Not Able to get dataset to generated random

```
In [12]: import pandas as pd
         import random
         # Create a list of users
         users = [f'user_{i}' for i in range(1, 51)] # 50 users
         # Generate random interactions between users
         interactions = []
         for _ in range(200): # Generating 200 random interactions
             user1 = random.choice(users)
             user2 = random.choice(users)
             while user1 == user2: # Avoid self-interactions
                 user2 = random.choice(users)
             interactions.append((user1, user2))
         # Create a DataFrame
         df_interactions = pd.DataFrame(interactions, columns=['user1', 'user2'])
         # Save to CSV (for your reference)
         df_interactions.to_csv('synthetic_social_network.csv', index=False)
         # Show the first few rows of the dataset
         df interactions.head()
             user1 user2
Out[12]:
```

	user1	user2
0	user_9	user_35
1	user_34	user_15
2	user_28	user_49
3	user_24	user_43
4	user_37	user_38

Load and Preprocess the Data

```
import pandas as pd
import networkx as nx

# Load the dataset (e.g., Twitter dataset with user interactions)
df = pd.read_csv('synthetic_social_network.csv')

# Inspect the dataset structure
df.head()

# Preprocessing: create edges for user interactions, remove duplicates, e
# Example: Let's say the dataset has 'user1' and 'user2' columns represen
edges = df[['user1', 'user2']].drop_duplicates()
```

```
# Construct a graph using NetworkX
G = nx.from_pandas_edgelist(edges, 'user1', 'user2')
```

Graph Construction and Representation

Apply Information Diffusion Model

```
In [9]: import random
        # Set transition probabilities
        beta = 0.3 # S -> E
        gamma = 0.1 \# E \rightarrow I
        delta = 0.05 \# I -> R
        # Randomly infect a small set of nodes initially
        for node in random.sample(list(G.nodes()), 5): # Convert G.nodes() to a
            G.nodes[node]['status'] = 'Infected'
        def update_status(G):
            new_status = {}
            for node in G.nodes():
                 status = G.nodes[node]['status']
                 if status == 'Susceptible':
                     # Check if any neighbor is infected
                     infected_neighbors = [n for n in G.neighbors(node) if G.nodes
                     if infected_neighbors and random.random() < beta:</pre>
                         new_status[node] = 'Exposed'
                 elif status == 'Exposed':
                     # Transition from Exposed to Infected
                     if random.random() < gamma:</pre>
                         new_status[node] = 'Infected'
                 elif status == 'Infected':
                     # Transition from Infected to Recovered
                     if random.random() < delta:</pre>
                         new_status[node] = 'Recovered'
            # Update statuses for the next round
            nx.set_node_attributes(G, new_status, 'status')
        # Run the simulation for a certain number of time steps
        time_steps = 20
        for t in range(time_steps):
            update_status(G)
```

Evaluate the Results

Final counts: {'Susceptible': 0, 'Exposed': 1, 'Infected': 12, 'Recovere d': 37}

Visualize the Graph

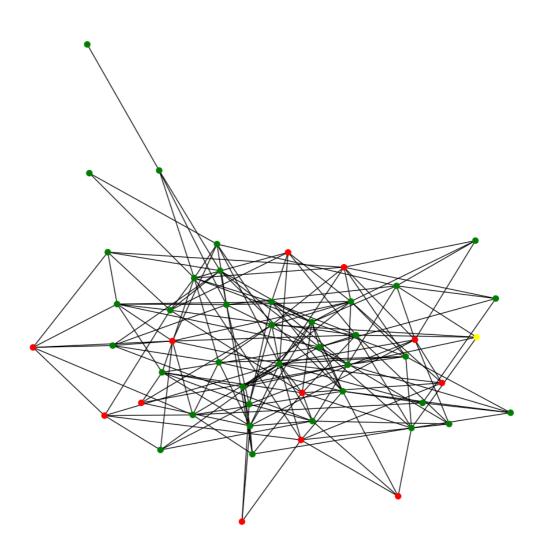
```
In [11]: import matplotlib.pyplot as plt

def draw_graph(G, t):
    color_map = {
        'Susceptible': 'blue',
        'Exposed': 'yellow',
        'Infected': 'red',
        'Recovered': 'green'
    }

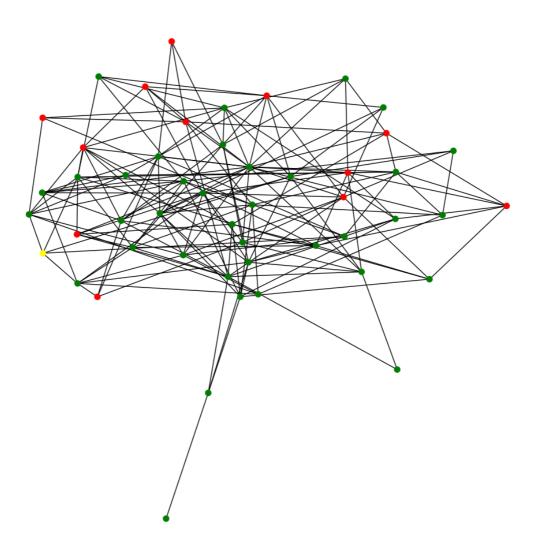
    colors = [color_map[G.nodes[node]['status']] for node in G.nodes()]
    plt.figure(figsize=(10, 10))
    nx.draw(G, node_color=colors, with_labels=False, node_size=50)
    plt.title(f'Time step {t}')
    plt.show()

# Visualize at different time steps
for t in [0, 5, 10, 15, 20]:
    draw_graph(G, t)
```

