">https://www.kaggle.com/c/FacebookRecruiting</a> data contains two columns source and destination eac edge in graph

```
- Data columns (total 2 columns):
- source_node : int64
- destination 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-message-attachments/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 ighest probability links

#### Performance metric for supervised learning:

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

```
#Importing Libraries
# please do go through this python notebook:
import warnings
warnings.filterwarnings("ignore")
{\bf import\ pandas\ as\ pd} \# pandas\ to\ create\ small\ data frames
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
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
```

```
Mounting Drive
In [0]:
!kill -9 -1
In [2]:
 from google.colab import drive
drive.mount('/content/drive')
Go to this URL in a browser: https://accounts.google.com/o/oauth2/auth?client id=947318989803-6bn6
\verb|qk8| qdgf4n4g3pfee6491hc0brc4i.apps.google user content.com&redirect\_uri=urn&3Aietf&3Awg&3Aoauth&3A2.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&accorder=0.0&ac
2Fauth%2Fdrive%20https%3A%2F%2Fwww.googleapis.com%2Fauth%2Fdrive.photos.readonly%20https%3A%2F%2Fww
ogleapis.com%2Fauth%2Fpeopleapi.readonly&response type=code
Enter your authorization code:
Mounted at /content/drive
In [3]:
 !pwd
/content
drive sample data
In [4]:
import os
 PATH = os.getcwd()
print(PATH)
/content
In [5]:
data path = PATH + '/drive/My Drive/AAIC/Case Studies/Facebook Friend Recommendation/'
data_path
Out[5]:
 '/content/drive/My Drive/AAIC/Case Studies/Facebook Friend Recommendation/'
```

```
print(nx.info(g))
Name:
Type: DiGraph
Number of nodes: 1862220
Number of edges: 9437519
Average in degree: 5.0679
                     5.0679
Average out degree:
      Displaying a sub graph
In [0]:
if not os.path.isfile(data_path + 'train_woheader_sample.csv'):
    pd.read_csv(data_path + 'data/train.csv', nrows=50).to_csv(data_path +
'train woheader sample.csv', header=False, index=False)
subgraph=nx.read_edgelist(data_path + 'train_woheader_sample.csv',delimiter=',',create_using=nx.Di
Graph(),nodetype=int)
{\#\ https://stackoverflow.com/questions/9402255/drawing-a-huge-graph-with-networkx-and-matplotlib}
pos=nx.spring_layout(subgraph)
nx.draw(subgraph,pos,node_color='#A0CBE2',edge_color='#00bb5e',width=1,edge_cmap=plt.cm.Blues,with_
labels=True)
plt.savefig("graph sample.pdf")
print(nx.info(subgraph))
Name:
Type: DiGraph
Number of nodes: 66
Number of edges: 50
Average in degree: 0.7576
Average out degree: 0.7576
                   6247296188602941522
6247296188602941522
1645927 20603278432
                                        1194519
                                        382654589497
    1657/60904
    690569
                                             1576703
                        23 5
   16468769
                                              1246523
   17718471
                                               677005
   296377
                                            456059456
    328678
      189226
                                           531778
```

# 1. Exploratory Data Analysis

x.prgraph(), nodecype=inc)

```
In [0]:
```

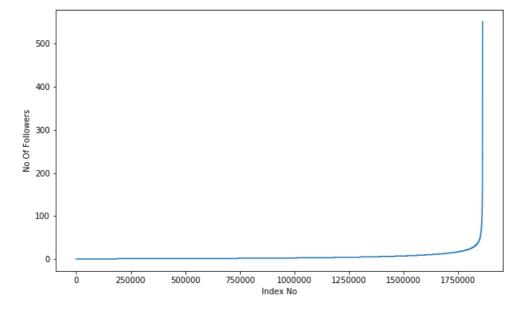
```
# No of Unique persons
print("The number of unique persons",len(g.nodes()))
```

The number of unique persons 1862220

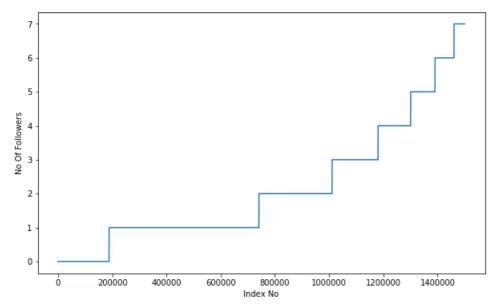
### 1.1 No of followers for each person

```
In [0]:
```

```
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()
```



```
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()
```



```
plt.boxplot(indegree_dist)
plt.ylabel('No Of Followers')
plt.show()
```

```
### 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.

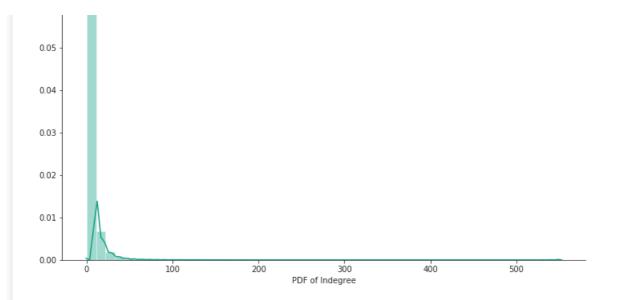
#### In [0]:

```
### 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
```

```
%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()
```

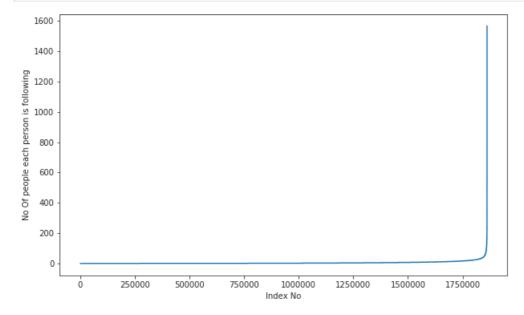
```
0.08 -
```



# 1.2 No of people each person is following

#### In [0]:

```
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()
```

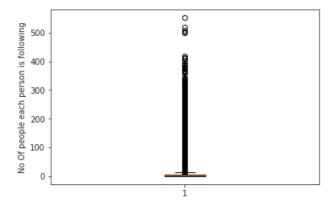


```
indegree_dist = list(dict(g.in_degree()).values())
indegree_dist.sort()
plt.figure(figsize=(10,6))
plt.plot(outdegree_dist[0:1500000])
plt.xlabel('Index No')
plt.ylabel('No Of people each person is following')
plt.show()
```

```
7 -
6 -
Bus
```

```
ON 1 - 0 200000 400000 600000 800000 1000000 1200000 1400000 Index No
```

```
plt.boxplot(indegree_dist)
plt.ylabel('No Of people each person is following')
plt.show()
```



#### In [0]:

```
### 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
```

```
### 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

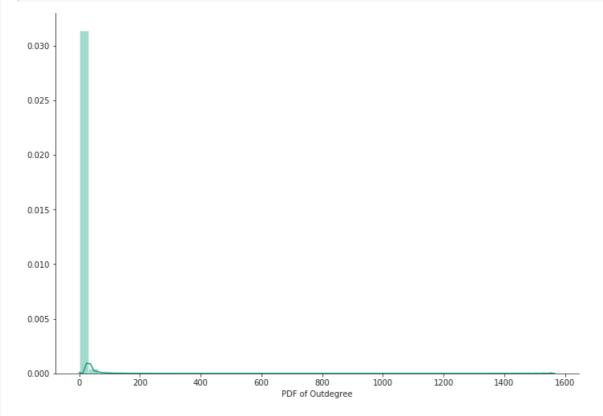
99.8 percentile value is 90.0

99.9 percentile value is 123.0

100.0
```

```
100.0 percentile value is 1566.0
```

```
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()
```



