Group K – Catch Me If You Can Intermediate Report

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1 CATCH ME IF YOU CAN - social network analysis

1.1 Introduction

Introduce the project..., this will serve as a guide throughout the project.



1.2 Group Members:

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1.3 Project Overview:

Section ?? – Understanding the network we are dealing with

Section ?? – Insights into local connectivity and overall cohesiveness.

Section ?? – Dealing with the centralities

1.4 Tools and Technologies Used:

- Programming Language: Python
- Libraries: Pandas, NumPy, Matplotlib, NetworkX, JSON, HeapQ
- Platform: Jupyter Notebook

LIBRARIES WE ARE GOING TO USE:

```
[63]: import pandas as pd
import networkx as nx
import matplotlib.pyplot as plt
import json
import heapq
import numpy as np
```

OPENING THE FILE:

```
[64]: nodes_df = pd.read_csv('a) Catch_me_if_you_can/nodes.csv')
edges_df = pd.read_csv('a) Catch_me_if_you_can/edges.csv')
```

The following line takes the 'viz' column of the nodes_df DataFrame and performs two operations:

First, it replaces all single quotes (') with double quotes (") to make the string JSON-compatible. Then, it parses the modified string as a JSON object, converting it into a *Python dictionary*.

```
[65]: nodes_df[' viz'] = nodes_df[' viz'].apply(lambda row: row.replace("'", '"')).

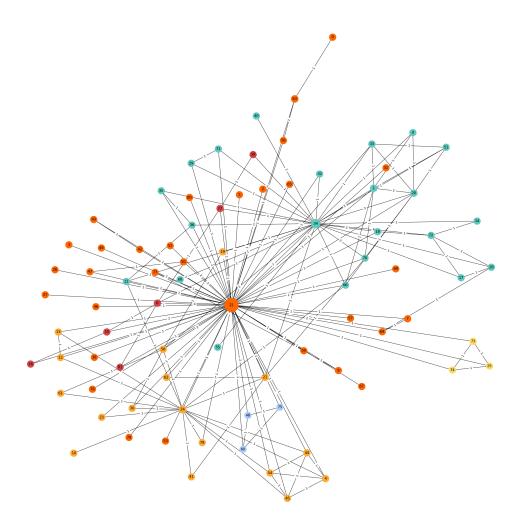
→apply(json.loads)
```

This section initializes an empty undirected graph G. It then iterates over the rows of nodes_df to add nodes to the graph, and over the rows of edges_df to add edges. Each node and edge is added with its associated attributes.

Graph G is now ready to be visualized.

```
[67]: plt.figure(figsize=(25, 25))
    nx.draw(G, pos=positions, node_color=colors, node_size=sizes, with_labels=True)
    edge_labels = {(u, v): d['weight'] for u, v, d in G.edges(data=True)}
    nx.draw_networkx_edge_labels(G, pos=positions, edge_labels=edge_labels)
    plt.title('CATCH ME IF YOU CAN', size=40, weight='bold')
    plt.show()
```

CATCH ME IF YOU CAN



NUMBER OF NODES, NUMBER OF EDGES, AVERAGE DEGREE AND THE DENSITY

```
[68]: n = G.number_of_nodes()
m = G.number_of_edges()
print("Number of nodes: ", n)
print("Number of edges: ", m)
```

Number of nodes: 82 Number of edges: 162

the degree k(v) of a node v is computed by iterating over each edge in the edges_df dataframe and incrementing the degree count for both the source and target nodes. Given an edge ((u, v)) from

the dataframe edges_df:

$$k(u) = k(u) + 1$$

$$k(v) = k(v) + 1$$

where (k(u)) and (k(v)) are the degrees of nodes (u) and (v) respectively.

The average degree k is computed by summing up the degrees of all nodes and dividing by the total number of nodes.

