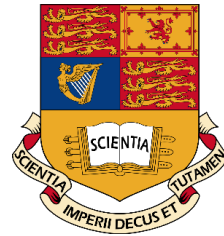


Rumour Source Detection in Social Networks using Partial Observations

Roxana Alexandru and Pier Luigi Dragotti

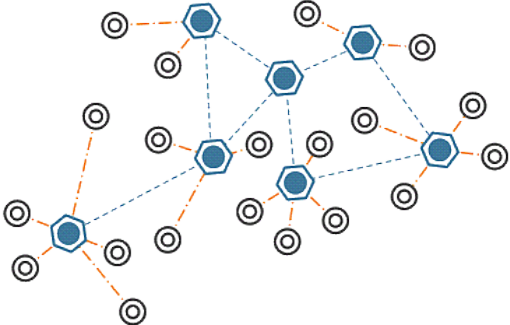
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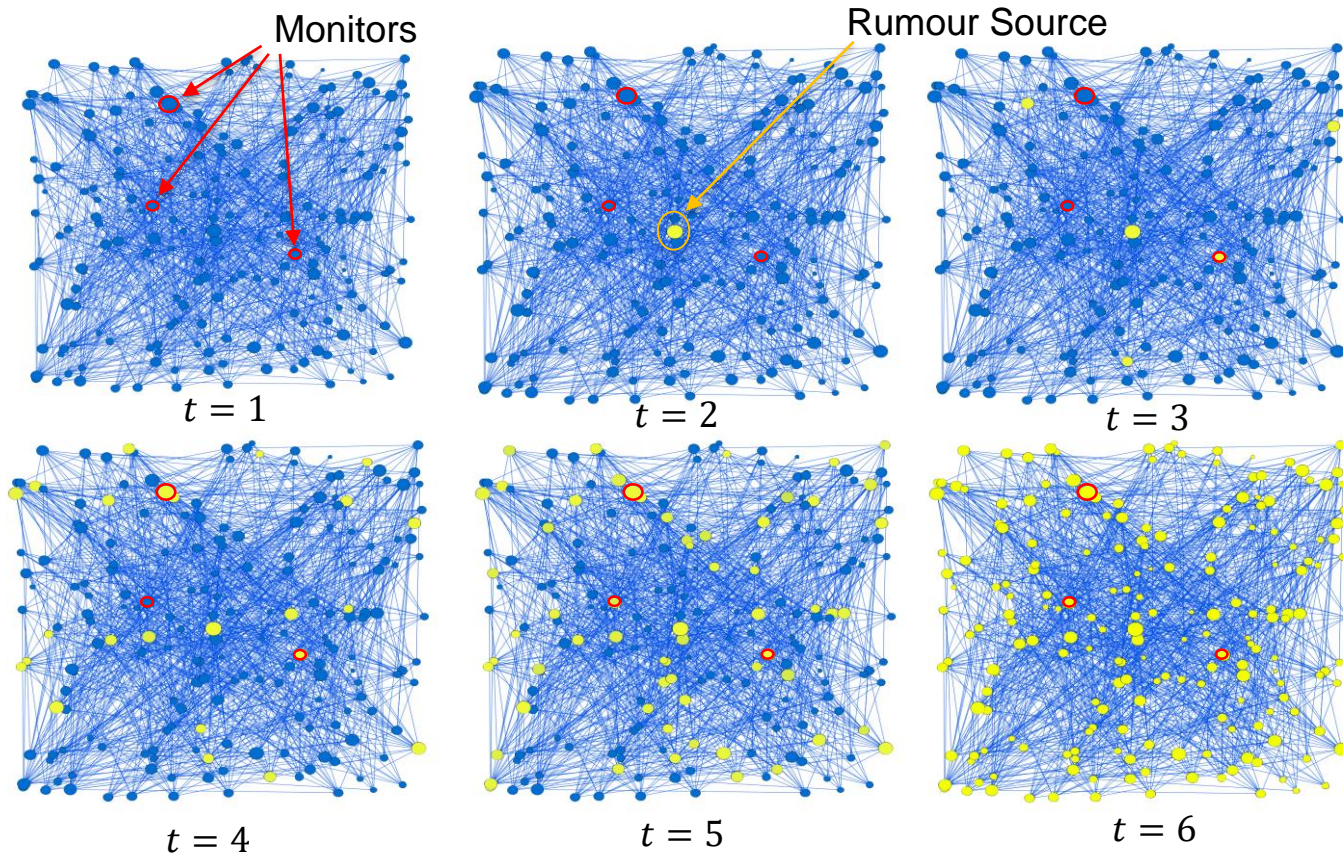
Content