#### In [0]:

No of persons those are not following anyone are 274512 and % is 14.741115442858524

#### In [0]:

No of persons having zero followers are 188043 and % is 10.097786512871734

#### In [0]:

No of persons those are not not following anyone and also not having any followers are 0

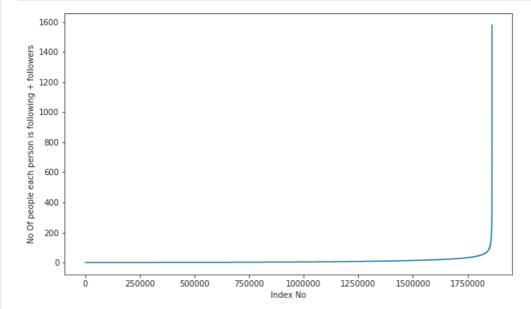
### 1.3 both followers + following

```
In [0]:
```

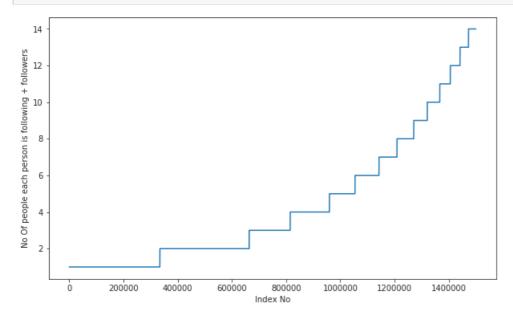
```
from collections import Counter
dict_in = dict(g.in_degree())
dict_out = dict(g.out_degree())
d = Counter(dict_in) + Counter(dict_out)
in_out_degree = np.array(list(d.values()))
```

#### In [0]:

```
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_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 [0]:
### 90-100 percentile
for i in range (0,11):
   print(90+i,'percentile value is',np.percentile(in_out_degree_sort,90+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 [0]:
### 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 [0]:
print('Min of no of followers + following is',in out degree.min())
print(np.sum(in out degree==in out degree.min()),' persons having minimum no of followers +
following')
Min of no of followers + following is 1
334291 persons having minimum no of followers + following
In [0]:
print('Max of no of followers + following is',in out degree.max())
print(np.sum(in out degree==in out degree.max()),' persons having maximum no of followers +
following')
Max of no of followers + following is 1579
1 persons having maximum no of followers + following
In [0]:
print('No of persons having followers + following less than 10 are',np.sum(in out degree<10))
No of persons having followers + following less than 10 are 1320326
In [0]:
print('No of weakly connected components',len(list(nx.weakly connected components(g))))
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 [0]:
%%time
###generating bad edges from given graph
import random
if not os.path.isfile(data path + 'data/after eda/missing edges final.p'):
    #getting all set of edges
    r = csv.reader(open(data_path + '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
                    missing_edges.add((a,b))
        else:
            continue
    pickle.dump (missing_edges,open (data_path + 'data/after_eda/missing_edges_final.p','wb'))
else:
    missing edges = pickle.load(open(data path + 'data/after eda/missing edges final.p','rb'))
CPU times: user 2.46 s, sys: 1.29 s, total: 3.75 s
Wall time: 5.65 s
In [0]:
len(missing edges)
Out[0]:
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

```
from sklearn.model_selection import train_test_split
if (not os.path.isfile(data_path + 'data/after_eda/train_pos_after_eda.csv')) and (not
os.path.isfile(data_path + 'data/after_eda/test_pos_after_eda.csv')):
    #reading total data df
    df_pos = pd.read_csv(data_path + '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 positive training data only for c
reating graph
    #and for feature generation
    X train pos, X test pos, y train pos, y test pos = train test split(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_split(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 po
s.shape[0])
   print("Number of nodes in the train data graph without edges", X train neg.shape[0],"=", y trai
n_neg.shape[0])
   print('='*60)
   print("Number of nodes in the test data graph with edges", X test pos.shape[0], "=",y test pos.s
hape[0])
    print ("Number of nodes in the test data graph without edges",
X test neg.shape[0], "=", y test neg.shape[0])
    #removing header and saving
    X train pos.to csv(data path + 'data/after eda/train pos after eda.csv',header=False, index=Fal
se)
    X test pos.to csv(data path + 'data/after eda/test pos after eda.csv',header=False, index=False
    X train neg.to csv(data path + 'data/after eda/train neg after eda.csv',header=False, index=Fal
    X_test_neg.to_csv(data_path + 'data/after_eda/test_neg_after_eda.csv',header=False, index=False
else:
    #Graph from Traing data only
    del missing edges
4
Number of nodes in the graph with edges 9437519
Number of nodes in the graph without edges 9437519
______
Number of nodes in the train data graph with edges 7550015 = 7550015
Number of nodes in the train data graph without edges 7550015 = 7550015
_____
Number of nodes in the test data graph with edges 1887504 = 1887504
Number of nodes in the test data graph without edges 1887504 = 1887504
In [0]:
if (os.path.isfile(data path + 'data/after eda/train pos after eda.csv')) and
(os.path.isfile(data path + 'data/after eda/test pos after eda.csv')):
    train graph=nx.read edgelist(data path +
'data/after eda/train pos after eda.csv',delimiter=',',create using=nx.DiGraph(),nodetype=int)
   test graph=nx.read edgelist(data path + 'data/after eda/test pos after eda.csv',delimiter=',',c
reate_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 -- ',trY teN)
    print('no of people present in test but not present in train -- ',teY trN)
    print(' % of people not there in Train but exist in Test in total Test data are {} %'.format(te
  trN/len(test nodes pos)*100))
                                                                                             ▶
Name:
```

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 are 7.1200735962845405 %
```

#### we have a cold start problem here

#### In [0]:

```
#final train and test data sets
if (not os.path.isfile(data path + 'data/after eda/train after eda.csv')) and \
(not os.path.isfile(data path + 'data/after eda/test after eda.csv')) and \
(not os.path.isfile(data path + 'data/train y.csv')) and \
(not os.path.isfile(data path + 'data/test y.csv')) and \
(os.path.isfile(data_path + 'data/after_eda/train_pos_after_eda.csv')) and \
(os.path.isfile(data_path + 'data/after_eda/test_pos_after_eda.csv')) and \
(os.path.isfile(data_path + 'data/after_eda/train_neg_after_eda.csv')) and \
(os.path.isfile(data_path + 'data/after_eda/test_neg_after_eda.csv')):
    X train pos = pd.read csv(data path + 'data/after eda/train pos after eda.csv', names=['source
node', 'destination node'])
   X test pos = pd.read csv(data path + 'data/after eda/test pos after eda.csv', names=['source no
de', 'destination node'])
   X_train_neg = pd.read_csv(data_path + 'data/after_eda/train_neg_after_eda.csv', names=['source_
node', 'destination node'])
   X test neg = pd.read csv(data path + 'data/after eda/test neg after eda.csv', names=['source no
de', '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 train neg.shape[0])
    print('='*60)
    print("Number of nodes in the test data graph with edges", X test pos.shape[0])
    print("Number of nodes in the test data graph without edges", X test 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 path + 'data/after eda/train after eda.csv',header=False,index=False)
    X test.to csv(data path + 'data/after eda/test after eda.csv',header=False,index=False)
    pd.DataFrame(y train.astype(int)).to csv(data path + 'data/train y.csv',header=False,index=False
e)
    pd.DataFrame(y test.astype(int)).to csv(data path + 'data/test y.csv',header=False,index=False)
```

