



# **Epidemic Source Detection:**





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A Benchmark Study

Evaluating Graph Neural Networks for

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#### Introduction

Epidemic source detection has been actively researched since Shah and Zaman [1] introduced the problem on static networks in 2010. More recently, several studies have explored using Graph Neural Networks (GNNs) for source identification [2-5].

#### **Problem formulation**

We consider static, undirected networks on which a continuous-time susceptibleinfectious-recovered (SIR) infection process unfolds. The SIR parameters are known, and a full snapshot of node states is observed at a known time T after the process begins.

Goal: Infer the single source of the epidemic based on the network and the observed node states (Fig. 1). This constitutes a graph prediction problem.

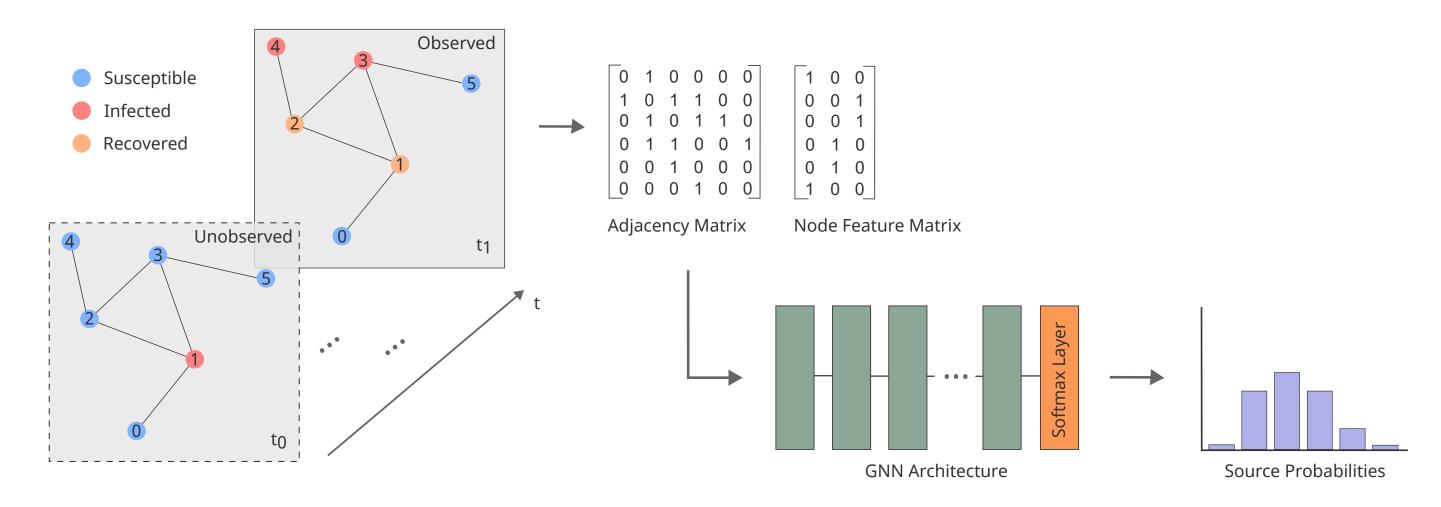


Figure 1. At time  $t_1 = t_0 + T$  we observe a full snapshot of the epidemic which translates into the two inputs for the GNN, the adjacency matrix of the graph and the one-hot encoded epidemic states of all nodes. The final output of the GNN is a source probability distribution.

#### Research question

Recent GNN-based approaches to source detection offer limited insight into how these models compare with traditional methods. This raises the central research question of our work: How do GNNs perform relative to traditional source detection methods, and what is their true potential for this task?

## **Our GNN architecture**

## Message-passing layers

Our architecture is primarily based on L graph convolution layers [6] that, for any node v, can be defined as follows:

$$\mathbf{h}_v^{(l)} = \text{ReLU}\left(\mathbf{B}_l \; \mathbf{h}_v^{(l-1)} + \mathbf{W}_l \sum_{u \in \mathcal{N}(v)} \mathbf{h}_u^{(l-1)}\right) \qquad l = 1 \dots, L$$

The initial embedding of a node is the one-hot encoded representation of its epidemic state, i.e.,  $\mathbf{h}_v^{(0)} \in \{0,1\}^3$ . The final convolutional layer returns the embedding matrix  $\mathbf{H}^{(L)} \in \mathbb{R}^{|V| \times s}$ , where s denotes the embedding dimension of the final layer.

## **Output layer**

The output layer constitutes of a linear transformation followed by a softmax activation, i.e.,  $\hat{\mathbf{y}} = \text{Log-Softmax} \left[ \mathbf{H}^{(L)} \mathbf{w} \right]$ , with  $\hat{\mathbf{y}}$  representing the (log-) source probability distribution over |V| nodes.

# **Training setup**

- Simulation of 500 training instances per node in the graph.
- Split in 90% training and 10% validation instances.
- Loss function: negative log likelihood loss.
- Mini-batch gradient descent (batch size: 128).
- Optimizer: Adam (learning rate: 0.001).
- Dropout rate up to layer L-1 is 0.05, for layer L it is 0.2.
- Early stopping based on validation loss with a patience of 5 epochs.

## **Benchmark methods**

- Random selection of the source (RANDOM).
- Jordan centrality (CENTRALITY) [7].
- Untrained GNN (GNN-UNTRAINED).
- Soft margin estimator (SME) [8].
- Factorized likelihood method utilizing Monte Carlo estimates of node state probabilities (MCMF) [9].

# References

[1] Shah, D. et al., SIGMETRICS Perform. Eval. Rev. 38, 1 (2010). 10.1145/1811099.1811063 [2] Dong, M. et al., CIKM, (2019). 10.1145/3357384.3357994

[3] Shah, C. et al., arXiv, (2020). https://arxiv.org/abs/2006.11913 [4] Sha, H. et al., IEEE DSAA, (2021). 10.1109/DSAA53316.2021.9564188

[5] Ru, X. et al., AAAI, (2023). 10.1609/aaai.v37i8.26152

[6] Morris, Ch. et al., AAAI, (2019). 10.1609/aaai.v33i01.33014602

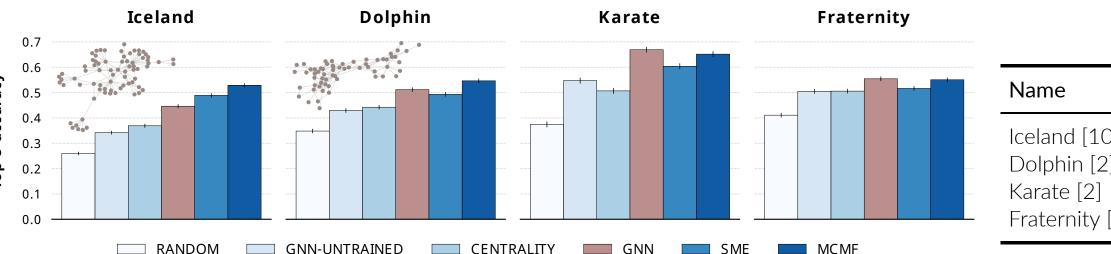
[7] Zhu, K. et al., *IEEE TNET* 24, 1 (2016). 10.1109/TNET.2014.2364972 [8] Antulov-Fantulin, N. et al., Phys. Rev. Lett. 114, 248701 (2015). 10.1103/PhysRevLett.114.248701

[9] Sterchi, M. et al., Scientific Reports 13, 1 (2023). 10.1038/s41598-023-38282-8

# Results

#### **Detection performance overall**

We simulate 200 test outbreaks per node in the graph. The test outbreaks have a basic reproduction number of  $R_0 \approx 2$  and the snapshots are taken when 40% of the nodes are infected, on average. The outbreaks in the train and test set have the same characteristics.



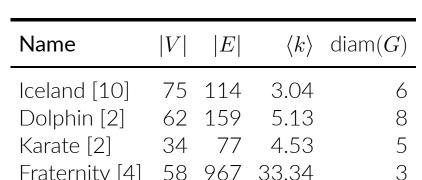


Figure 2. Top-5 accuracy of GNN and benchmark methods for the four empirical networks.

Table 1. Overview of networks.