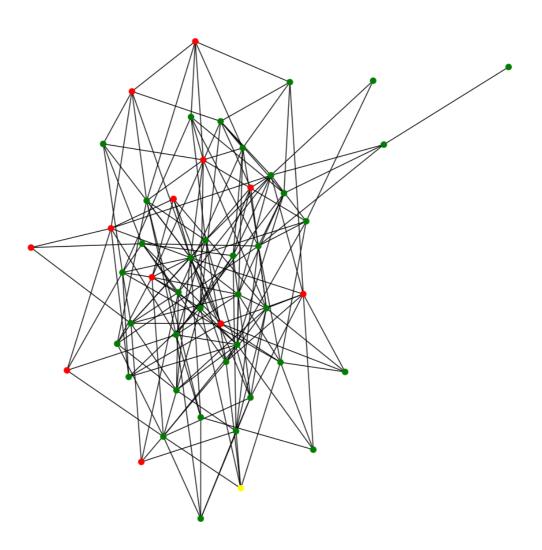
Time step 0



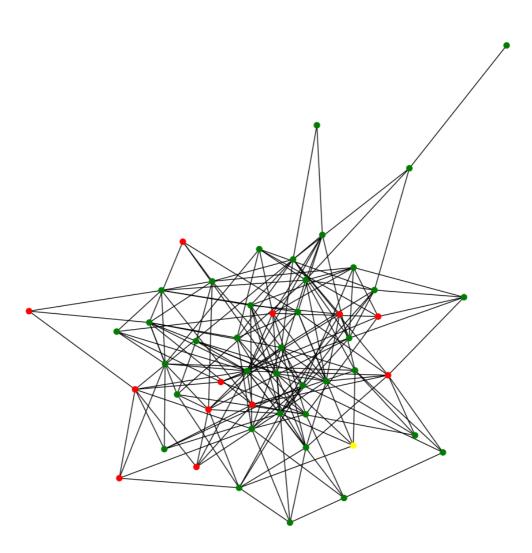
Time step 5



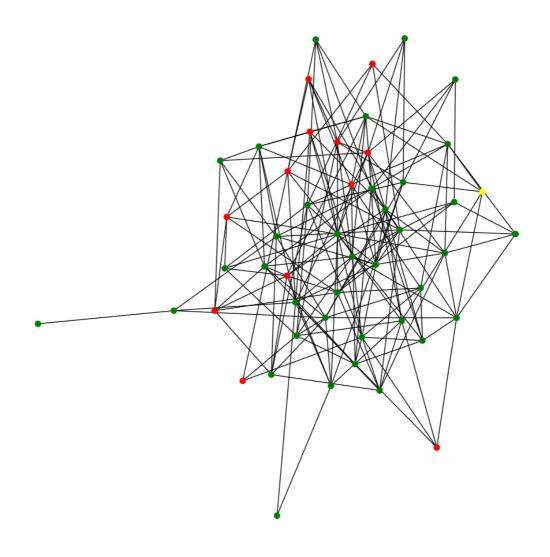
Time step 10



Time step 15



Time step 20



In []: