

# NGSA: Network Science Analytics

## Assignment 1

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### 1 Graph Theory and Graph Properties

- **Question 1**

*Let  $\mathbf{A}$  be the adjacency matrix of an undirected graph (unweighted, with no self-loops) and  $\mathbf{1}$  be the column vector whose elements are all 1. In terms of these quantities and simple matrix operations like matrix transpose and matrix trace, write expressions for:*

- (a) *The vector  $\mathbf{k}$  whose elements are the degrees  $k_i$  of the nodes.*

$$\mathbf{k} = \mathbf{A} * \mathbf{1}$$

- (b) *The number  $m$  of edges in the graph.*

$$m = \frac{\text{sum}(\mathbf{A} * \mathbf{1})}{2} = \frac{\mathbf{1}^T * \mathbf{A} * \mathbf{1}}{2}$$

- (c) *The matrix  $\mathbf{N}$  whose element  $N_{ij}$  is equal to the number of common neighbors of nodes  $i$  and  $j$ .*

$$\mathbf{N} = \mathbf{A}^2$$

- **Question 2**

*Consider a bipartite network, with its two types of nodes (type 1 and 2), and suppose that there are  $n_1$  nodes of type 1 and  $n_2$  nodes of type 2. Show that the mean degrees  $c_1$  and  $c_2$  of the two types are related by*

$$c_2 = \frac{n_1}{n_2} * c_1$$

If we consider a bipartite network, then each edge has one vertex in each type of nodes. Let us call  $\mathbf{N}$  the total number of edges in the graph. There are  $\mathbf{N}$  edges connected to both types of nodes.

By definition of the mean degrees  $c_1$  and  $c_2$  we have:

$$c_1 = \frac{N}{n_1} \text{ and } c_2 = \frac{N}{n_2}$$

From that we can derive:

$$n_1 * c_1 = \mathbf{N} = n_2 * c_2$$

i.e.

$$c_2 = \frac{n_1}{n_2} * c_1$$

• **Question 3**

Let  $G = (V, E)$  be an unweighted, undirected graph with no self-loops.

The  $(i, j)$ -th element of  $\mathbf{A}^l$  (i.e., the adjacency matrix raised to the  $l$ -th power) counts the number of paths of length  $l$  that start from node  $i$  and end at node  $j$ .

A triangle in a graph corresponds to a clique of three nodes.

- (a) Using simple matrix operations, express the total number of triangles in the graph  $\Delta(G)$ , as a function of the adjacency matrix  $\mathbf{A}$ .

We simply take the sum over all the diagonal elements of  $\mathbf{A}^3$ , but we remember to divide by 6 in order to not count the same triangle multiple times.

The number 6 comes from the 3 choices of starting points times the 2 possible direction of going along the triangle.

$$\Delta(G) = \frac{\text{Tr}(\mathbf{A}^3)}{6}$$

Note: if  $\mathbf{S}$  is a symmetric matrix composed of 0s and 1s then  $\text{Tr}(\mathbf{S}^2) = \text{sum}(\mathbf{S})$

- (b) Similarly, express the total number of triangles in the graph  $\Delta(G)$  as a function of the eigenvalues  $\lambda_i, \forall i \in V$  of  $\mathbf{A}$ .

We know that for a given matrix  $\mathbf{A}$  and its eigenvalues  $(\lambda_i)_i$ ,  $\text{Tr}(\mathbf{A}) = \sum \lambda_i$ .

We also know that if  $\lambda$  is an eigenvalue of  $\mathbf{A}$  then  $\lambda^n$  is an eigenvalue of  $\mathbf{A}^n$ .

We can derive:

$$\Delta(G) = \frac{\sum \lambda_i^3}{6}$$

- (c) Let  $\Delta_i, \forall i \in V$  be the number of triangles that node  $i$  participates in. Express  $\Delta_i$  as a function of the spectrum (i.e., eigenvalues and/or eigenvectors) of the adjacency matrix  $\mathbf{A}$ .

The number of triangles that  $i$  participates in is the  $i$ -th diagonal component of  $\mathbf{A}^3$  divided by 2 (2 possible directions to form triangle from a 3-way path from  $i$  to  $i$ ).