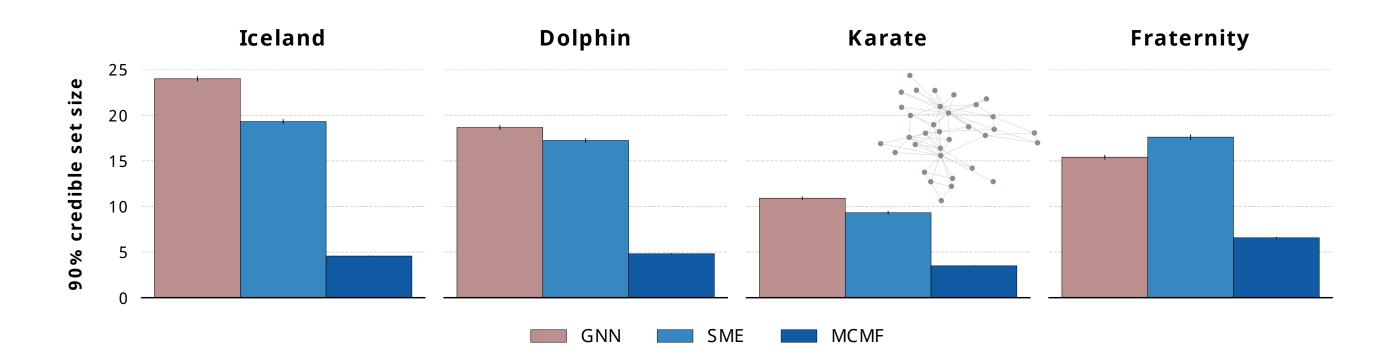


Figure 3. 90% credible set size for the methods that output a distribution.

#### Detection performance by outbreak size

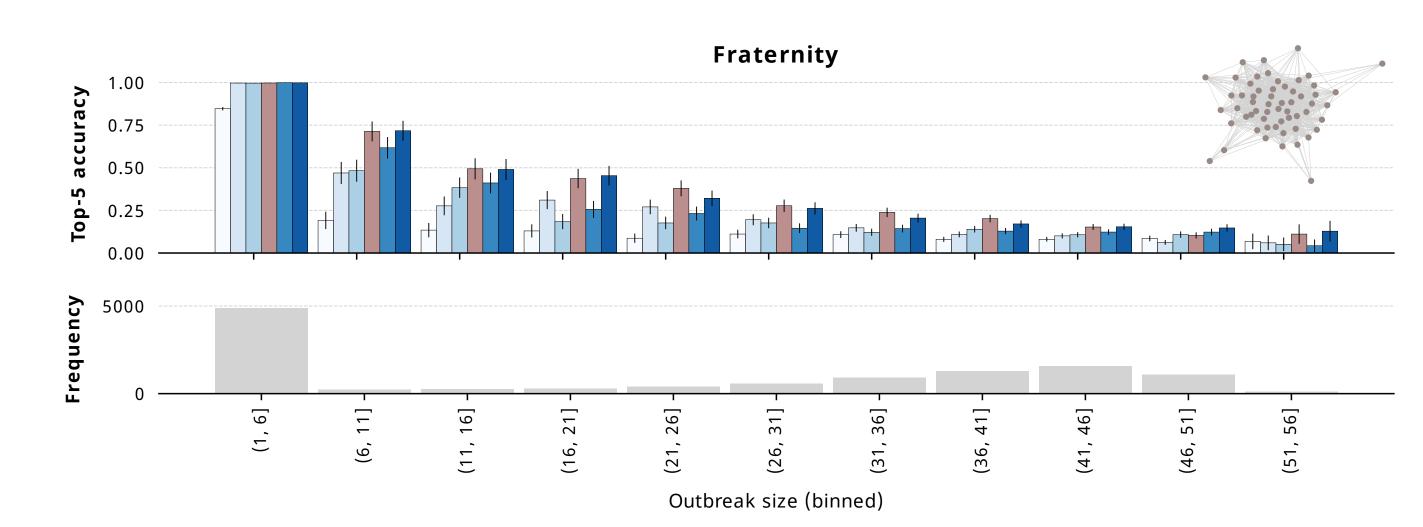


Figure 4. Top-5 accuracy of all methods for the Fraternity network, categorized by outbreak size.

