Epidemics Graph Neural Network Node Classification and Link Prediction

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Abstract

The COVID-19 pandemic showed that contact tracing helped mitigate the spread of the virus. However, manual contact tracing is slow and can be inaccurate. Thus, this project aims to automate contact tracing by utilizing Graph Neural Networks (GNNs). Our preliminary work on network analysis showed that the contact network is a mix of an exponential and scale-free network. Also, our simulation showed that during the first 12 hours, the infection does not spread much, but then it starts spreading steadily.
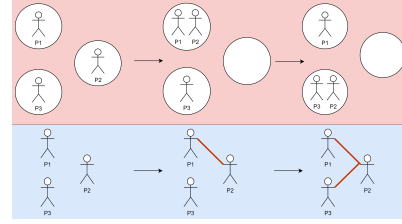
1. Introduction and Motivation

When COVID-19 first appeared, manual tracing was deployed to mitigate the initial outbreak. Contact tracing is the process of tracking how the virus spreads by identifying people who may have come in contact with an infected person, and then asking them to isolate and get tested.

However, the pandemic revealed that the COVID-19 virus spread faster than manual contact tracing [2]. Thus, this project's objective is to automate contact tracing by incorporating machine learning using GNNs to hopefully increase the mitigation of the spread of COVID-19 when compared to manual contact tracing.

2. Previous Work

Mathematical models, classical ML models, and graph-based ML models have been used to predict virus spread.

The SEIRD model is a mathematical model that predicts the change in Susceptible, Exposed, Infected, Recovered, and Deceased people over time by using differential equations. [3]. SIR is simpler version of the SIERD model [11].

The LSTM is an ML model that has been used to predict the number of cases over time [3]. A hybrid of SIRD and LSTM helps account for time dependent parameters of the SIRD model [1]. Also, GNNs, which are graph-based ML models, have been used on mobility data to predict virus spread and for link prediction for contact tracing [8][9].

Figure 1: Example of contact network generation

3. Approach

3.1 Network Generation and Analysis

We used the foursquare dataset to build a contact graph of Austin, TX [7]. Each entry contains a device ID, a location ID, UTC date and hour, and a dwell time, which tell us when and how long a person visited a location. Given this data, we generated a contact graph of Austin.

Firstly, we used data from July 1st, 2020 to July 5th, 2020 to create a sample contact network. Our nodes were all the unique device IDs in the dataset, which correspond to people. For our edges, we used the following logic: we ignored entries with a dwell time less than 60 minutes, as we assumed this not enough time to make significant contact with others. Then, we used the UTC date and hour with the dwell time to determine the arrival and departure time interval for each entry. We then compared every entry with every other entry. If the entries' locations were the same and if intervals overlapped by at least 60 minutes, we considered this as a contact between the two people and added an edge between them. We will call this the 5-Day Contact Network. This network captures the meaningful contacts that occurred from July 1st, 2020 to July 5th, 2020. Figure 1 shows an example of a sample contact network generation.

Then we also created a set of 62 contact networks, one for each day from July 1st, 2020 to August 31st, 2020. We used the same logic as before to create these individual contact networks. We will call this set of networks the Temporal Contact Networks. Each day's network cap-

tures the meaningful contacts that occurred for that day. Furthermore, we analyzed how the Clustering Coefficient and Average Node Degree changed over time by calculating these metrics for each network in the Temporal Contact Networks.

We also created a Susceptible-Infected-Recovered simulation using the Temporal Contact Networks. We ran the simulation from July 1st, 2020 to August 31st, 2020. Here are the parameters and assumptions that were made for the simulation:

- Contact between people that is less than 60 minutes is not considered significant enough to spread the virus.

- If a susceptible person came into contact with an infected person for at least one hour, then they get infected with a probability of 0.30. This is called the infection rate (IR).

- The IR is constant throughout the simulation.

- An infected person will recover after seven days. This is called the recovery period (RP).

- The RP is constant throughout the simulation.

- A person can only be infected if they were previously susceptible, and a person can only be recovered if they were previously infected.

- Initially, 20% of the people, chosen at random, are infected. The rest are susceptible.

- Only infected people can infect others.

3.2 Machine Learning

After generating and analyzing the 5-Day Contact Network, Temporal Contact Networks, and performing the SIR simulation, we moved towards leveraging graph learning techniques to perfom link prediction, which is the fundamental task behind automating contact tracing. For this milestone, we focused on performing link prediction on a static graph. In order to have enough data to train and evaluate our model(s), we generated a contact network for the first five days of July. We used the same technique described in building our July 1st contact network to build the 5 day contact network; so the nodes are people who have a dwell time of at least 60 minutes at one location, and edges represent people who have come into contact with each other in the 5 day period.