#### In [0]:

```
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)
```

D-F- --;-F- ;- F--;- 3-F- (16100000 0)

```
Data points in train data (15100030, 2)
Data points in test data (3775008, 2)
Shape of traget variable in train (15100030,)
Shape of traget variable in test (3775008,)
```

# 3. Reading Data

```
In [0]:
```

```
if os.path.isfile(data_path + 'data/after_eda/train_pos_after_eda.csv'):
    train_graph=nx.read_edgelist(data_path +
    'data/after_eda/train_pos_after_eda.csv',delimiter=',',create_using=nx.DiGraph(),nodetype=int)
    print(nx.info(train_graph))
else:
    print("please run the FB_EDA.ipynb or download the files from drive")
```

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

# 4. Similarity measures

### 4.1 Jaccard Distance:

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

\begin{equation} j = \frac{|X\cap Y|}{|X \cup Y|} \end{equation}

```
In [0]:
```

```
In [0]:
```

```
#one test case
print(jaccard_for_followees(273084,1505602))
```

0.0

#### In [0]:

```
#node 1635354 not in graph
print(jaccard_for_followees(273084,1505602))
```

0.0

```
if len(set(train_graph.predecessors(a))) == 0 | len(set(g.predecessors(b))) == 0:
       sim = (len(set(train_graph.predecessors(a)).intersection(set(train_graph.predecessors(b)))))
)/\
                                (len(set(train graph.predecessors(a)).union(set(train graph.predec
ssors(b)))))
       return sim
    except:
       return 0
4
In [0]:
print(jaccard for followers(273084,470294))
0
In [0]:
#node 1635354 not in graph
print(jaccard for followees(669354,1635354))
0
4.2 Cosine distance
In [0]:
#for followees
def cosine for followees(a,b):
       if len(set(train graph.successors(a))) == 0 | len(set(train graph.successors(b))) == 0:
           return 0
        sim = (len(set(train graph.successors(a)).intersection(set(train graph.successors(b)))))/\
(math.sqrt(len(set(train_graph.successors(a)))*len((set(train_graph.successors(b))))))
       return sim
    except:
       return 0
In [0]:
print(cosine for followees(273084,1505602))
0.0
In [0]:
print(cosine for followees(273084,1635354))
0
In [0]:
def cosine for followers(a,b):
       if len(set(train graph.predecessors(a))) == 0 | len(set(train graph.predecessors(b))) == 0
•
       sim = (len(set(train graph.predecessors(a)).intersection(set(train graph.predecessors(b))))
)/\
                                    (math.sqrt(len(set(train graph.predecessors(a)))) * (len(set(tra
n_graph.predecessors(b)))))
       return sim
```

```
return 0

In [0]:

print (cosine_for_followers(2,470294))

0.02886751345948129

In [0]:

print (cosine_for_followers(669354,1635354))

0
```

# 5. 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.

### 5.1 Page Ranking

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

```
In [0]:
if not os.path.isfile(data path + 'data/fea sample/page rank.p'):
    pr = nx.pagerank(train graph, alpha=0.85)
    pickle.dump(pr,open(data_path + 'data/fea_sample/page_rank.p','wb'))
    pr = pickle.load(open(data_path + 'data/fea_sample/page_rank.p','rb'))
In [0]:
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 2.7098251341935827e-05
mean 5.615699699389075e-07
In [0]:
#for imputing to nodes which are not there in Train data
mean_pr = float(sum(pr.values())) / len(pr)
print(mean_pr)
```

### 6. Other Graph Features

5.615699699389075e-07

### 6.1 Shortest path:

In [0]:

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

```
#if has direct edge then deleting that edge and calculating shortest path
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)
            p= nx.shortest_path_length(train_graph,source=a,target=b)
        return p
    except:
        return -1
In [0]:
compute shortest path length (77697, 826021)
Out[0]:
10
In [0]:
#testing
compute_shortest_path_length(669354,1635354)
Out[0]:
```

### 6.2 Checking for same community

```
In [0]:
```

-1

```
#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):
       {f 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 [0]:
belongs_to_same_wcc(861, 1659750)

Out[0]:
0

In [0]:
belongs_to_same_wcc(669354,1635354)

Out[0]:
0
```

### 6.3 Adamic/Adar Index:

except:

return 0

Adamic/Adar measures is defined as inverted sum of degrees of common neighbours for given two vertices.  $A(x,y)=\sum_{u \in N(y)}\frac{1}{\log(|N(u)|)}$ 

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

# 6.4 Is person was following back:

```
def follows_back(a,b):
    if train_graph.has_edge(b,a):
        return 1
    else:
```

```
return 0
 In [0]:
 follows back(1,189226)
Out[0]:
1
 In [0]:
 follows back (669354, 1635354)
Out[0]:
6.5 Katz Centrality:
https://en.wikipedia.org/wiki/Katz_centrality
https://www.geeksforgeeks.org/katz-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 = \alpha \cdot i = \alpha \cdot i + \beta \cdot i + 
where A is the adjacency matrix of the graph G with eigenvalues $$\lambda$$.
The parameter $$\beta$$ controls the initial centrality and
\ \lambda \frac{1}{\lambda \max}\.\$
 In [0]:
 if not os.path.isfile(data path + 'data/fea sample/katz.p'):
                      katz = nx.katz.katz centrality(train graph,alpha=0.005,beta=1)
                      pickle.dump(katz,open(data_path + 'data/fea_sample/katz.p','wb'))
 else:
                      katz = pickle.load(open(data_path + 'data/fea_sample/katz.p','rb'))
 In [0]:
print('min', katz[min(katz, key=katz.get)])
 print('max', katz[max(katz, key=katz.get)])
 print('mean',float(sum(katz.values())) / len(katz))
min 0.0007313532484065916
max 0.003394554981699122
mean 0.0007483800935562018
```

```
In [0]:
```

```
mean_katz = float(sum(katz.values())) / len(katz)
print(mean_katz)
```

0.0007483800935562018

#### 6.6 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 [0]:
```

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

```
print('min',hits[0][min(hits[0], key=hits[0].get)])
print('max',hits[0][max(hits[0], key=hits[0].get)])
print('mean',float(sum(hits[0].values())) / len(hits[0]))

min 0.0
max 0.004868653378780953
mean 5.615699699344123e-07
```

### 7. Featurization

### 7. 1 Reading a sample of Data from both train and test

Number of rows we are going to elimiate in test data are 3725006

#### In [0]:

```
import random
if os.path.isfile(data_path + 'data/after_eda/train_after_eda.csv'):
    filename = data_path + "data/after_eda/train_after_eda.csv"
    # you uncomment this line, if you dont know the lentgh 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 [0]:

```
if os.path.isfile(data_path + 'data/after_eda/train_after_eda.csv'):
    filename = data_path + "data/after_eda/test_after_eda.csv"
    # you uncomment this line, if you dont know the lentph 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 [0]:

```
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(skip_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(skip_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
```

```
df_final_train = pd.read_csv(data_path + 'data/after_eda/train_after_eda.csv', skiprows=skip_train,
names=['source_node', 'destination_node'])
df_final_train['indicator_link'] = pd.read_csv(data_path + 'data/train_y.csv', skiprows=skip_train,
names=['indicator_link'])
print("Our train matrix size ",df_final_train.shape)
df_final_train.head(2)
```

our crum mucrin orde (100002, 0,

#### Out[0]:

	source_node	destination_node	indicator_link	
0	273084	1505602	1	

1543415

#### In [0]:

832016