- Motivation
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 - Mathematical Models of Diffusion
 - Single Diffusion Source Detection Algorithm
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-

Motivation



Problem Statement



Problem Statement and Assumptions

Network topology

- General graph with small-world property.

Epidemic model

- Discrete-time version of susceptible-infected model.
- Constant transmission rate within the network.

Observation model

- Known graph topology.
 - Monitoring of a small fraction of nodes.
-

Problem Statement and Assumptions

Source localisation problem

- A source emits R rumours, at $t_0 = 0$.
- We observe some monitors, at discrete times $t \in \{0, 1, \dots, T\}$.
- The probability of infection of a monitor i at time t is given by:

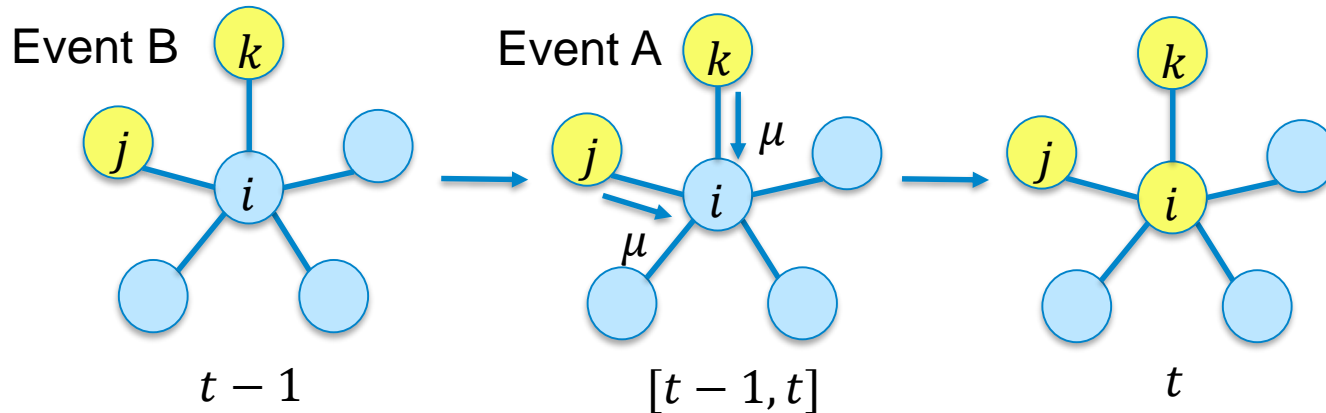
$$\tilde{F}_i(t) = \frac{R_i(t)}{R},$$

where $R_i(t)$ is the number of rumours which have reached i by time t .

- We aim to leverage the divergence of the monitor measurements from an analytical probability of infection.

Approach I to Model Diffusion in a Network

What is the probability a node i gets first infected at time t , $f_i(t)$?



$$f_i(t) = P(A \cap B) = P(A|B)P(B)$$

μ is the constant transmission rate

Derivation in spirit with the methods presented in:

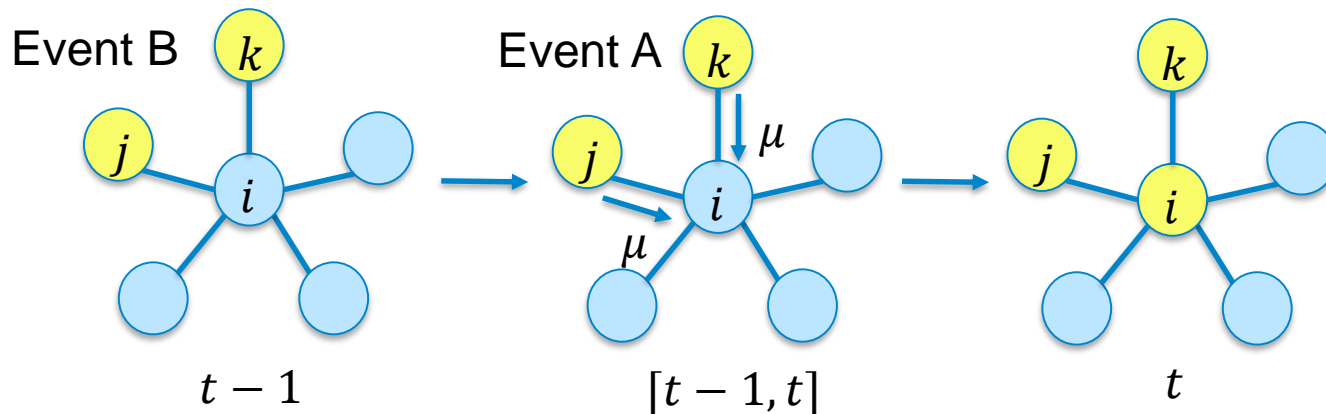
[1] M. Gomez-Rodriguez, D. Balduzzi, B. Schölkopf. *Uncovering the Temporal Dynamics of Diffusion Networks*.

[2] A. Lokhov, M. Mézard, H. Ohta, L. Zdeborová. *Inferring the origin of an epidemic with a dynamic message-passing algorithm*.

[3] N. Ruhi, H. Ahn, B. Hassibi. *Analysis of Exact and Approximated Epidemic Models over Complex Networks*.

Approach I to Model Diffusion in a Network

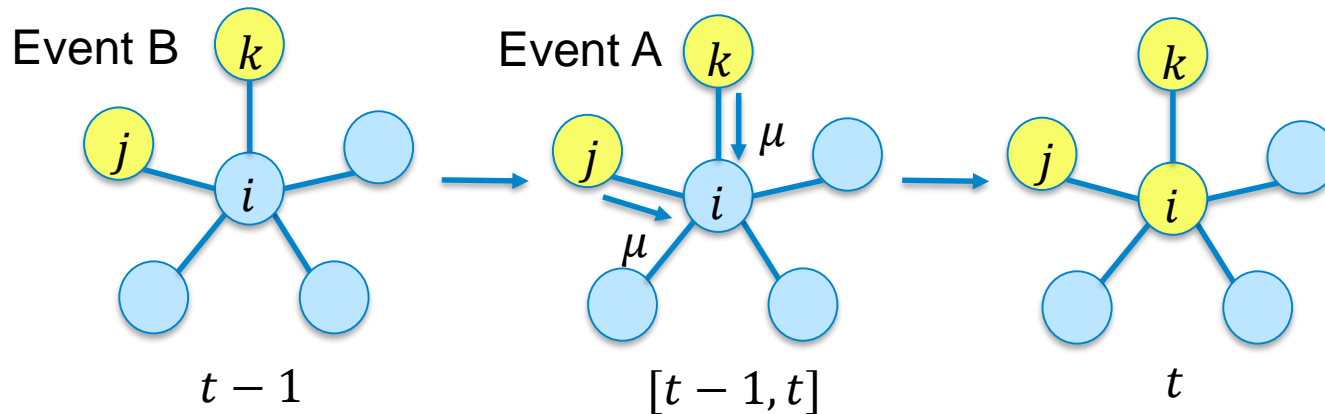
What is the probability a node i gets first infected at time t , $f_i(t)$?