If we define  $\mathbf{1}_i$  the the vector containing only 0s and a 1 in the  $i$ -th position, we have:

$$\Delta_i = \frac{\mathbf{1}_i^T * \mathbf{A}^3 * \mathbf{1}_i}{2}$$

$\mathbf{A}$  is a symmetric matrix so we can diagonalize it (Spectral Theorem) as follows, where  $\mathbf{P}$  is an orthogonal matrix of eigenvectors:

$$\mathbf{A} = \mathbf{P}^{-1} * \mathbf{D} * \mathbf{P} = \mathbf{P}^T * \mathbf{D} * \mathbf{P}$$

thus

$$\Delta_i = \frac{\mathbf{1}_i^T * (\mathbf{P}^T * \mathbf{D} * \mathbf{P})^3 * \mathbf{1}_i}{2} = \frac{(\mathbf{P} * \mathbf{1}_i)^T * \mathbf{D}^3 * (\mathbf{P} * \mathbf{1}_i)}{2} = \frac{\mathbf{V}_i^T * \mathbf{D}^3 * \mathbf{V}_i}{2}$$

where  $\mathbf{D}$  is the diagonal matrix of eigenvalues and  $\mathbf{V}_i$  is the  $i$ -th eigenvector.

## 2 Graph Models

### • Question 4

Consider the random graph  $G_{n,p}$  with average degree  $c$ .

- (a) Show that in the limit of large  $n$ , the expected number of triangles in the graph is  $\frac{1}{6}c^3$ . In other words, show that the number of triangles is constant, neither growing nor vanishing in the limit of large  $n$ .

We have seen in the class Lecture 2A that for a random graph and large  $n$ :

- For a given node there are on average  $c^2$  nodes at distance 2. (Slide 34)
- The probability of existing for a given edge is  $p = \frac{c}{n-1} \approx \frac{c}{n}$ . (Slide 27)

The expected number of triangles is then:

$$\frac{\text{nodes in the graph} \times \text{nodes at distance 2} \times P(\text{last edge})}{3 \text{ vertices} \times 2 \text{ directions}} = \frac{n \times c^2 \times \frac{c}{n}}{6} = \frac{c^3}{6}$$

- (b) A connected triplet is defined as a triplet of nodes  $uvw$ , with edges  $(u,v)$  and  $(v,w)$  (the edge  $(u,w)$  can be present or not). Show that the expected number of connected triplets in the graph is  $\frac{1}{2}n \times c^2$ .

Similarly we obtain the expected number of triplets:

$$\frac{\text{nodes in the graph} \times \text{nodes at distance 2}}{\text{number of possible end vertices}} = \frac{n \times c^2}{2}$$

- (c) The clustering coefficient of a graph can also be expressed as

$$C = \frac{(\text{number of triangles}) \times 3}{(\text{number of connected triplets})}$$

Calculate the clustering coefficient of the  $G_{n,p}$  random graph using the above formula based on (a) and (b), and confirm that for large  $n$  it agrees with the value shown in class (Lecture 2A; slide number 32).

From the previous questions we can derive:

$$C = \frac{\frac{c^3}{6} \times 3}{\frac{n \times c^2}{2}} = \frac{c}{n}$$

which is the expected result from the Lecture 2A.

### 3 Centrality Criteria

- **Question 5**

Suppose that we define a new centrality criterion  $x_i, \forall i \in V$  to be a sum of contributions as follows: 1 for node  $i$  itself,  $\alpha$  for each node at (geodesic) distance 1 from  $i$ ,  $\alpha^2$  for each node at distance 2, and so forth, where  $\alpha < 1$  is a given constant.

- (a) Write an expression for  $x_i$  in terms of  $\alpha$  and the geodesic distances  $d_{ij}$  between node pairs.

$$x_i = \sum_{j \in V} \alpha^{d_{ij}}$$

- (b) Describe briefly (max 3 lines) an algorithm for computing this centrality measure. What is the complexity of calculating  $x_i$  for all  $i \in V$ ?

We could proceed as follows:

- Compute the pairwise shortest path for all  $(i, j)$  and store it in a matrix  $\mathbf{M}$ .  
This could be achieved with the Dijkstra algorithm;
- For each column in  $\mathbf{M}$  compute  $x_i$  as defined above.

As seen in Lecture 3B Slide 14, the complexity would be that of Dijkstra's algorithm i.e.  $O(n^2 \log n + nm)$ .