```
df_final_test = pd.read_csv(data_path + 'data/after_eda/test_after_eda.csv', skiprows=skip_test,
names=['source_node', 'destination_node'])
df_final_test['indicator_link'] = pd.read_csv(data_path + 'data/test_y.csv', skiprows=skip_test, na
mes=['indicator_link'])
print("Our test matrix size ",df_final_test.shape)
df_final_test.head(2)
```

Our test matrix size (50002, 3)

#### Out[0]:

### source\_node destination\_node indicator\_link

0	848424	784690	1
1	483294	1255532	1

### 7.2 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. num\_followers\_s6. num\_followees\_s
- 7. num\_followers\_d
- 8. num followees d
- 9. inter followers
- 10. inter\_followees

```
if not os.path.isfile(data_path + 'data/fea_sample/storage_sample_stagel.h5'):
    #mapping_jaccrd_followers to train_and_test_data
    df_final_train['jaccard_followers'] = df_final_train.apply(lambda_row:

jaccard_for_followers(row['source_node'],row['destination_node']),axis=1)
    df_final_test['jaccard_followers'] = df_final_test.apply(lambda_row:

jaccard_for_followers(row['source_node'],row['destination_node']),axis=1)

    #mapping_jaccrd_followees_to_train_and_test_data
    df_final_train['jaccard_followees'] = df_final_train.apply(lambda_row:

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_jaccrd_followers_to_train_and_test_data
    df_final_train['cosine_followers'] = df_final_train.apply(lambda_row:

cosine_for_followers(row['source_node'],row['destination_node']),axis=1)
    df_final_test['cosine_followers'] = df_final_test.apply(lambda_row:
```

```
cosine for followers(row['source node'], row['destination node']), axis=1)
    #mapping jaccrd followees to train and test data
    df final train['cosine followees'] = df final train.apply(lambda row:
cosine for followees(row['source node'], row['destination node']), axis=1)
    df_final_test['cosine_followees'] = df_final_test.apply(lambda row:
cosine for followees(row['source node'],row['destination node']),axis=1)
In [0]:
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():
            s1=set(train graph.predecessors(row['source node']))
            s2=set(train graph.successors(row['source node']))
        except:
            s1 = set()
            s2 = set()
        trv:
            d1=set(train graph.predecessors(row['destination node']))
            d2=set(train graph.successors(row['destination node']))
        except:
            d1 = set()
            d2 = set()
        {\tt 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 followees d, inter followers, int
er followees
4
                                                                                                   | b |
In [0]:
if not os.path.isfile(data_path + 'data/fea_sample/storage_sample_stage1.h5'):
    df_final_train['num_followers_s'], df_final_train['num_followers_d'], \
    {\tt 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(d
f_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 stage1(df final test)
    hdf = HDFStore('data/fea sample/storage sample stage1.h5')
    hdf.put('train_df',df_final_train, format='table', data_columns=True)
    hdf.put('test df',df final test, format='table', data columns=True)
    hdf.close()
else:
    df final train = read hdf(data path + 'data/fea sample/storage sample stage1.h5', 'train df',mo
de='r')
    df final test = read hdf(data path + 'data/fea sample/storage sample stage1.h5', 'test df', mode
```

# 7.3 Adding new set of features

4

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

```
if not os.path.isfile(data_path + 'data/fea_sample/storage_sample_stage2.h5'):
    #mapping adar index on train
    df final train['adar index'] = df final train.apply(lambda row: calc 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'],r
ow['destination node']),axis=1)
    #mapping followback or not on train
    df final train['follows back'] = df final train.apply(lambda row:
follows_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: follows 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: belongs 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: belongs_to_same_wcc(row['source_no
de'], 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']), axis=1)
    #mapping shortest path on test
    df final test['shortest path'] = df final test.apply(lambda row: compute shortest path length(r
ow['source node'], row['destination node']), axis=1)
    hdf = HDFStore(data path + 'data/fea sample/storage sample stage2.h5')
    hdf.put('train_df',df_final_train, format='table', data_columns=True)
    hdf.put('test_df',df_final_test, format='table', data_columns=True)
   hdf.close()
else:
   df_final_train = read_hdf(data_path + 'data/fea_sample/storage_sample_stage2.h5', 'train_df',mo
de='r')
   df final test = read hdf(data path + 'data/fea sample/storage sample stage2.h5', 'test df',mode
='r')
```

# 7.4 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

 $\label{eq:weighted} $$ \operatorname{quation} W = \frac{1}{\sqrt{1+|X|}} \end{equation} $$$ 

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

#### In [0]:

```
#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()))
100%|
                                                                          | 1780722/1780722
[00:11<00:00, 152682.24it/s]
```

#### In [0]:

```
if not os.path.isfile(data path + '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,m
ean weight in))
   df final train['weight out'] = df final train.source node.apply(lambda x: Weight out.get(x,mean
weight out))
    #mapping to pandas test
   df_final_test['weight_in'] = df_final_test.destination_node.apply(lambda x: Weight in.get(x,mea
n weight in))
   df_final_test['weight_out'] = df_final_test.source_node.apply(lambda x: Weight_out.get(x,mean_w
eight out))
    #some features engineerings on the in and out weights
   df final train['weight_f1'] = df_final_train.weight_in + df_final_train.weight_out
   df final train['weight f2'] = df final train.weight in * df final train.weight out
   df final train['weight f3'] = (2*df final train.weight in + 1*df final train.weight out)
   df final train['weight f4'] = (1*df final train.weight in + 2*df final train.weight out)
    #some features engineerings on the in and out weights
   df_final_test['weight_f1'] = df_final_test.weight_in + df_final_test.weight_out
   df final test['weight f2'] = df final test.weight in * df final test.weight out
   df_final_test['weight_f3'] = (2*df_final_test.weight_in + 1*df_final_test.weight_out)
   df_final_test['weight_f4'] = (1*df_final_test.weight_in + 2*df_final_test.weight_out)
```

```
if not os.path.isfile(data_path + '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.apply(lambda x:pr.get(x,mean pr))
    df final train['page rank d'] = df final train.destination node.apply(lambda x:pr.get(x,mean pr
))
    df final test['page rank s'] = df final test.source node.apply(lambda 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(lambda 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(lambda 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(lambda 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(lambda 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].get(x,0))
    df_final_train['authorities_d'] = df_final_train.destination_node.apply(lambda x: hits[1].get(x
, 0))
     \texttt{df final test['authorities s']} = \texttt{df\_final\_test.source\_node.apply(lambda x: hits[1].get(x,0))} 
    df final test['authorities d'] = df final test.destination node.apply(lambda x: hits[1].get(x,0
) )
    hdf = HDFStore(data_path + 'data/fea_sample/storage_sample_stage3.h5')
    hdf.put('train df', df final train, format='table', data columns=True)
    hdf.put('test df',df final test, format='table', data columns=True)
   hdf.close()
else:
   df final train = read hdf(data path + 'data/fea sample/storage sample stage3.h5', 'train df',mo
de='r')
   df final test = read hdf(data path + 'data/fea sample/storage sample stage3.h5', 'test df', mode
='r')
4
                                                                                                 •
```

### 7.5 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 [0]:

def svd(x, S):
    try:
        z = sadj_dict[x]
        return S[z]
    except:
        return [0,0,0,0,0,0]
```

```
In [0]:

#for svd features to get feature vector creating a dict node val and inedx in svd vector
sadj_col = sorted(train_graph.nodes())
sadj_dict = { val:idx for idx,val in enumerate(sadj_col)}
```

```
Adj = nx.adjacency_matrix(train_graph,nodelist=sorted(train_graph.nodes())).asfptype()
```