B is the event of node i being in a susceptible state at time $t - 1$:

$$P(B) = \prod_{\tau=1}^{t-1} (1 - f_i(\tau))$$

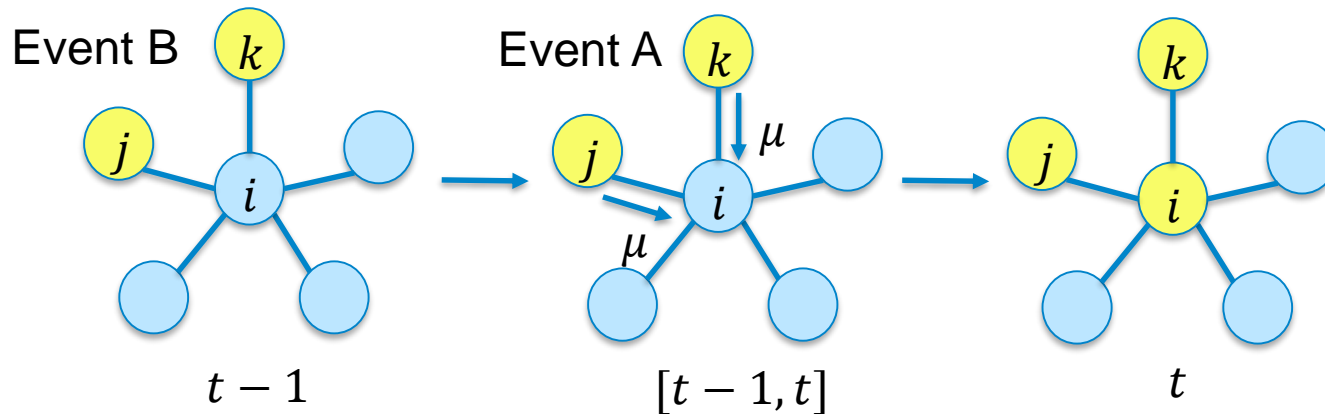
Approach I to Model Diffusion in a Network



$$P(A) = 1 - \prod_{j \in N_i} [1 - \underbrace{\mu \times F(x_j(t-1) = 1)}_{\text{neighbour } j \text{ infected}}]_{\text{neighbour } j \text{ does not transmit}}$$

none of neighbours transmit

Approach I to Model Diffusion in a Network

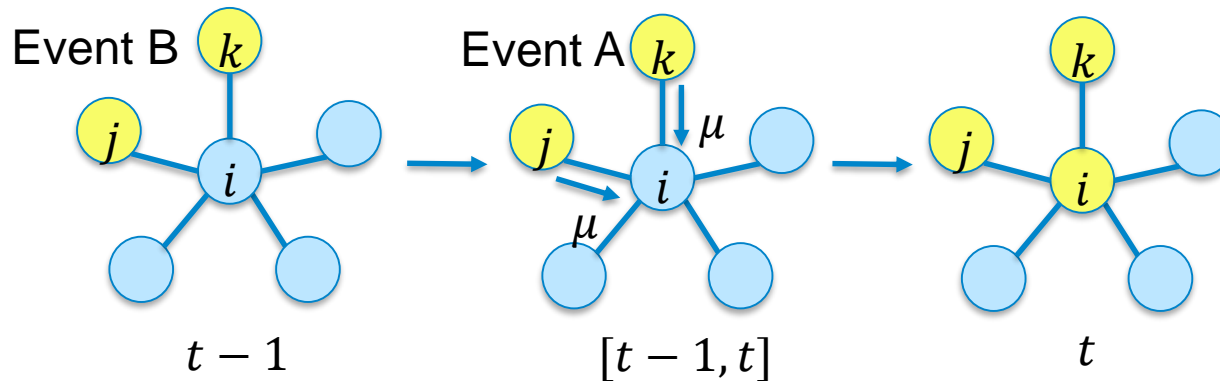


$$P(A) = 1 - \prod_{j \in N_i} [1 - \mu \times F(x_j(t-1) = 1)]$$

The probability i gets the rumour from at least one neighbour, given i was previously in a susceptible state is:

$$P(A|B) = 1 - \prod_{j \in N_i} [1 - \mu \times F(x_j(t-1) = 1 | x_i(t-1) = \mathbf{0})]$$