## Single oubreak scenario on Karate network

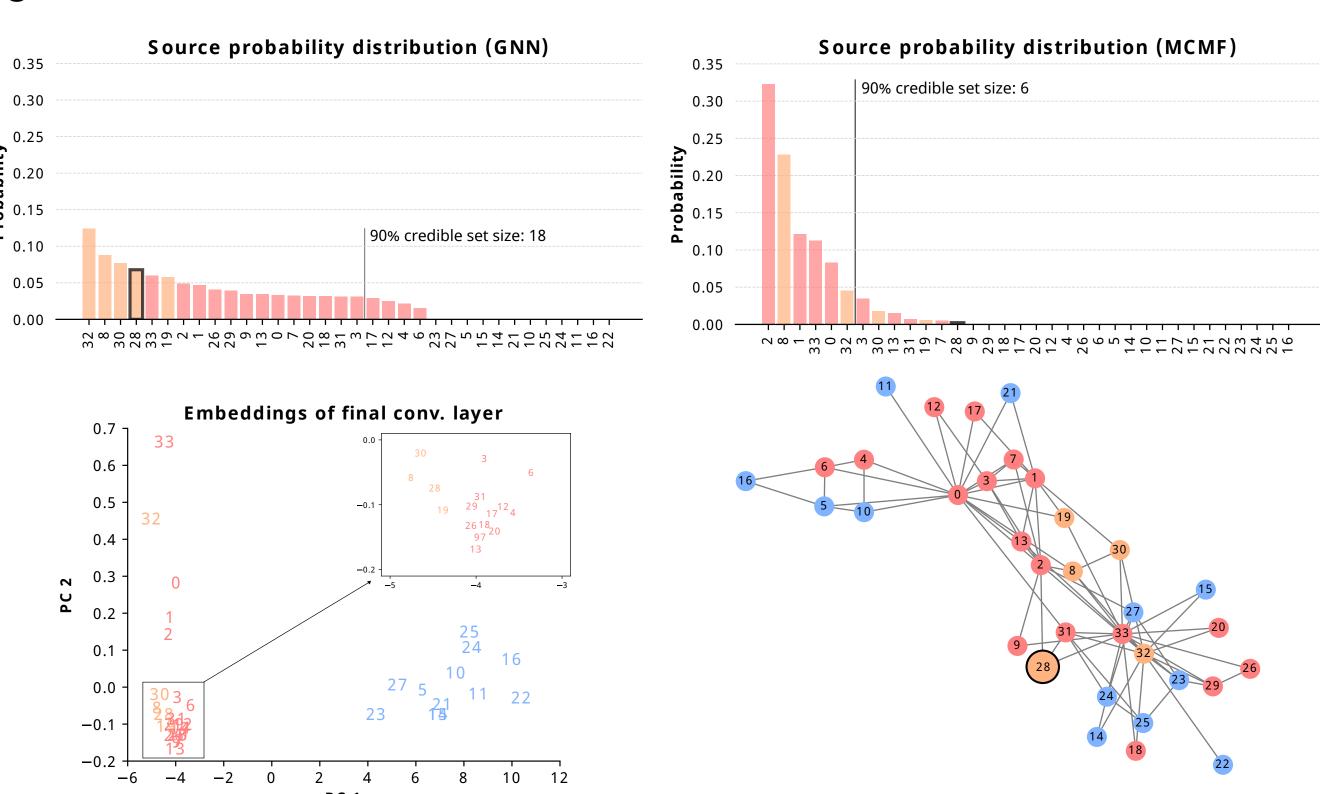


Figure 5. Top left: Output distribution of GNN. Top right: Output distribution of MCMF. Bottom left: First two principal component scores of the final convolutional layer embeddings of GNN. **Bottom** right: Karate network with nodes colored according to epidemic state. Node 28 is the true source.

## **Conclusions**

GNNs perform on par with, or slightly better than, traditional methods. However, their full potential may remain untapped, as the GNN design space has yet to be systematically explored. A key drawback is that GNNs tend to produce larger credible sets.

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