• **Question 6**

Consider an undirected, unweighted graph of  $n$  nodes that is composed by exactly two sub-graphs of size  $n_A$  and  $n_B$ , which are connected by a single edge  $(A, B)$ . Show that the closeness centralities  $C_A$  and  $C_B$  of nodes  $A$  and  $B$  respectively, are related by

$$\frac{1}{C_A} + \frac{n_A}{n} = \frac{1}{C_B} + \frac{n_B}{n}$$

Let  $G_A$  and  $G_B$  be the subgraphs of size  $n_A$  and  $n_B$ . By definition, we have:

$$\frac{1}{C_A} = \frac{\sum_{j \in G} d_{Aj}}{n} = \frac{\sum_{j \in G_A} d_{Aj} + \sum_{j \in G_B} d_{Aj}}{n} = \frac{\sum_{j \in G_A} d_{Aj}}{n} + \frac{\sum_{j \in G_B} (1 + d_{Bj})}{n}$$

i.e.

$$\frac{1}{C_A} - \frac{n_B}{n} = \frac{\sum_{j \in G_A} d_{Aj}}{n} + \frac{\sum_{j \in G_B} (d_{Bj})}{n}$$

And we can similarly derive:

$$\frac{1}{C_B} - \frac{n_A}{n} = \frac{\sum_{j \in G_A} d_{Aj}}{n} + \frac{\sum_{j \in G_B} (d_{Bj})}{n}$$

Thus we have:

$$\frac{1}{C_A} - \frac{n_B}{n} = \frac{1}{C_B} - \frac{n_A}{n}$$

i.e.

$$\frac{1}{C_A} + \frac{n_A}{n} = \frac{1}{C_B} + \frac{n_B}{n}$$

## 4 Analyzing a Real Network

### • Question 7

(a) Basic properties of the network:

1. Number of nodes: 5 242; number of edges: 14 496.
2. Number of CCs: 355. The distribution of the CCs' sizes can be seen in Figure 1.

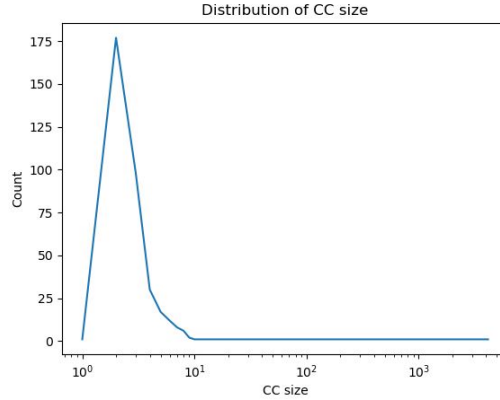


Figure 1: Size of connected components.

3. GCC: Number of nodes in the GCC: 4 158 (79.3% of total).  
Number of edges: 13 428(92.6% of total).

We can see that almost all the edges are concentrated within the GCC, more so than the nodes (13.3% difference). We can see that there are a high number of 2-connected components from Figure 1: those count as 1 edge but 2 nodes, which explains the difference in percentages.

(b) Degree distribution:

- Minimum degree: 1;
- Maximum degree: 81;
- Median degree: 3;
- Mean degree: 5.5.

We can see that the median degree is quite smaller than the mean degree, which is understandable given the fact that the degree distribution is right-skewed.

We can see that there are nodes that are alone and that the most-connected node has only 81 connections out of 5 242. This is understandable since no scientist will have collaborated with a high percentage of the community.

Using the `powerlaw` Python library I obtained a power law with coefficient  $\alpha = 2.09$ . The degree distribution is plotted in Figure 2.

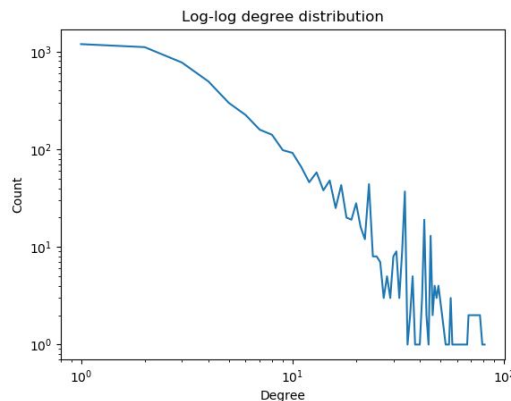


Figure 2: Degree distribution.

(c) *Triangles*

There are 47 779 triangles in the network.

The triangle participation is plotted in Figure 3.

It is interesting to see that there are 3 times more nodes that participate in 3 triangles than nodes that participate in 2 triangles (probably because if you participate in 2 separate triangles there is a high chance of participating in a 3rd composed of vertices of the first 2).

