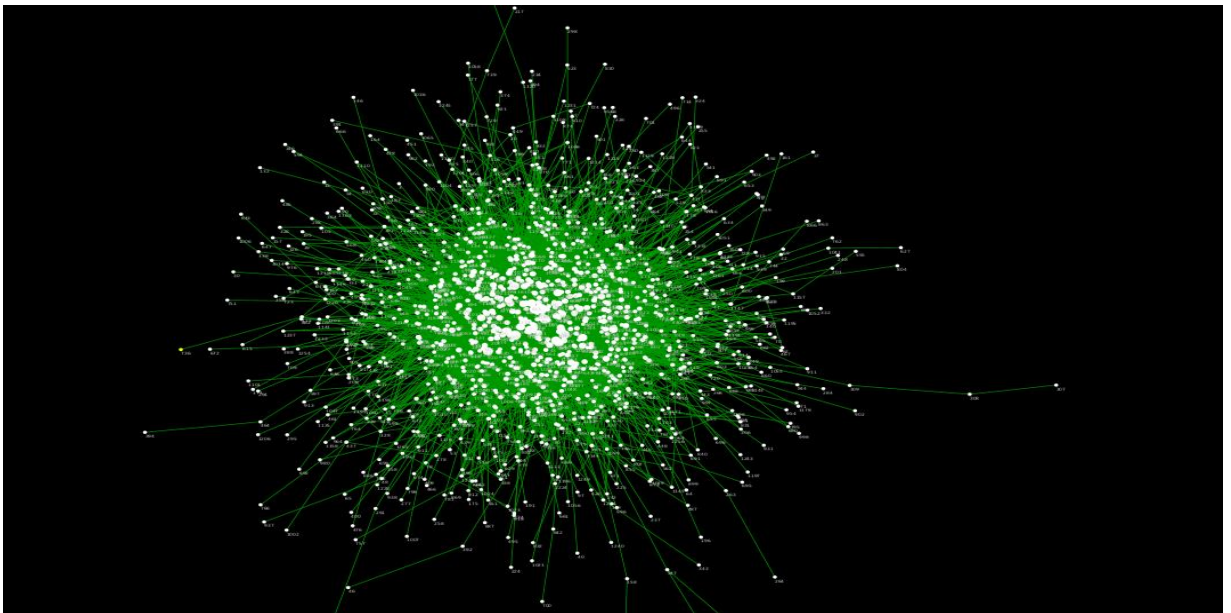
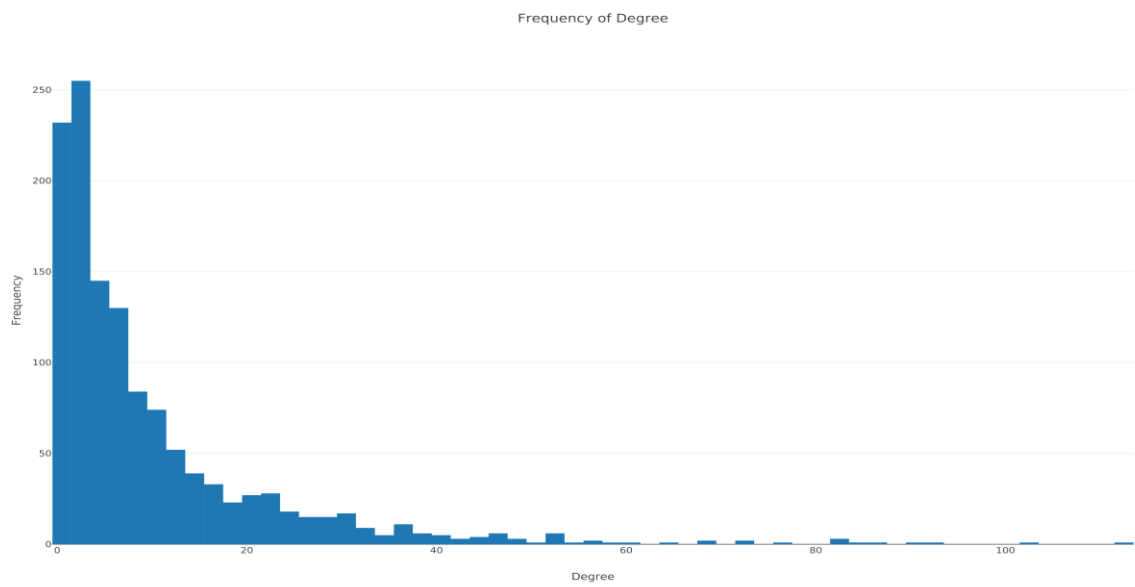