 $\langle k \rangle = \frac{\sum_{v \in V} k(v)}{N}$

where (N) is the total number of nodes in the graph and (k(v)) is the degree of node (v).

```
[69]: def compute_degrees(edges_df):
    degrees = {}

    for index, row in edges_df.iterrows():
        source = row['# source']
        target = row[' target']

        degrees[source] = degrees.get(source, 0) + 1
        degrees[target] = degrees.get(target, 0) + 1

    return degrees

def compute_average_degree(edges_df):
    degrees = compute_degrees(edges_df)
    total_degree = sum(degrees.values())
    N = len(degrees)

return total_degree / N
```

```
[70]: print("Average degree: ", compute_average_degree(edges_df))
```

Average degree: 3.951219512195122

The formula for the density (D) of an undirected graph is:

$$D = \frac{2M}{N(N-1)}$$

Where: - (M) is the number of edges in the graph. - (N) is the number of nodes in the graph.

```
[71]: def density(G):
    n = len(G.nodes())
    m = len(G.edges())
    if n == 0 or n == 1:
        return 0.0
    density = (2 * m) / (n * (n - 1))
```

```
print('Density:', density(G))
```

Density: 0.04878048780487805

WEEK 2 ANALYSIS -

While considering the largest component of your network. Depending on what you prefer/seems more relevant in your graph

- Compute Average clustering and Transitivity number
- Implement a function computing the transitivity using basic function of networkx

DIJKSTRA'S ALGORITHM

```
[72]: def dijkstra_generic(G, start_node, use_weights=True):
          distances = {node: float('inf') for node in G.nodes()}
          distances[start node] = 0
          pq = [(0, start_node)]
          while pq:
              current_distance, terrent_node = heapq.heappop(pq)
              for neighbor, edge_attr in G[current_node].items():
                  weight = edge_attr['weight'] if use_weights else 1
                  new_distance = current_distance + 1/weight
                  if new_distance < distances[neighbor]:</pre>
                      distances[neighbor] = new_distance
                      heapq.heappush(pq, (new_distance, neighbor))
          return distances
      all_unweighted_shortest_path_lengths = {node: dijkstra_generic(G, node,_
       →use weights=False)
                                               for node in G.nodes()}
      all_weighted_shortest_path_lengths = {node: dijkstra_generic(G, node,_
       →use weights=True)
                                             for node in G.nodes()}
```

Given the nature of the data (same-scene appearances), the average clustering and transitivity seemed be more relevant to us. These metrics would provide a clearer picture of character interactions and how characters are grouped or clustered in the narrative. It would highlight which characters frequently share scenes and are thus likely central to the movie's main plot or subplots.

We, thus, implemented a first function which retrieves the number of triangles in the graph. This is done by iterating over each node in the graph and counting the number of triangles that the node is part of. The number of triangles is then divided by the number of connected triples in the graph to get the transitivity.

TRIANGLES -

 $\{u,v,w\}$ forms a triangle if (u,v),(v,w), and (u,w) are edges in the graph

TRANSITIVITY -

$$T = \frac{3 \times \text{triangles_count}}{\text{wedges}}$$

Where: 'TRIANGLES COUNT' is the number of unique triangles in the graph and 'WEDGES' is the sum of the number of pairs of neighbors for each node in the graph.

```
[73]: def find_triangles(G):
          triangles = set()
          for u, v in G.edges():
               # Find common neighbors
               common_neighbors = set(G.neighbors(u)).intersection(G.neighbors(v))
               for w in common neighbors:
                   triangle = tuple(sorted([u, v, w]))
                   triangles.add(triangle)
          return triangles
      triangles_in_graph = find_triangles(G)
      def custom_transitivity_from_triangles(G, triangles_set):
          triangles_count = len(triangles_set) # Count of unique triangles
          triples = sum(len(list(nx.ego_graph(G, v))) * (len(list(nx.ego_graph(G, u))) * (len(list(nx.ego_graph(G, u))))
       \rightarrowv))) - 1) for v in G) / 2
          if triples == 0:
               return 0
          return (3 * triangles count) / triples
      custom_trans_from_triangles = custom_transitivity_from_triangles(G,_
       →triangles_in_graph)
```

We then implemented a second function which computes the average clustering coefficient of the graph. This is done by iterating over each node in the graph and computing the clustering coefficient of the node.

Clustering Coefficient for a Node: For a given node (v) with degree (k):

$$C(v) = \frac{2T(v)}{k(k-1)}$$

Where (T(v)) is the number of triangles involving node (v).