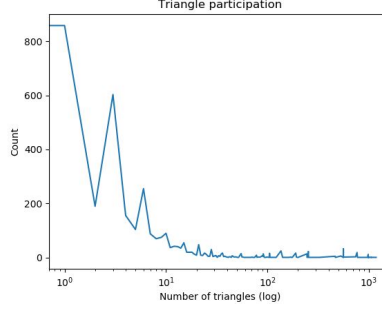


Figure 3: Triangle participation.

(d) *Spectral counting of triangles*

Finding eigenvalues involves inverting matrices and here we are manipulating quite large matrices ( $\approx 5\,000 \times 5\,000$ ).

By looking at the eigenvalue distribution in Figure 4, we can see that the major contributions to  $\frac{\sum_i \lambda_i^3}{6}$  will be contained in the first eigenvalues. Additionally, the eigenvalues tend to compensate each other starting around the 300-th eigenvalue.

We plot the respective errors and the computation times associated to various values of  $k$  in Figure 5.  $k = 500$  seems like a good value.

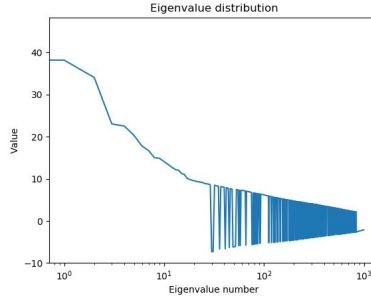


Figure 4: The eigenvalues of the adjacency matrix.

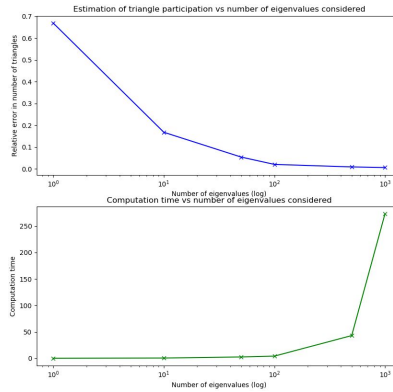


Figure 5: Triangle participation.



## Code for question 7

```

import networkx as nx
import matplotlib.pyplot as plt
import numpy as np
import scipy
import powerlaw
from datetime import datetime

questions = [1,2,3]

G = nx.read_edgelist("CA-GrQc.txt")
connected_components = sorted(nx.connected_component_subgraphs(G),
                               key = len, reverse=True)
gcc = connected_components[0]
if 1 in questions:
    print("Number_of_nodes: %i" % G.number_of_nodes())
    print("Number_of_edges: %i" % G.number_of_edges())

    print("\nNumber_of_CC: %i" % nx.number_connected_components(G))

    cc_size = [g.number_of_nodes() for g in connected_components]
    values = sorted(np.unique(cc_size))
    counts = [cc_size.count(val) for val in values]

    please_plot = True
    if please_plot:
        plt.plot(values, counts)
        ax = plt.gca()
        ax.set_xscale('log')
        plt.title("Distribution_of_CC_size")
        plt.xlabel("CC_size")
        plt.ylabel("Count")
        plt.show()

    print("Number_of_nodes_in_GCC: %i i.e. %.1f%% of total_graph"
          % (gcc.number_of_nodes(),
             100*float(gcc.number_of_nodes())/float(G.number_of_nodes())))
    print("Number_of_edges_in_GCC: %i i.e. %.1f%% of total_graph"
          % (gcc.number_of_edges(),
             100*float(gcc.number_of_edges())/float(G.number_of_edges())))
elif 2 in questions:
    nodes = G.nodes
    degrees = [d for n, d in nx.degree(G)]
    print("Min_degree: %i" % np.min(degrees))
    print("Max_degree: %i" % np.max(degrees))
    print("Mean_degree: %.1f" % np.mean(degrees))
    print("Median_degree: %i" % np.median(degrees))

    values = sorted(np.unique(degrees))
    counts = [degrees.count(val) for val in values]
    fit = powerlaw.Fit(degrees, discrete=True)
    print('Power_law!_alpha= ', fit.power_law.alpha, ' _sigma= ', fit.power_law.sigma)
    please_plot = True
    if please_plot:
        plt.plot(values, counts)
        ax = plt.gca()
        ax.set_xscale('log')
        ax.set_yscale('log')
        plt.title("Log-log_degree_distribution")
        plt.xlabel("Degree")
        plt.ylabel("Count")
        plt.show()
elif 3 in questions:
    to_remove = []
    for edge in gcc.edges:
        if edge[0] == edge[1]:
            to_remove += [edge[0]]
    for n in to_remove:
        gcc.remove_edge(n, n)
    A = np.matrix(nx.to_numpy_matrix(gcc))
    B = A ** 3
    num_triangles = int(np.trace(B)/6)
    print("Total_number_of_triangles: %i" % num_triangles)