SNA Assignment 1 Solution

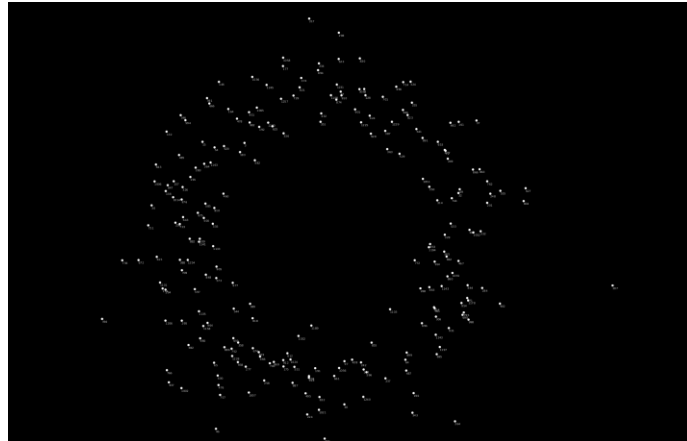
Tool Used - Cytoscape

Part B

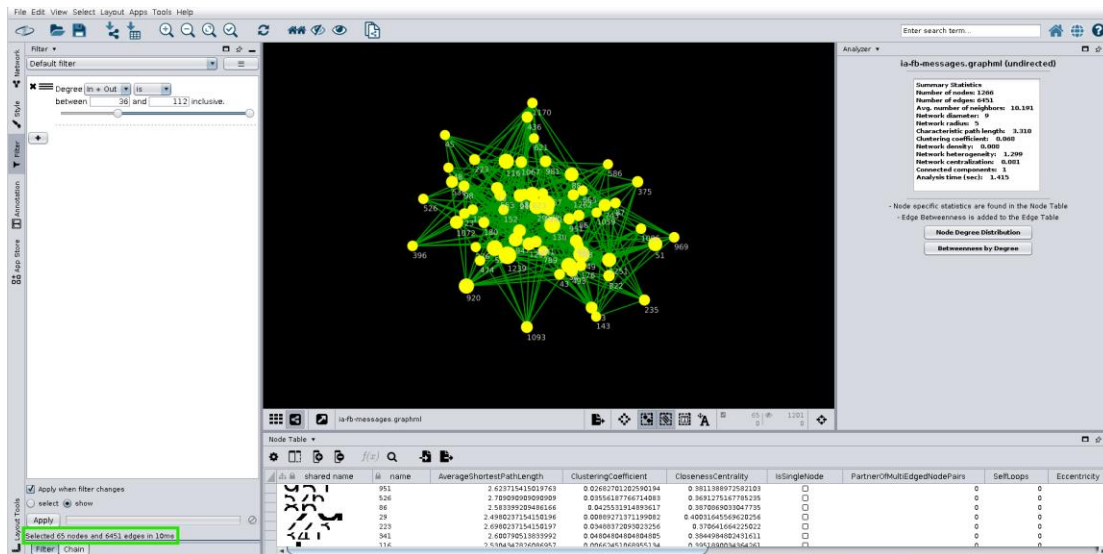
1. Degree Distribution of graph



2. Bottom 10% of nodes and the connection among them.



Top 5% of nodes and the connections among them.