Our first goal was to create a baseline link prediction model. We used the node2vec algorithm to generate node embeddings for each node in the graph [4]. We then had to generate our dataset of edges. In order to perform link prediction, we need the set of positive edges, which are the
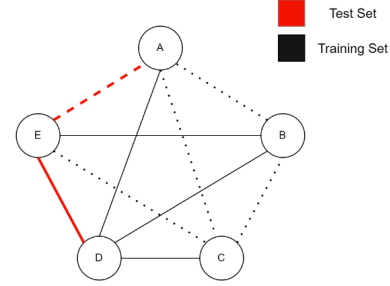

Figure 2: Example of positive and negative edges for train and test data

edges present in the network, and we need a set of negative edges, which are the edges not present the network. This allows us to boil down the link prediction to a binary classification problem. Given our network, we created a set of negative edges that was equal in size to the set of positive edges to ensure balanced training. Using the node2vec embeddings and the set of positive and negative edges, we trained a GraphSAGE model to perform link prediction on the static 5 day graph [5].

A sample example of training and testing data is shown in Figure 2. The black lines representing training data and red lines represent testing data. The solid lines represent positive edges and dotted lines represent negative edges. The training data is used to train the model, and the testing data is used to evaluate the model.

After creating the baseline model, we searched for ways to improve the model's performance on the graph. This would include performing feature engineering techniques to add dimensions to our node embeddings and exploring the use of other GNN architectures such as the Graph Convolutional Network (GCN) and/or the Graph Attention Network (GAT) [6] [10]. We hoped to be able to finetune the model and improve its performance to the point we could use it to perform link prediction on the Temporal Contact Networks.

4. Experimental Setup and Results

In Figure 3, we can see the 5-Day Contact Network. Network properties for this network were also calculated. The average node degree is 4.278, the network diameter is 20, the average clustering coefficient is 0.432, and the average path length is 6.95. In addition to this, the degree distribution was mainly an exponential distribution with subtle hints at a power-law as shown in Figure 4. This can also be seen from the network itself, as we can see the presence of a few hubs in the network. This makes sense, as we should expect a social network to be scale-free, but we do not have all the data points, so it is not fully scale-free on the sample network.
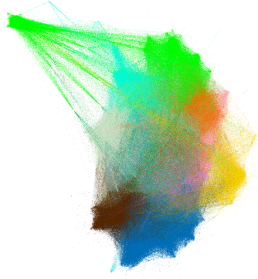
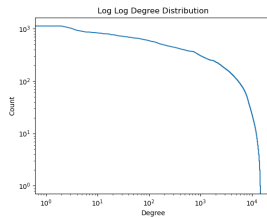Figure 3: 5-Day Contact Network from July 1st, 2020 to July 5th, 2020

Figure 4: 5-Day Contact Network Degree Distribution

Figure 5: SIR Simulation Results

Figure 6: Clustering Coefficient and Average Node Degree for Temporal Contact Networks

In addition, the simulation results are shown in Figure 5. The simulation results show that the number of infected people initially increases, but then decreases significant and approached 0. Whereas the number of recovered people exhibit the opposite behavior. This makes sense since the infected people recover after seven days can no longer become susptible again. Also, the number of susceptible people decreases over time, which makes sense since they are getting infected and recovering. The maximum number of infected people on any given day is about 11,500 and the total number of people infected over the simulation is about 14,300.

It is important to note that this simulation was run on limited data. The total number of nodes in the simulation, which is the total number of people, is around 45,000, whereas the actual population in Austin, TX in 2020 was around 2 million. Furthermore, this model assumes a closed population, which is not the case in Austin, TX. People are constantly moving in and out of Austin, TX. Thus, the simulation results are not representative of the actual spread of the virus in Austin, TX. However, this simulation does show that the virus can spread quickly, as seen in the first week of the simulation. Thus, it is important to have an efficient and accurate contact tracing system to mitigate the spread of the virus.

In addition, the analysis of Clustering Coefficient and Average Node Degree for the Temporal Contact Networks are shown in Figure 6. The results demonstrate a periodic patte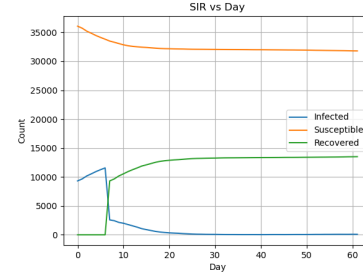rn. It is evident that the period is 7 days, and the peaks occur on the weekends. These re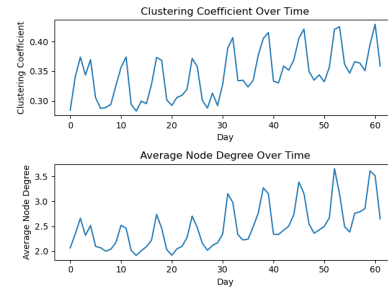sults makes sense since people are more likely to go out and socialize on the weekends. This increase in socialization on the weekends leads to more contacts, which leads to a higher clustering coefficient and average node degree.