```

```

triangle_participation = [B[i, i]/2 for i in range(len(B))]
values = sorted(np.unique(triangle_participation))
counts = [triangle_participation.count(val) for val in values]

please_plot = False
if please_plot:
    plt.plot(values, counts)
    ax = plt.gca()
    ax.set_xscale('log')
    plt.title("Triangle_participation")
    plt.xlabel("Number_of_triangles_(log)")
    plt.ylabel("Count")
    plt.show()

eigenvalues, eigenvectors = np.array(scipy.sparse.linalg.eigs(A, 1000))
eigenvalues = eigenvalues.real
plt.plot(eigenvalues)
ax = plt.gca()
ax.set_xscale('log')
plt.title("Eigenvalue_distribution")
plt.xlabel("Eigenvalue_number")
plt.ylabel("Value")
plt.show()

ready_for_long_computation = False
if ready_for_long_computation:
    num_eig = [1, 10, 50, 100, 500, 1000]
    errors = []
    delays = []
    for k in num_eig:
        start = datetime.now()
        eigenvalues, eigenvectors = np.array(scipy.sparse.linalg.eigs(A, k))
        eigenvalues = eigenvalues.real
        errors += [abs((np.sum(np.power(eigenvalues, 3)) / 6) - num_triangles)
                    / num_triangles]
        time_computation = datetime.now() - start
        delays += [time_computation.minutes
                    + time_computation.seconds
                    + time_computation.microseconds / 1000000]

    plt.subplot(211)
    plt.plot(num_eig, errors, '-bx')
    ax = plt.gca()
    ax.set_xscale('log')
    plt.title("Estimation_of_triangle_participation_vs_number_of_eigenvalues")
    plt.xlabel("Number_of_eigenvalues_(log)")
    plt.ylabel("Relative_error_in_number_of_triangles")

    plt.subplot(212)
    plt.plot(num_eig, delays, '-gx')
    ax = plt.gca()
    ax.set_xscale('log')
    plt.title("Computation_time_vs_number_of_eigenvalues_considered")
    plt.xlabel("Number_of_eigenvalues_(log)")
    plt.ylabel("Computation_time")
    plt.show()

```

• Question 8

- (a) In Lecture 2A (Slide 27) we saw that  $p = \frac{c}{n-1}$  so the mean degree should be  $\approx 8.9$ .
- (b) We saw in Lecture 2A (Slide 45) that the graph is connected if  $c > \ln(n)$ . Here  $\ln(1000) = 6.9$  so the graph is connected.
- (c) The computed mean degree of the graph is 8.93. The degree distribution is plotted in Figure 6 and does look like a Poisson distribution with parameter  $c$  (see Figure 7 which was computed from <https://homepage.divms.uiowa.edu/~mbognar/applets/pois.html>).

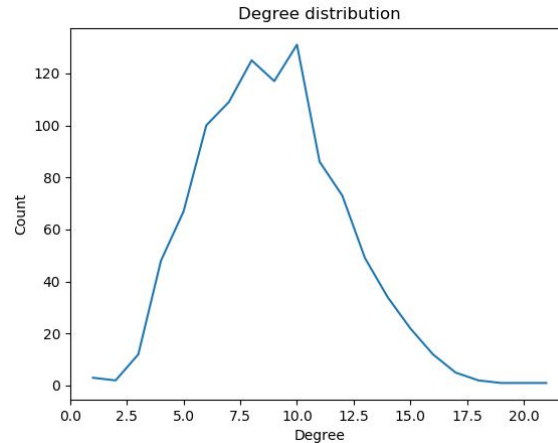


Figure 6: Degree distribution.