Approach I to Model Diffusion in a Network



The probability a node i gets first infected at time t , $f_i(t)$ is:

$$f_i(t) = \underbrace{\left[1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1 | x_i(t-1) = 0))\right]}_{P(A|B)} \times \underbrace{\prod_{\tau=1}^{t-1} (1 - f_i(\tau))}_{P(B)}$$

Approach I to Model Diffusion in a Network

- The probability a node i gets first infected at time t is:

$$f_i(t) = \underbrace{\left[1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1 | x_i(t-1) = 0))\right]}_{P(A|B)} \times \underbrace{\prod_{\tau=1}^{t-1} (1 - f_i(\tau))}_{P(B)}$$

- We make the approximation:

$$f_i(t) \approx \left[1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1))\right] \times \prod_{\tau=1}^{t-1} (1 - f_i(\tau))$$

- The approximate probability a node i is infected at time t is:

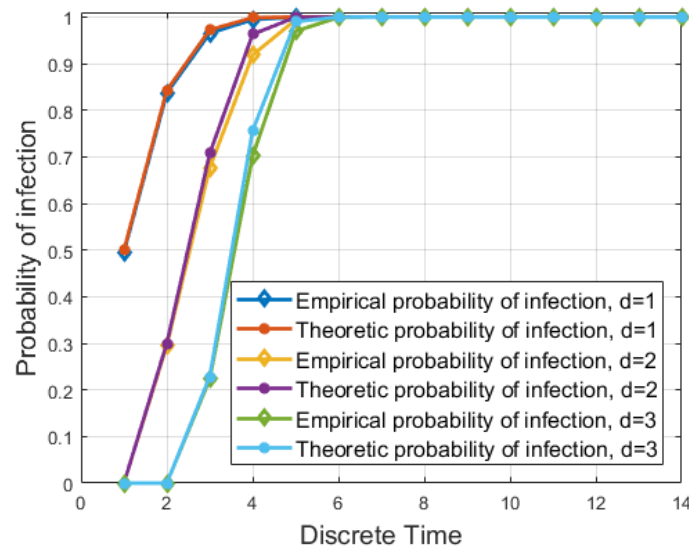
$$F_i(\tau) \approx \sum_{t=1}^{\tau} \left[1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1))\right] \times \prod_{\theta=1}^{t-1} (1 - f_i(\theta))$$

Approach I to Model Diffusion in a Network

- The approximate probability a node i is infected at time t is:

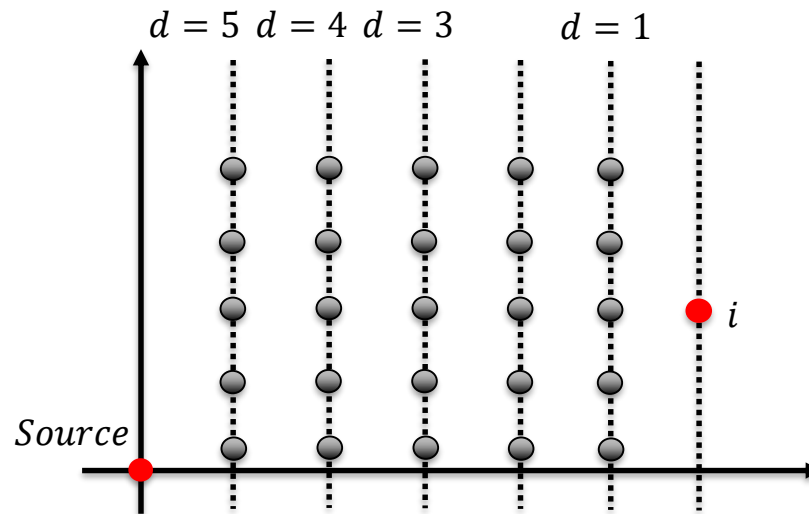
$$F_i(\tau) \approx \sum_{t=1}^{\tau} \left[1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1)) \right] \times \prod_{\theta=1}^{t-1} (1 - f_i(\theta))$$