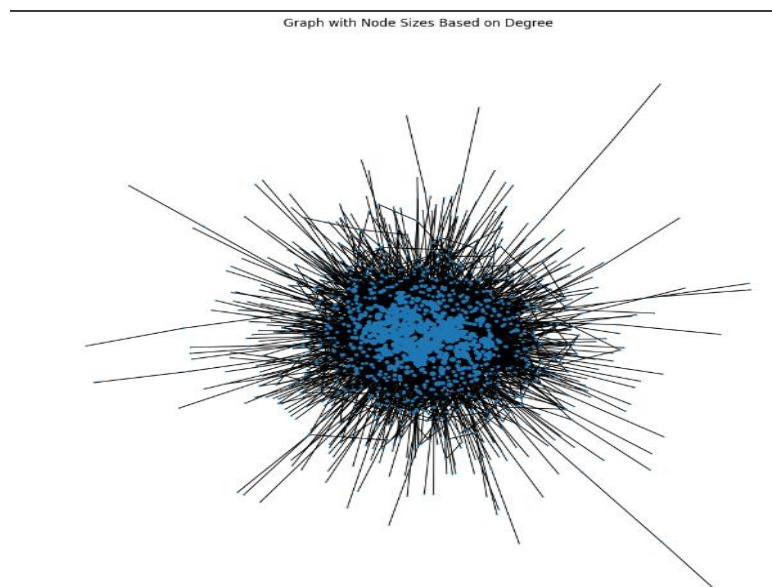
3. Size of connected component



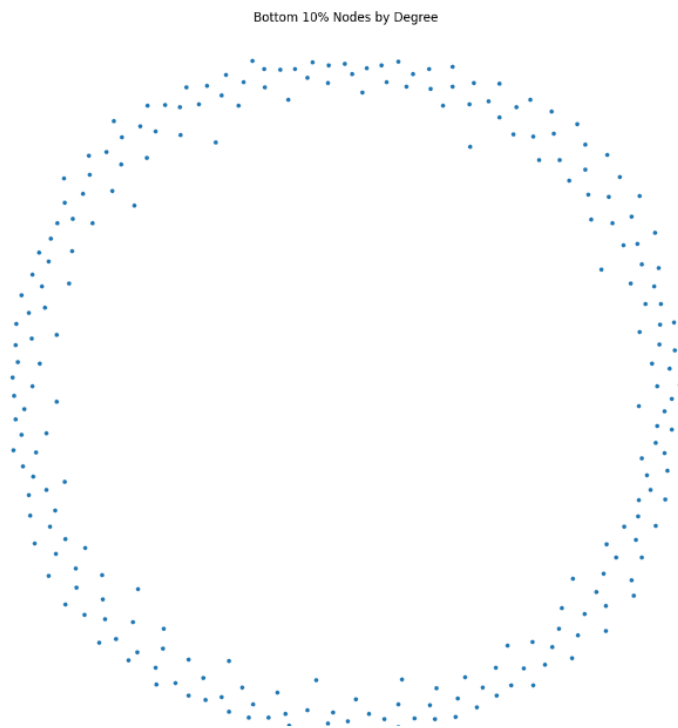
Part C (Code in Python)

<https://colab.research.google.com/drive/1P1ZpobT07-3KXiRK0REkp8pOxn4gA1EP?usp=sharing>

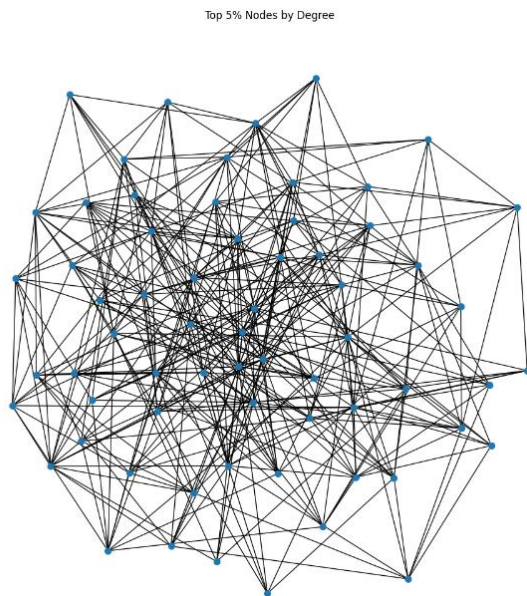
1. Degree Distribution of graph.



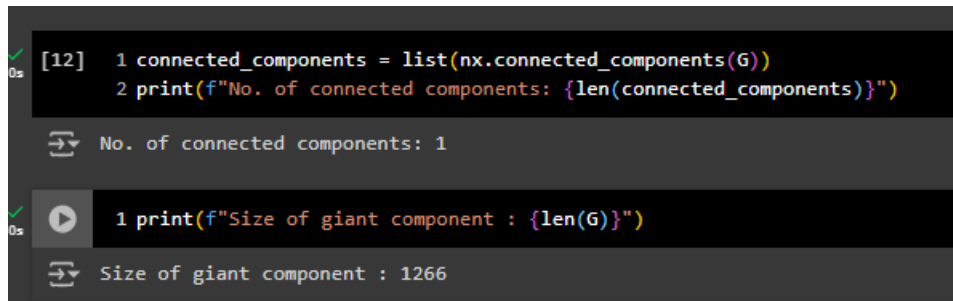
2. Bottom 10% of nodes and the connection among them.



3. Top 5% nodes and connection between them.



4. All the connected components of the network.

A screenshot of a Jupyter Notebook interface showing two code cells. The first cell, labeled [12], contains two lines of Python code: `1 connected_components = list(nx.connected_components(G))` and `2 print(f"No. of connected components: {len(connected_components)}")`. The output of this cell is `No. of connected components: 1`. The second cell contains a single line of code: `1 print(f"Size of giant component : {len(G)}")`. The output of this cell is `Size of giant component : 1266`.

```
[12] 1 connected_components = list(nx.connected_components(G))
      2 print(f"No. of connected components: {len(connected_components)}")

No. of connected components: 1

1 print(f"Size of giant component : {len(G)}")

Size of giant component : 1266
```

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