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[['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 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[['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 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[['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.source\_node.apply(lambda x: svd(x, V.T)).apply(pd.Series)

df final test.source node.apply(lambda x: svd(x, U)).apply(pd.Series)

df final test.destination node.apply(lambda x: svd(x, U)).apply(pd.Series)

df final test.source node.apply(lambda x: svd(x, V.T)).apply(pd.Series)

hdf = HDFStore(data\_path + 'data/fea\_sample/storage\_sample\_stage4.h5')
hdf.put('train\_df',df\_final\_train, format='table', data\_columns=True)
hdf.put('test df',df final test, format='table', data columns=True)

hdf.close()

4

df final test.destination node.apply(lambda x: svd(x, V.T)).apply(pd.Series)

df final train.destination node.apply(lambda x: svd(x, V.T)).apply(pd.Series)

# 8. Modelling the data

```
In [0]:
#reading
from pandas import read hdf
df final train = read hdf(data path + 'data/fea sample/storage sample stage4.h5', 'train df', mode='
df final test = read hdf(data path + 'data/fea sample/storage sample stage4.h5',
'test df',mode='r')
In [0]:
df final train.columns
Out[0]:
Index(['source node', 'destination node', 'indicator link',
        'jaccard_followers', 'jaccard_followees', 'cosine_followers', 'cosine_followees', 'num_followers_s', 'num_followees_s',
        'num followees d', 'inter_followers', 'inter_followees', 'adar_index',
        'follows_back', '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'],
       dtype='object')
In [0]:
y train = df final train.indicator link
y_test = df_final_test.indicator_link
In [0]:
df_final_train.drop(['source_node', 'destination_node','indicator_link'],axis=1,inplace=True)
df final test.drop(['source node', 'destination_node', 'indicator_link'], axis=1, inplace=True)
8.1 Random Forest
In [0]:
estimators = [10,50,100,250,450]
train scores = []
test scores = []
for i in estimators:
     clf = RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
              max depth=5, max features='auto', max leaf nodes=None,
              min_impurity_decrease=0.0, min_impurity_split=None,
              min samples leaf=52, min samples split=120,
              min weight fraction leaf=0.0, n estimators=i, n jobs=-1,random state=25,verbose=0,warm
start=False)
    clf.fit(df final train,y train)
    train_sc = f1_score(y_train,clf.predict(df_final_train))
    test_sc = f1_score(y_test,clf.predict(df_final_test))
     test scores.append(test sc)
    train_scores.append(train_sc)
     print('Estimators = ',i,'Train Score',train sc,'test Score',test sc)
plt.plot(estimators, train scores, label='Train Score')
```

Estimators = 10 Train Score 0.9063252121775113 test Score 0.8745605278006858 Estimators = 50 Train Score 0.9205725512208812 test Score 0.9125653355634538 Estimators = 100 Train Score 0.9238690848446947 test Score 0.9141199714153599

plt.plot(estimators,test\_scores,label='Test Score')

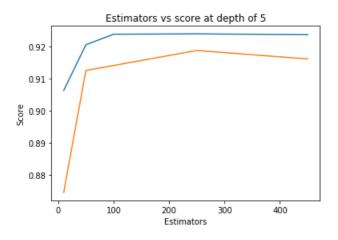
plt.title('Estimators vs score at depth of 5')

plt.xlabel('Estimators')
plt.ylabel('Score')

```
Estimators = 250 Train Score 0.9239789348046863 test Score 0.9188007232664732 Estimators = 450 Train Score 0.9237190618658074 test Score 0.9161507685828595
```

#### Out[0]:

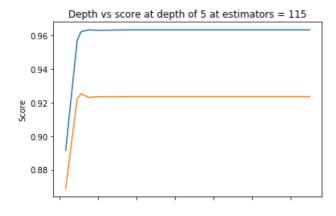
Text(0.5,1,'Estimators vs score at depth of 5')



#### In [0]:

```
depths = [3, 9, 11, 15, 20, 35, 50, 70, 130]
train scores = []
test_scores = []
for i in depths:
    clf = RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
            max_depth=i, max_features='auto', max_leaf_nodes=None,
            min impurity decrease=0.0, min impurity split=None,
            min samples leaf=52, min samples split=120,
            min_weight_fraction_leaf=0.0, n_estimators=115, n_jobs=-1,random_state=25,verbose=0,war
m start=False)
    clf.fit(df final train,y train)
    train sc = f1 score(y train,clf.predict(df final train))
    test sc = f1 score(y test,clf.predict(df final test))
    test_scores.append(test_sc)
    train_scores.append(train_sc)
    print('depth = ',i,'Train Score',train_sc,'test Score',test_sc)
plt.plot(depths,train_scores,label='Train Score')
plt.plot(depths,test scores,label='Test Score')
plt.xlabel('Depth')
plt.ylabel('Score')
plt.title('Depth vs score at depth of 5 at estimators = 115')
plt.show()
```

depth = 3 Train Score 0.8916120853581238 test Score 0.8687934859875491
depth = 9 Train Score 0.9572226298198419 test Score 0.9222953031452904
depth = 11 Train Score 0.9623451340902863 test Score 0.9252318758281279
depth = 15 Train Score 0.9634267621927706 test Score 0.9231288356496615
depth = 20 Train Score 0.9631629153051491 test Score 0.9235051024711141
depth = 35 Train Score 0.9634333127085721 test Score 0.9235601652753184
depth = 50 Train Score 0.9634333127085721 test Score 0.9235601652753184
depth = 70 Train Score 0.9634333127085721 test Score 0.9235601652753184
depth = 130 Train Score 0.9634333127085721 test Score 0.9235601652753184



```
20 40
             60
                       100
                            120
                  80
             Depth
```

```
In [0]:
from sklearn.metrics import f1_score
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import f1 score
from sklearn.model_selection import RandomizedSearchCV
from scipy.stats import randint as sp_randint
from scipy.stats import uniform
param_dist = {"n_estimators":sp_randint(105,125),
              "max depth": sp randint(10,15),
              "min_samples_split": sp_randint(110,190),
              "min_samples_leaf": sp_randint(25,65)}
clf = RandomForestClassifier(random state=25,n jobs=-1)
rf_random = RandomizedSearchCV(clf, param_distributions=param_dist,
                                   n_iter=5,cv=10,scoring='f1',random_state=25)
rf_random.fit(df_final_train,y_train)
print('mean test scores',rf_random.cv_results_['mean_test_score'])
print('mean train scores',rf random.cv results ['mean train score'])
mean test scores [0.96225043 0.96215493 0.96057081 0.96194015 0.96330005]
mean train scores [0.96294922 0.96266735 0.96115674 0.96263457 0.96430539]
In [0]:
print(rf_random.best_estimator_)
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
            max depth=14, max features='auto', max leaf nodes=None,
            min_impurity_decrease=0.0, min_impurity_split=None,
            min_samples_leaf=28, min_samples_split=111,
            min_weight_fraction_leaf=0.0, n_estimators=121, n_jobs=-1,
            oob score=False, random state=25, verbose=0, warm start=False)
In [0]:
clf = RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
            max depth=14, max features='auto', max leaf nodes=None,
            min_impurity_decrease=0.0, min_impurity_split=None,
            min samples leaf=28, min samples split=111,
            min weight fraction leaf=0.0, n estimators=121, n jobs=-1,
            oob score=False, random state=25, verbose=0, warm start=False)
In [0]:
clf.fit(df final train, y train)
y_train_pred = clf.predict(df_final_train)
```

```
y test pred = clf.predict(df final test)
```

```
from sklearn.metrics import f1 score
print('Train f1 score', f1 score(y train, y train pred))
print('Test f1 score',f1_score(y_test,y_test_pred))
```