- Spreading of 1000 Rumors, small-world network, 200 Nodes, for distances 1, 2, and 3 from the source:



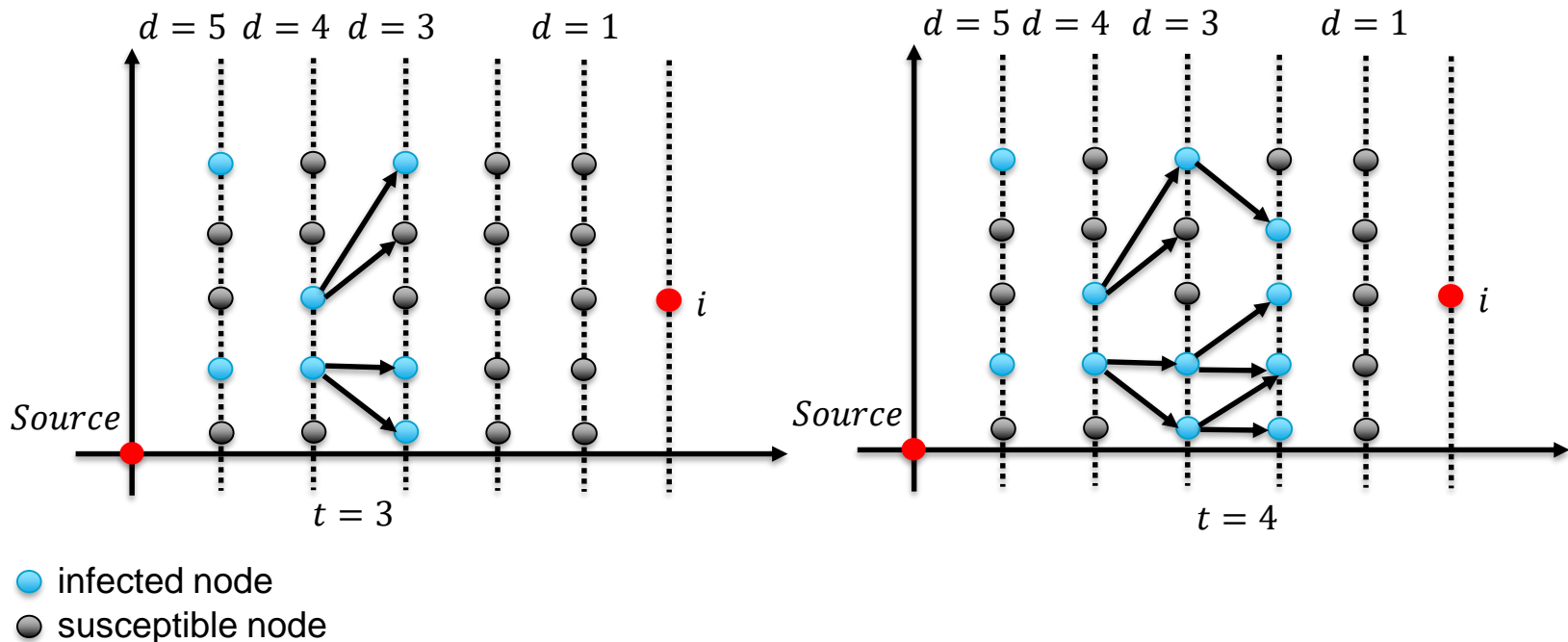
Approach II to Model Diffusion in a Network

- Probability of infection based on the shortest distance to the source.
- Arrange the nodes according to the shortest distance to the destination.
- What is the probability of first infection of a node i at distance d , at time t ?