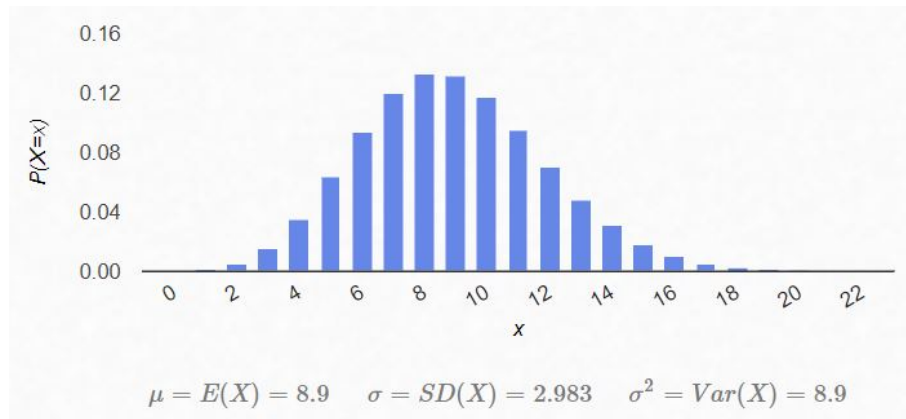


Figure 7: Poisson distribution of parameter  $c = 8.9$ .

### Code for question 8

```
import networkx as nx
import matplotlib.pyplot as plt
import numpy as np

n = 1000
p = 0.009
G = nx.fast_gnp_random_graph(n, p)
degrees = [d for n, d in nx.degree(G)]

print("Mean_degree: %.2f" % np.mean(degrees))

values = sorted(np.unique(degrees))
counts = [degrees.count(val) for val in values]

please_plot = True
if please_plot:
    plt.plot(values, counts)
    ax = plt.gca()
    plt.title("Degree_distribution")
    plt.xlabel("Degree")
    plt.ylabel("Count")
    plt.show()
```

• **Question 9**

While building the Kronecker graph, I chose to take  $k = 13$  and end up with up to 8 192 nodes potentially. Naturally, some of these nodes will end up not existing (if all their edges are picked as 0): I ended up with 6 440 nodes whereas the real network has 5 242 nodes<sup>1</sup>.

- (a) The produced Kronecker graph is not connected because  $0.26 + 0.53 < 1$  and the GCC is not of size  $\Theta(n)$  because  $(0.99 + 0.26)(0.26 + 0.53) = 0.9875 < 1$ . Those rules are extracted from Lecture 3A Slide 44 but we can for example see that if  $b + c < 1$  then the degree probability of the last node (last row of the matrix) is  $(b + c)^k \rightarrow 0$ .

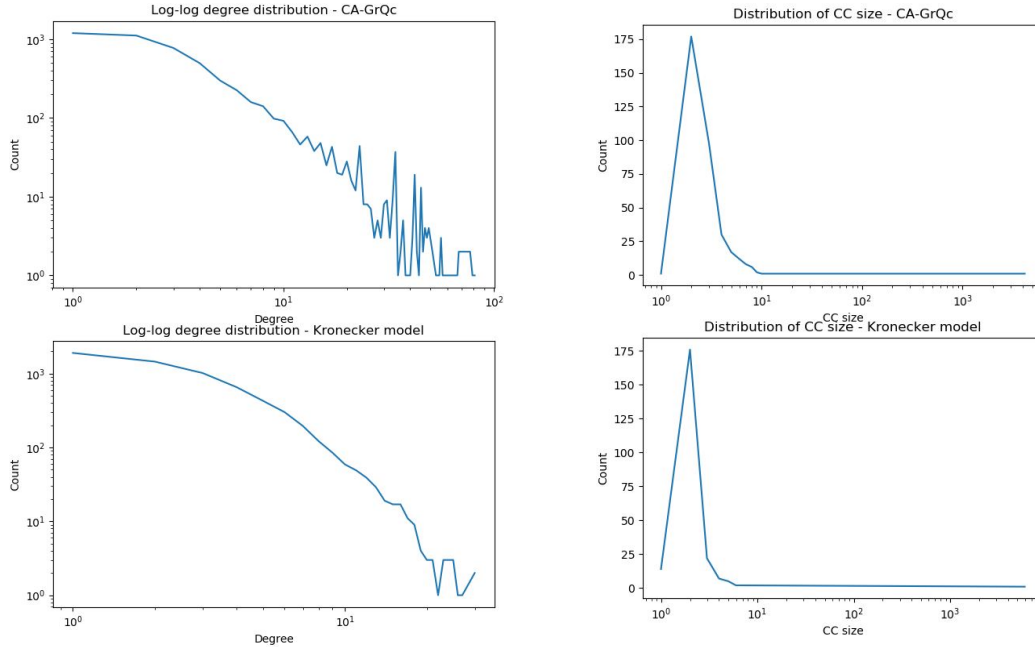
*Note: see [Mahdian and Xu '07](#) for more complete proof, section 2.2 and 2.3.*

- (b) We plot below in Figure 8 the degree distribution as well as the size of the connected components.