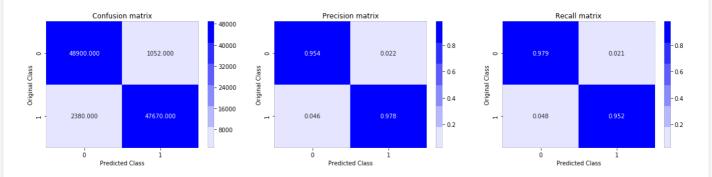
Train f1 score 0.9652533106548414 Test f1 score 0.9241678239279553

```
from sklearn.metrics import confusion matrix
def plot_confusion_matrix(test_y, predict_y):
   C = confusion matrix(test y, predict y)
```

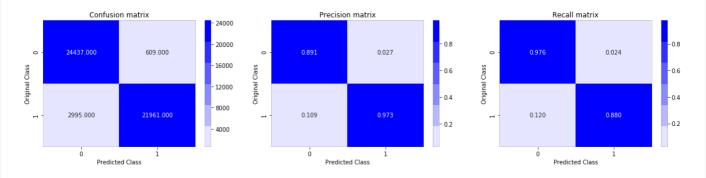
```
A = (((C.T) / (C.sum(axis=1))).T)
B = (C/C.sum(axis=0))
plt.figure(figsize=(20,4))
labels = [0,1]
# representing A in heatmap format
cmap=sns.light_palette("blue")
plt.subplot(1, 3, 1)
sns.heatmap(C, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels, yticklabels=labels)
plt.xlabel('Predicted Class')
plt.ylabel('Original Class')
plt.title("Confusion matrix")
plt.subplot(1, 3, 2)
sns.heatmap(B, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels, yticklabels=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()
```

```
print('Train confusion_matrix')
plot_confusion_matrix(y_train_pred)
print('Test confusion_matrix')
plot_confusion_matrix(y_test,y_test_pred)
```

Train confusion matrix

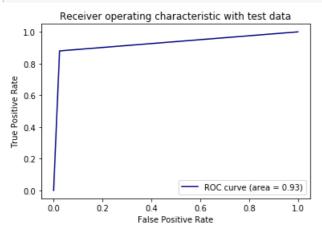


Test confusion\_matrix

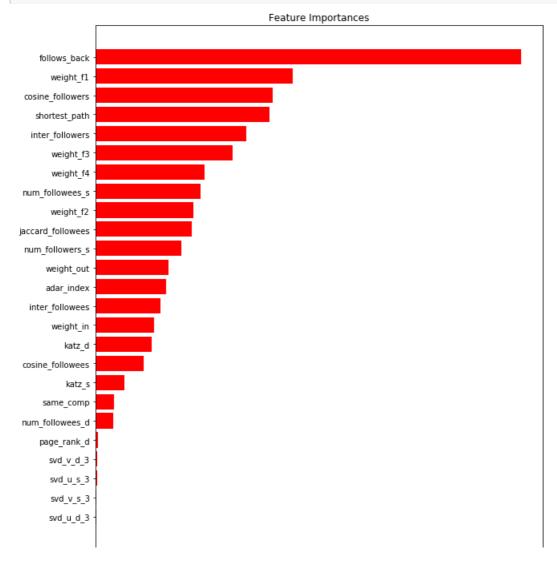


```
from sklearn.metrics import roc_curve, auc
fpr,tpr,ths = roc_curve(y_test,y_test_pred)
auc_sc = auc(fpr, tpr)
plt.plot(fpr, tpr, color='navy',label='ROC curve (area = %0.2f)' % auc_sc)
```

```
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic with test data')
plt.legend()
plt.show()
```



```
features = df_final_train.columns
importances = clf.feature_importances_
indices = (np.argsort(importances))[-25:]
plt.figure(figsize=(10,12))
plt.title('Feature Importances')
plt.barh(range(len(indices)), importances[indices], color='r', align='center')
plt.yticks(range(len(indices)), [features[i] for i in indices])
plt.xlabel('Relative Importance')
plt.show()
```



# 9. Assignments:

- 1. Add another feature called Preferential Attachment with followers and followees data of vertex. you can check about Preferential Attachment in below link <a href="http://be.amazd.com/link-prediction/">http://be.amazd.com/link-prediction/</a>
- Add feature called svd\_dot. you can calculate svd\_dot as Dot product between sourse node svd and destination node svd features. you can read about this in below pdf <a href="https://storage.googleapis.com/kaggle-forum-message-attachments/2594/supervised link prediction.pdf">https://storage.googleapis.com/kaggle-forum-message-attachments/2594/supervised link prediction.pdf</a>
- 3. Tune hyperparameters for XG boost with all these features and check the error metric.

```
In [0]:
```

```
#reading
from pandas import read_hdf
df_final_train = read_hdf(data_path + 'data/fea_sample/storage_sample_stage4.h5', 'train_df',mode='
r')
df_final_test = read_hdf(data_path + 'data/fea_sample/storage_sample_stage4.h5',
'test_df',mode='r')
```

#### In [7]:

```
df_final_train.columns
```

#### Out[7]:

#### In [0]:

```
y_train = df_final_train.indicator_link
y_test = df_final_test.indicator_link
```

#### In [9]:

```
df_final_train.head()
```

#### Out[9]:

	source_node	destination_node	indicator_link	jaccard_followers	jaccard_followees	cosine_followers	cosine_followees	num_followe
0	273084	1505602	1	0	0.000000	0.000000	0.000000	
1	832016	1543415	1	0	0.187135	0.028382	0.343828	
2	1325247	760242	1	0	0.369565	0.156957	0.566038	
3	1368400	1006992	1	0	0.000000	0.000000	0.000000	
4	140165	1708748	1	0	0.000000	0.000000	0.000000	

```
source_node destination_node indicator_link jaccard_followers jaccard_followees cosine_followers cosine_followees num_followe
4
In [10]:
if os.path.isfile(data_path + 'data/after_eda/train_pos_after_eda.csv') and
os.path.isfile(data path + 'data/after eda/test pos after eda.csv'):
    train graph=nx.read edgelist(data path +
'data/after eda/train pos after eda.csv',delimiter=',',create using=nx.DiGraph(),nodetype=int)
    test_graph=nx.read_edgelist(data_path + 'data/after_eda/test_pos_after_eda.csv',delimiter=',',c
reate_using=nx.DiGraph(),nodetype=int)
    print(nx.info(train graph))
    print(nx.info(test_graph))
                                                                                                   •
Name:
Type: DiGraph
Number of nodes: 1780722
Number of edges: 7550015
Average in degree: 4.2399
                     4.2399
Average out degree:
Name:
Type: DiGraph
Number of nodes: 1144623
Number of edges: 1887504
Average in degree: 1.6490
Average out degree: 1.6490
```