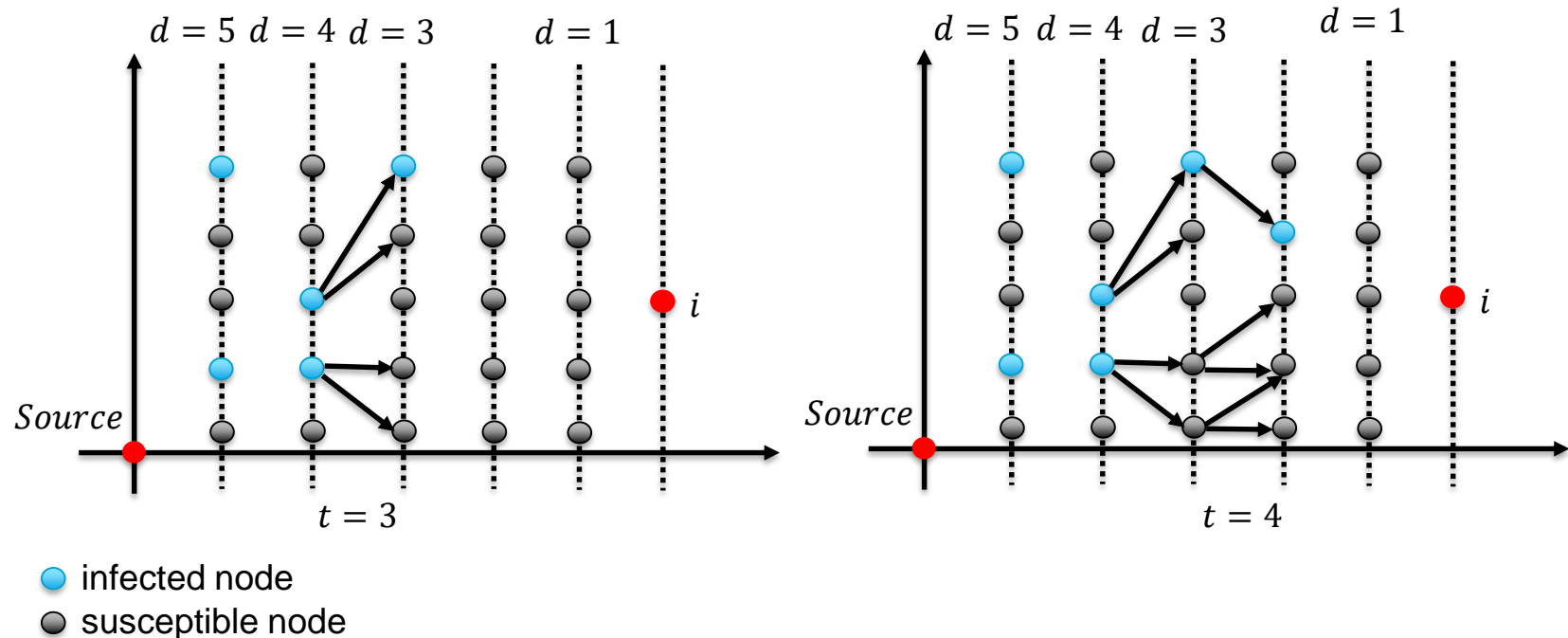
Approach II to Model Diffusion in a Network

- What is the probability of first infection of a node i at distance d , at time t ?
- Success: move closer to node i .



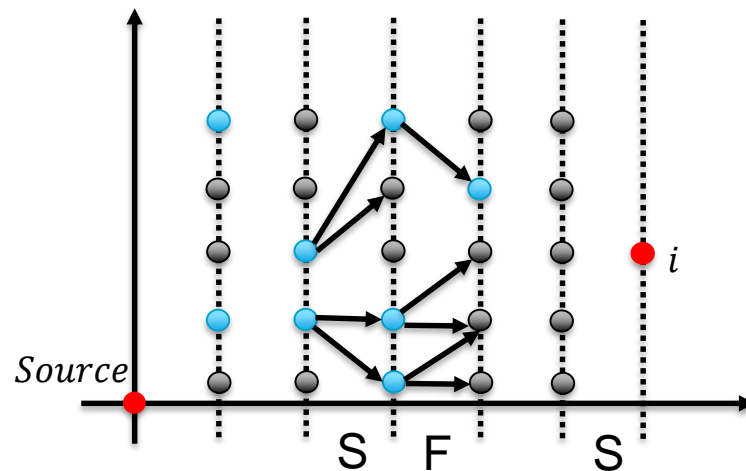
Approach II to Model Diffusion in a Network

- What is the probability of infection of a node i at distance d , at time t ?
- Failure: not spreading the rumour to a sufficient number of nodes closer to the destination.