We can see that the degree distribution isn't as granular but has a very similar shape.

The distribution of the connected components is very close as well.

Lastly, the size of the connected components of size 2 is the same (value of the peak of Figure 8b), while the share of the GCC is similar (97.3% of the edges in the Kronecker model vs 92.6% in the real network).



(a) Degree distribution

(b) Size of connected components.

Figure 8: Comparing the Kronecker model with the real CA-GrQc graph.

<sup>1</sup>I could have added the nodes with 0 edges as nodes of degree 0 but decided against it, given that the original network doesn't have any such node. Indeed, the original graph is constructed solely on edges.

### Code for question 9

```

import numpy as np
import networkx as nx
import matplotlib.pyplot as plt
from datetime import datetime
from tqdm import tqdm

def generate_graph(M):
    g = nx.Graph()
    n = len(M)
    for l in tqdm(range(n)):
        for j in range(n - l):
            if np.random.choice([False, True], p=[1 - M[l, j], M[l, j]]):
                g.add_edge(l, j)
    return g.to_undirected()

A_1 = np.array([[0.99, 0.26], [0.26, 0.53]])

# k = 11 => 4096 nodes
k = 12

start = datetime.now()
A = A_1
for i in range(k):
    A = np.kron(A, A_1)
time_computation = datetime.now() - start
print("Computation_time: %s" % time_computation)

start = datetime.now()
k_G = generate_graph(A)
time_computation = datetime.now() - start
print("Reading_time: %s" % time_computation)

connected_components_k_G = sorted(nx.connected_component_subgraphs(k_G),
                                   key = len, reverse=True)
gcc_k_G = connected_components_k_G[0]

print("Number_of_edges_in_GCC_(Kro._model): %i i.e. %1f%% of total_graph (%i edges)"
      % (gcc_k_G.number_of_edges(), 100 * float(gcc_k_G.number_of_edges())
        / float(k_G.number_of_edges()), k_G.number_of_edges()))

print("Number_of_nodes_in_GCC_(Kro._model): %i i.e. %1f%% of total_graph (%i nodes)"
      % (gcc_k_G.number_of_nodes(), 100 * float(gcc_k_G.number_of_nodes())
        / float(k_G.number_of_nodes()), k_G.number_of_nodes()))

degrees_k_G = [d for n, d in nx.degree(k_G)]
values_k_G = sorted(np.unique(degrees_k_G))
counts_k_G = [degrees_k_G.count(val) for val in values_k_G]

cc_size_k_G = [g.number_of_nodes() for g in connected_components_k_G]
values_cc_k_G = sorted(np.unique(cc_size_k_G))
counts_cc_k_G = [cc_size_k_G.count(val) for val in values_cc_k_G]

G = nx.read_edgelist("CA-GrQc.txt")
connected_components_G = sorted(nx.connected_component_subgraphs(G),
                                key = len, reverse=True)
gcc_G = connected_components_G[0]

degrees_G = [d for n, d in nx.degree(G)]
values_G = sorted(np.unique(degrees_G))
counts_G = [degrees_G.count(val) for val in values_G]

```

```

cc_size_G = [g.number_of_nodes() for g in connected_components_G]
values_cc_G = sorted(np.unique(cc_size_G))
counts_cc_G = [cc_size_G.count(val) for val in values_cc_G]

please_plot = True
if please_plot:
    plt.subplot(211)
    plt.plot(values_G, counts_G)
    ax = plt.gca()
    ax.set_xscale('log')
    ax.set_yscale('log')
    plt.title("Log-log_degree_distribution_-_CA-GrQc")
    plt.xlabel("Degree")
    plt.ylabel("Count")

    plt.subplot(212)
    plt.plot(values_k_G, counts_k_G)
    ax = plt.gca()
    ax.set_xscale('log')
    ax.set_yscale('log')
    plt.title("Log-log_degree_distribution_-_Kronecker_model")
    plt.xlabel("Degree")
    plt.ylabel("Count")
    plt.show()

    plt.subplot(211)
    plt.plot(values_cc_G, counts_cc_G)
    ax = plt.gca()
    ax.set_xscale('log')
    plt.title("Distribution_of_CC_size_-_CA-GrQc")
    plt.xlabel("CC_size")
    plt.ylabel("Count")

    plt.subplot(212)
    plt.plot(values_cc_k_G, counts_cc_k_G)
    ax = plt.gca()
    ax.set_xscale('log')
    plt.title("Distribution_of_CC_size_-_Kronecker_model")
    plt.xlabel("CC_size")
    plt.ylabel("Count")
    plt.show()

```

- **Question 10** My results are displayed on Figure 9.  
We can see that an attack has a bigger impact than a failure, in terms of splitting up the GCC into isolated components.