After performing the analysis on the 5-Day Contact Network and Temporal Contact Networks, we moved towards performing link prediction on the 5-Day Contact Network. For our baseline model, as stated in the approach, we used a GraphSAGE model with node2vec embeddings. We trained the model for 1000 epochs on a 90/10 train/test split with the Adam optimizer. The model achieved an AUC score of 0.49 on the test data. We can see that this performs worse than random guessing. Part of this could be due to the fact that no internal node features were used in the node embeddings.

To improve upon this, we performed feature engineering to generate additional features for each node. Here are the additional featues the generated for every node (person) in the graph:

- Average location travelled to per day

- Average distance travelled per day

- Gender

- SAG Score

- Age

By adding these features to the node2vec embeddings, we were able to improve the AUC score to 0.63, which is a significant improvement over the baseline model. We can see that the additional features helped the model learn the graph structure better.

We then attempted to improve upon this by taking the features we created and normalizing them before appending them. We saw that the features were all of different scales compared to each other and the node2vec embeddings, so we hypothesized that normalizing the features would improve the model's performance. This, again, improved our model's performance, as we were able to achieve an AUC score of 0.75.

Finally, we attempted different model architectures to see if we could improve the model's performance. More specifically, we tried using the GCN model. With the same features and data, we were able to achieve an AUC score of 0.91, which is a significant improvement over the previous model.

From these performance metrics, we can see that performing link prediction on the Temporal Contact Networks is a feasible task. We can also see that the GCN model performs significantly better than the GraphSAGE model. Thus, we plan to use the GCN model to perform link prediction on the Temporal Contact Networks for M3.

5. Conclusion and Short-Term Plans

Through the analysis of the network, we were able to determine the network of contacts is a combination of an exponential and scale-free network. Some people came into contact with many other people whereas others stayed within their cliques. The simulation showed that the number of infected people initially stayed relatively constant, but after 12 hours, infections began increasing steadily.

For M2, we plan to do a more complete analysis of the network by considering both July and August data. We also plan to run more realistic simulations by accounting for the incubation period and deceased/recovered people. We also plan to implement GNNs at a small scale to perform link prediction. Specifically, we will use GNNs to predict the people that come into contact with previously infected people, allowing us to predict the spread of the virus.

For this milestone, Afnan generated and analyzed the contact network for July 1st, 2020, and Jaykumar created and analyzed the simulation of the hourly spread of the virus.

References

[1] A. Bousquet, W. H. Conrad, S. O. Sadat, N. Vardanyan, and Y. Hong. Deep learning forecasting using time-varying parameters of the sird model for covid-19, February 22 2022.

[2] S. Flaxman, S. Mishra, A. Gandy, H. J. T. Unwin, T. A. Mellan, H. Coupland, and et al. Estimating the number of infections and the impact of non-pharmaceutical interventions on covid-19 in european countries: technical description update, 2020.

[3] T. Geroski, A. Blagojevic, D. M. Cvetković, A. M. Cvetković, I. Lorencin, S. B. Šegota, D. Milovanovic, D. Baskic, Z. Car, and N. Filipovic. Epidemiological predictive modeling of covid-19 infection: Development, testing, and implementation on the population of the benelux union, October 28 2021.

[4] A. Grover and J. Leskovec. node2vec: Scalable feature learning for networks, 2016.

[5] W. L. Hamilton, R. Ying, and J. Leskovec. Inductive representation learning on large graphs, 2018.

[6] T. N. Kipf and M. Welling. Semi-supervised classification with graph convolutional networks, 2017.

[7] C. D. Lab. Foursquare Community Mobility Data with Basemap (US), 2020.

[8] K. Skianis, G. Nikolentzos, B. Gallix, R. Thiebaut, and G. Exarchakis. Predicting covid-19 positivity and hospitalization with multi-scale graph neural networks, March 31 2023.

[9] C. W. Tan, P.-D. Yu, S. Chen, and H. V. Poor. Deeptrace: Learning to optimize contact tracing in epidemic networks with graph neural networks, 2023.

[10] P. Veličković, G. Cucurull, A. Casanova, A. Romero, P. Liò, and Y. Bengio. Graph attention networks, 2018.

[11] R. S. Yadav. Mathematical modeling and simulation of sir model for covid-2019 epidemic outbreak: A case study of india, May 21 2020.