The clustering coefficient of the node is then added to the total clustering coefficient of the graph. The total clustering coefficient is then divided by the number of nodes in the graph to get the average clustering coefficient.

Average Clustering Coefficient: For the entire graph (G):

$$C = \frac{1}{n} \sum_{v \in V} C(v)$$

Where (n) is the number of nodes in (G) and (V) is the set of nodes.

```
[74]: def avg_clustering(G):
    triangles_set = find_triangles(G)
    total_clustering_coeff = 0
```

```
for node in G.nodes():
    neighbors = list(G.neighbors(node))
    k = len(neighbors)
    if k < 2:
        continue
    # Count triangles involving the node
    T = sum(1 for triangle in triangles_set if node in triangle)
        total_clustering_coeff += 2 * T / (k * (k - 1))
    return total_clustering_coeff / len(G.nodes())

avg_clustering_custom = avg_clustering(G)
avg_clustering_custom</pre>
```

[74]: 0.5990508780501608



WEEK 3 ANALYSIS -

Depending on what seems more relevant in your graph, pick one of the following local notions

- Decay centrality
- Betweeness centrality
- Closeness centrality
- Clustering
- Any other notions that you invent
- 1) Provide a code computing the given centrality using basic functions of networks (you are not allowed to use directly nx."what you want").
- 2) Discuss why you picked this measure and who is the most central in your network based on your choice.
- 3) Provide the cumulative distribution for this centrality and give a graphical representation of your graph(log-log, log or normal representation as you think it is more relevant).

CLOSENESS CENTRALITY could be a relevant approach in our analysis. It would allow us to identify the characters who are most central to the narrative. These characters would be the ones who are most likely to be involved in the main plot or subplots of the movie.

Therefore, given the following formula for the Closeness Centrality:

$$C(u) = \frac{n-1}{\sum_{v \neq u} d(v, u)}$$

This is the corresponding implementation:

```
[75]: def custom_closeness_centrality(G):
    closeness_centrality = {}
```

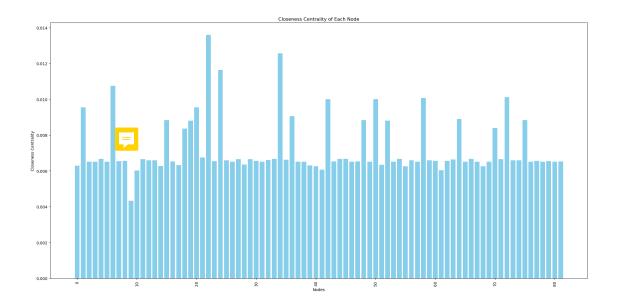
Most central character: 22 Closeness centrality: 0.013585198602665282

It results that our protagonist (node 22) has relatively low closeness centrality, shich suggest that the character is not closely connected to many other characters in the movie, indicating a potential narrative focus on isolated or introspective journeys, distinct subplots or a fragmented narrative.

DISPLAYING THE CLOSENESS CENTRALITY FOR EACH NODE

```
[76]: nodes = list(closeness_centrality_custom.keys())
    closeness_values = list(closeness_centrality_custom.values())

plt.figure(figsize=(20,10))
    plt.bar(nodes, closeness_values, color='skyblue')
    plt.xlabel('Nodes')
    plt.ylabel('Closeness Centrality')
    plt.title('Closeness Centrality of Each Node')
    plt.xticks(rotation=90)
    plt.tight_layout()
    plt.show()
```



DISPLAYING THE CUMULATIVE DISTRIBUTION FOR THE CLOSENESS CENTRALITY

```
[77]: closeness_values = sorted(list(closeness_centrality_custom.values()))

ccdf_y = np.arange(1, len(closeness_values) + 1)[::-1] / len(closeness_values)

ccdf_x = closeness_values

plt.figure(figsize=(12,8))
 plt.plot(ccdf_x, ccdf_y, "x", color='blue')
 plt.xlabel('Closeness Centrality')
 plt.ylabel('Complementary Cumulative Distribution')
 plt.title('Complementary Cumulative Distribution of Closeness Centrality')
 plt.grid(True, which="both")
 plt.show()
```