## 9.1 Addigng SVD Dot product as one of the feature

In [12]:

```
from tqdm import tqdm
svd_dot_u = []
svd_dot_v = []
for i in tqdm(range(len(df_final_train))):
    svd_s_u = list(df_final_train[df_final_train.columns[30:36]].iloc[i])
    svd_s_v = list(df_final_train[df_final_train.columns[42:48]].iloc[i])
    svd_d_u = list(df_final_train[df_final_train.columns[36:42]].iloc[i])
    svd_d_v = list(df_final_train[df_final_train.columns[48:]].iloc[i])
    svd_dot_u.append(np.dot(svd_s_u,svd_d_u))
    svd_dot_v.append(np.dot(svd_s_v,svd_d_v))
print(len(svd_dot_u), len(svd_dot_v))
100%| | 100002/1000002 [15:28<00:00, 107.67it/s]
```

100002 100002

```
In [13]:

df_final_train['svd_dot_u'] = svd_dot_u
df_final_train['svd_dot_v'] = svd_dot_v
df_final_train.head()
```

Out[13]:

	source_node	destination_node	indicator_link	jaccard_followers	jaccard_followees	cosine_followers	cosine_followees	num_followe
0	273084	1505602	1	0	0.000000	0.000000	0.000000	
1	832016	1543415	1	0	0.187135	0.028382	0.343828	
2	1325247	760242	1	0	0.369565	0.156957	0.566038	
3	1368400	1006992	1	0	0.000000	0.000000	0.000000	

```
source_node destination_node indicator_link jaccard_followers jaccard_followees cosine_followers cosine_followees num_followe
4
         140165
                        1708748
                                                            0
                                                                       0.000000
                                                                                       0.000000
                                                                                                        0.000000
                                                                                                                          ▶
In [14]:
svd dot u = []
svd_dot_v = []
for i in tqdm(range(len(df_final_test))):
  svd_s_u = list(df_final_test[df_final_test.columns[30:36]].iloc[i])
  svd_s_v = list(df_final_test[df_final_test.columns[42:48]].iloc[i])
svd_d_u = list(df_final_test[df_final_test.columns[36:42]].iloc[i])
  svd d v = list(df final test[df final test.columns[48:]].iloc[i])
  svd dot u.append(np.dot(svd s u,svd d u))
  svd_dot_v.append(np.dot(svd_s_v,svd_d_v))
df_final_test['svd_dot_u'] = svd_dot_u
df_final_test['svd_dot_v'] = svd_dot_v
df_final_test.head()
           | 50002/50002 [06:04<00:00, 137.11it/s]
Out[14]:
```

	source_node	destination_node	indicator_link	jaccard_followers	jaccard_followees	cosine_followers	cosine_followees	num_followe
0	848424	784690	1	0	0.0	0.029161	0.000000	
1	483294	1255532	1	0	0.0	0.000000	0.000000	
2	626190	1729265	1	0	0.0	0.000000	0.000000	
3	947219	425228	1	0	0.0	0.000000	0.000000	
4	991374	975044	1	0	0.2	0.042767	0.347833	
4								Þ

# 9.2 Adding Preferential Attachment as one feature

```
In [0]:
```

```
def compute features(df final, graph):
    #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=[]
    for i,row in df final.iterrows():
            s1=set(graph.predecessors(row['source_node']))
            s2=set(graph.successors(row['source node']))
        except:
            s1 = set()
            s2 = set()
            d1=set(graph.predecessors(row['destination node']))
            d2=set(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))
```

```
return num_followers_s, num_followers_d, num_followees_s, num_followees_d
In [0]:
followers_s, followers_d, followees_s, followees_d = compute_features(df_final_train, train_graph)
In [17]:
df_final_train['pref_attachment_followers'] = [a*b for a,b in zip(followers_s,followers_d)]
df_final_train['pref_attachment_followees'] = [a*b for a,b in zip(followees_s,followees_d)]
df_final_train.head()
Out[17]:
   source_node destination_node indicator_link jaccard_followers jaccard_followees cosine_followers cosine_followees num_followe
0
        273084
                      1505602
                                                               0.000000
                                                                              0.000000
                                                                                             0.000000
        832016
                      1543415
                                                      0
                                                               0.187135
                                                                              0.028382
                                                                                             0.343828
1
                                       1
```

#### In [18]:

2

3

1325247

1368400

140165

760242

1006992

1708748

followers\_s, followers\_d, followees\_s, followees\_d = compute\_features(df\_final\_test, test\_graph)
df\_final\_test['pref\_attachment\_followers'] = [a\*b for a,b in zip(followers\_s,followers\_d)]
df\_final\_test['pref\_attachment\_followees'] = [a\*b for a,b in zip(followees\_s,followees\_d)]
df\_final\_test.head()

0

0.369565

0.000000

0.000000

0.156957

0.000000

0.000000

0.566038

0.000000

0.000000

Out[18]:

	source_node	destination_node	indicator_link	jaccard_followers	jaccard_followees	cosine_followers	cosine_followees	num_followe
0	848424	784690	1	0	0.0	0.029161	0.000000	
1	483294	1255532	1	0	0.0	0.000000	0.000000	
2	626190	1729265	1	0	0.0	0.000000	0.000000	
3	947219	425228	1	0	0.0	0.000000	0.000000	
4	991374	975044	1	0	0.2	0.042767	0.347833	
4								Þ

#### In [0]:

```
df_final_train.drop(['source_node', 'destination_node','indicator_link'],axis=1,inplace=True)
df_final_test.drop(['source_node', 'destination_node','indicator_link'],axis=1,inplace=True)
```

# 9.3 Applying XGBoost

```
from xgboost import XGBClassifier
from sklearn.metrics import accuracy_score
from sklearn.model_selection import RandomizedSearchCV
from datetime import datetime
```

#### In [21]:

```
estimators = [10,50,100,250,450]
train error = []
test error = []
for i in estimators:
   model = XGBClassifier(max depth=10,
                            subsample=0.33,
                            n estimators=i,
                            learning rate = 0.01)
    model.fit(df_final_train,y_train)
    train_er = 1 - accuracy_score(y_train, model.predict(df_final_train))
    test_er = 1 - accuracy_score(y_test,model.predict(df_final_test))
    test_error.append(test_er)
    train error.append(train er)
    print('Estimators = ',i,'Train Error',train_er,'test Error',test_er)
plt.plot(estimators,train_error,label='Train Error')
plt.plot(estimators, test error, label='Test Error')
plt.xlabel('Estimators')
plt.ylabel('Error')
plt.title('Estimators vs Error at depth of 10')
```

```
Estimators = 10 Train Error 0.024259514809703786 test Error 0.06373745050197988 

Estimators = 50 Train Error 0.023449531009379854 test Error 0.06259749610015597 

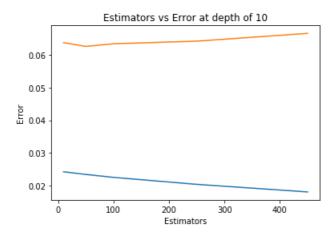
Estimators = 100 Train Error 0.022549549009019842 test Error 0.06339746410143599 

Estimators = 250 Train Error 0.020419591608167864 test Error 0.06419743210271589 

Estimators = 450 Train Error 0.018099638007239904 test Error 0.06657733690652379
```

#### Out[21]:

Text(0.5, 1.0, 'Estimators vs Error at depth of 10')