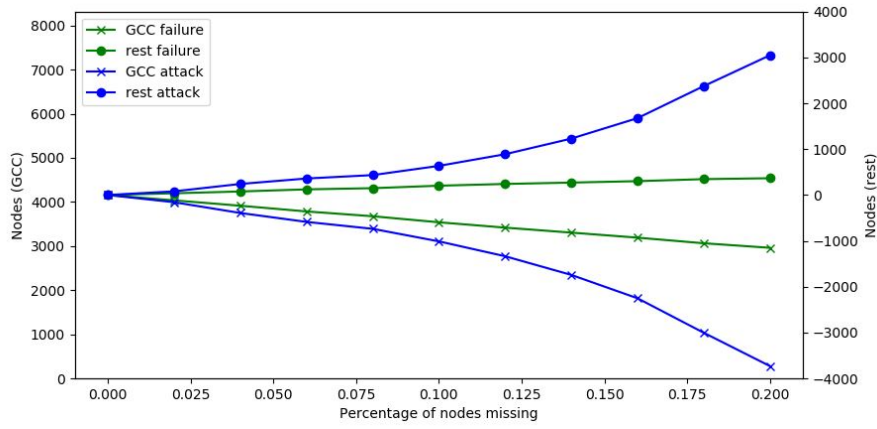


Figure 9: Comparing the impact of loss of nodes on the GCC and the rest of the nodes, in case of an attack and a failure scenario.



## Code for question 10

```

import numpy as np
import networkx as nx
import matplotlib.pyplot as plt
import random

G = nx.read_edgelist("CA-GrQc.txt")
G = max(nx.connected_component_subgraphs(G), key=len)

nodes = [n for n, d in nx.degree(G)]
degrees = [d for n, d in nx.degree(G)]

# sorts the nodes based on degrees
nodes = [n for _, n in sorted(zip(degrees, nodes), reverse=True)]
n = len(nodes)

# keep 20%
large_nodes = nodes[:int(0.2*n)]

G1 = G.copy()
G2 = G.copy()

starting_length = len(nodes)
current_length = starting_length
step = int(0.02 * starting_length)
GCCs_failure = []
rests_failure = []
totals = []
while current_length >= 0.8 * starting_length:
    gcc = max(nx.connected_component_subgraphs(G1), key=len)
    GCCs_failure += [len(gcc.nodes)]
    rests_failure += [len(nodes) - len(gcc.nodes)]
    totals += [len(nodes)]

    to_remove = random.sample(nodes, k=step)
    for node in to_remove:
        G1.remove_node(node)
    nodes = [n for n, d in nx.degree(G1)]
    current_length = len(nodes)

GCCs_attack = []
rests_attack = []
steps = []
current_steps = 0
current_length = starting_length
nodes = [n for n, d in nx.degree(G2)]
while len(large_nodes) > 0:
    gcc = max(nx.connected_component_subgraphs(G2), key=len)
    GCCs_attack += [len(gcc.nodes)]
    rests_attack += [len(nodes) - len(gcc.nodes)]
    totals += [len(nodes)]
    steps += [current_steps]
    current_steps += 0.02

    to_remove = random.sample(large_nodes, k=min(step, len(large_nodes)))
    for node in to_remove:
        G2.remove_node(node)
        large_nodes.remove(node)
    nodes = [n for n, d in nx.degree(G2)]
    current_length = len(nodes)

ax1 = plt.gca()
plt.ylabel("Nodes_(GCC)")
plt.xlabel("Percentage_of_nodes_missing")
ax2 = ax1.twinx()
plt.ylabel("Nodes_(rest)")
ax1.set_ylim([0, 2*GCCs_attack[0]])
ax2.set_ylim([-4000, 4000])
g_f, = ax1.plot(steps, GCCs_failure, '-xg', label='GCC_failure')
r_f, = ax2.plot(steps, rests_failure, '-og', label='rest_failure')
g_a, = ax1.plot(steps, GCCs_attack, '-xb', label='GCC_attack')
r_a, = ax2.plot(steps, rests_attack, '-ob', label='rest_attack')
plt.legend(handles=[g_f, r_f, g_a, r_a])
plt.show()

```