#### In [22]:

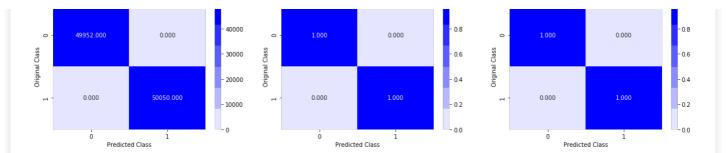
```
depths = [3,9,11,15,20,35,50,70,130]
train error = []
test error = []
for i in depths:
   model = XGBClassifier(max depth=i,
                            subsample=0.33,
                            n estimators=50,
                            learning rate = 0.01)
    model.fit(df final train,y train)
    train_er = 1 - accuracy_score(y_train, model.predict(df_final_train))
    test_er = 1 - accuracy_score(y_test,model.predict(df_final_test))
    test_error.append(test_er)
    train_error.append(train_er)
    print('Depth = ',i,'Train Error',train er,'test Error',test er)
plt.plot(depths,train_error,label='Train Error')
plt.plot(depths,test error,label='Test Error')
plt.xlabel('Depths')
```

```
plt.ylabel('Error')
plt.title('Depths vs Error at 50 Estimators')
Depth = 3 Train Error 0.07985840283194334 test Error 0.0836766529338826
Depth = 9 Train Error 0.026239475210495744 test Error 0.06423743050277986
Depth = 11 Train Error 0.022119557608847873 test Error 0.06297748090076394
Depth = 15 Train Error 0.01885962280754383 test Error 0.06301747930082802
Depth =
         20 Train Error 0.01857962840743188 test Error 0.0632974681012759
         35 Train Error 0.018549629007419854 test Error 0.0632974681012759
Depth =
         50 Train Error 0.018549629007419854 test Error 0.0632974681012759
Depth = 70 Train Error 0.018549629007419854 test Error 0.0632974681012759
Depth = 130 Train Error 0.018549629007419854 test Error 0.0632974681012759
              Depths vs Error at 50 Estimators
  0.08
  0.07
  0.06
Ē 0.05
  0.04
  0.03
  0.02
            20
                  40
                        60
                                   100
                                         120
                             80
                        Depths
In [0]:
params = {
        'n_estimators': [5, 10, 50, 100, 200, 500],
        'gamma': [0.5, 1, 1.5, 2, 5],
        'colsample_bytree': [0.6, 0.8, 1.0],
        'learning_rate': [0.01,0.1,0.3,0.5,1],
        'max depth': [2, 5, 9, 10, 11, 13, 15, 20]
#https://www.kaggle.com/tilii7/hyperparameter-grid-search-with-xgboost
def timer(start time=None):
    if not start time:
        start_time = datetime.now()
       return start_time
    elif start time:
        thour, temp sec = divmod((datetime.now() - start time).total seconds(), 3600)
        tmin, tsec = divmod(temp_sec, 60)
        print('\n Time taken: %i hours %i minutes and %s seconds.' % (thour, tmin, round(tsec, 2)))
4
In [25]:
xqb = XGBClassifier()
clf = RandomizedSearchCV(xgb, params, cv=5, scoring='f1')
start_time = timer(None) # timing starts from this point for "start_time" variable
clf.fit(df_final_train, y_train)
timer(start time) # timing ends here for "start time" variable
Time taken: 1 hours 11 minutes and 39.79 seconds.
In [26]:
clf.best params
Out [26]:
```

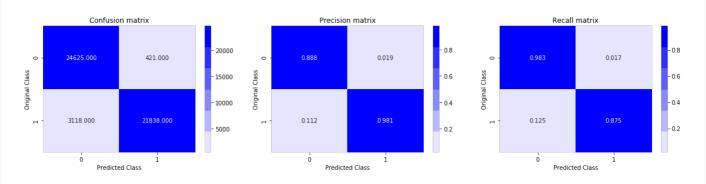
{'colsample bytree': 0.6,

```
'gamma': ⊥,
 'learning_rate': 0.1,
 'max depth': 9,
 'n estimators': 500}
In [0]:
clf = XGBClassifier(learning rate=0.1, colsample bytree=0.6, max depth=9, gamma=1, n estimators=500
In [0]:
clf.fit(df final train,y train)
y train pred = clf.predict(df final train)
y test pred = clf.predict(df final test)
In [29]:
from sklearn.metrics import f1 score
print('Train f1 score', f1_score(y_train, y_train_pred))
print('Test f1 score',f1_score(y_test,y_test_pred))
Train f1 score 1.0
Test f1 score 0.9250450068834057
In [0]:
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, yticklabels=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 [31]:
print('Train confusion matrix')
plot confusion_matrix(y_train,y_train_pred)
print('Test confusion matrix')
plot_confusion_matrix(y_test,y_test_pred)
Train confusion_matrix
```

Confusion matrix Precision matrix Recall matrix -10

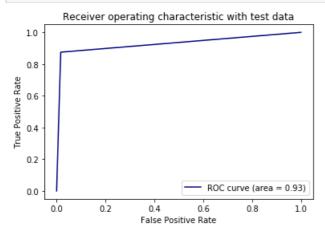


Test confusion\_matrix



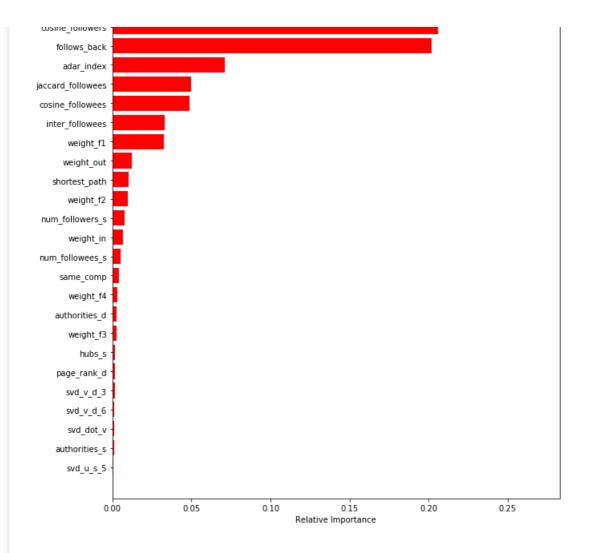
#### In [32]:

```
from sklearn.metrics import roc_curve, auc
fpr,tpr,ths = roc_curve(y_test,y_test_pred)
auc_sc = auc(fpr, tpr)
plt.plot(fpr, tpr, color='navy',label='ROC curve (area = %0.2f)' % auc_sc)
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic with test data')
plt.legend()
plt.show()
```



#### In [33]:

```
features = df_final_train.columns
importances = clf.feature_importances_
indices = (np.argsort(importances))[-25:]
plt.figure(figsize=(10,12))
plt.title('Feature Importances')
plt.barh(range(len(indices)), importances[indices], color='r', align='center')
plt.yticks(range(len(indices)), [features[i] for i in indices])
plt.xlabel('Relative Importance')
plt.show()
```



## 10 Conclusion

```
In [0]:
```

```
from prettytable import PrettyTable
x = PrettyTable()
x.field_names = ["Algorithm", "Hyper Parameters", "F1 Score-Train", "F1 Score-Test", "ROC-AUC Score
(test)"]
```

#### In [35]:

#### Steps that were followed during this case study

- 1. The data was analysed for the initial steps and the actual data had to be prepared to apply models on top of that.

- $2. \ \ \text{ I he need was to have certain features to help train the model and so proper featurization was done.}$
- 3. Features were then extracted from the data available and a numerous type of featurizations were applied which eventually gave 47 Features at the end
- 4. These features are then used for the purpose of modelling.
- 5. 2 models have been applied after hyperparameter tuning which are XGBoost and Random Forest and the results seen are mentioned above.