Approach II to Model Diffusion in a Network

- Number of paths is the number of ways to choose $d - 1$ success steps of $t - 1$ time steps: $\binom{t-1}{d-1}$.
- Probability of success: p_S .
- Probability of each path: $p_S^d \times (1 - p_S)^{t-d}$.
- Approximate $p_S = \alpha_d \mu$, where μ is the constant transmission rate in the graph.



Approach II to Model Diffusion in a Network

- Number of paths is: $\binom{t-1}{d-1}$.
- Probability of each path: $p_S^d \times (1 - p_S)^{t-d}$.
- Set $p_S = \alpha_d \mu$, where μ is the constant transmission rate in the graph.
- The probability of first infection is:

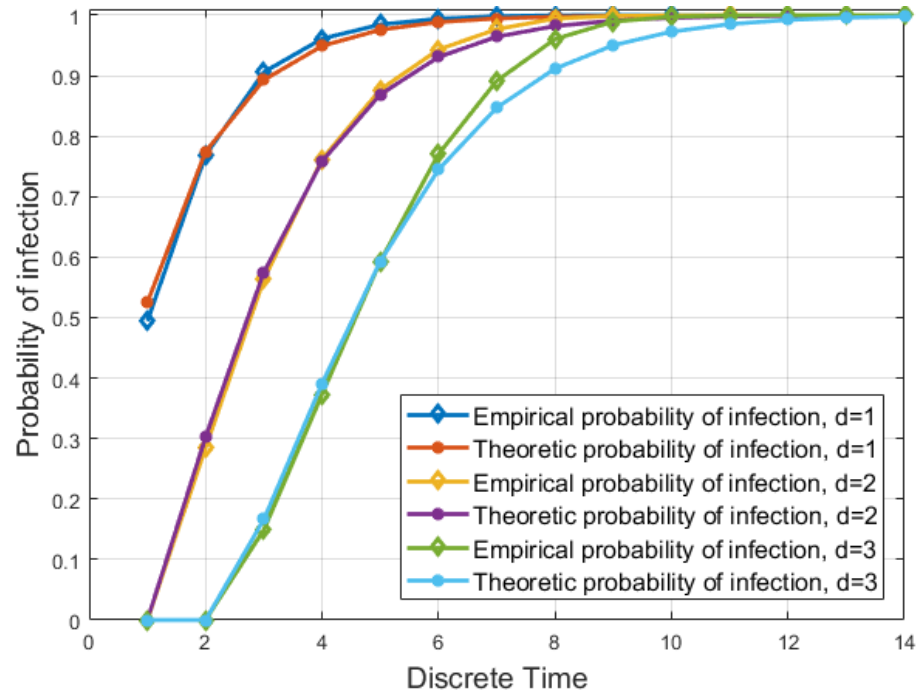
$$f_d(t) = \underbrace{(\alpha_d \mu)^d}_{p_S} \times (1 - \alpha_d \mu)^{t-d} \times \underbrace{\binom{t-1}{d-1}}_{\text{\# of paths from source to destination}}$$

- The probability of infection of a node at distance d from the source at time τ is:

$$F_d(\tau) \approx \sum_{t=d}^{\tau} (\alpha_d \mu)^d \times (1 - \alpha_d \mu)^{t-d} \times \binom{t-1}{d-1}$$

Approach II to Model Diffusion in a Network

- 1000 Rumors, small-world network, 200 Nodes:



Single Diffusion Source Detection Algorithm

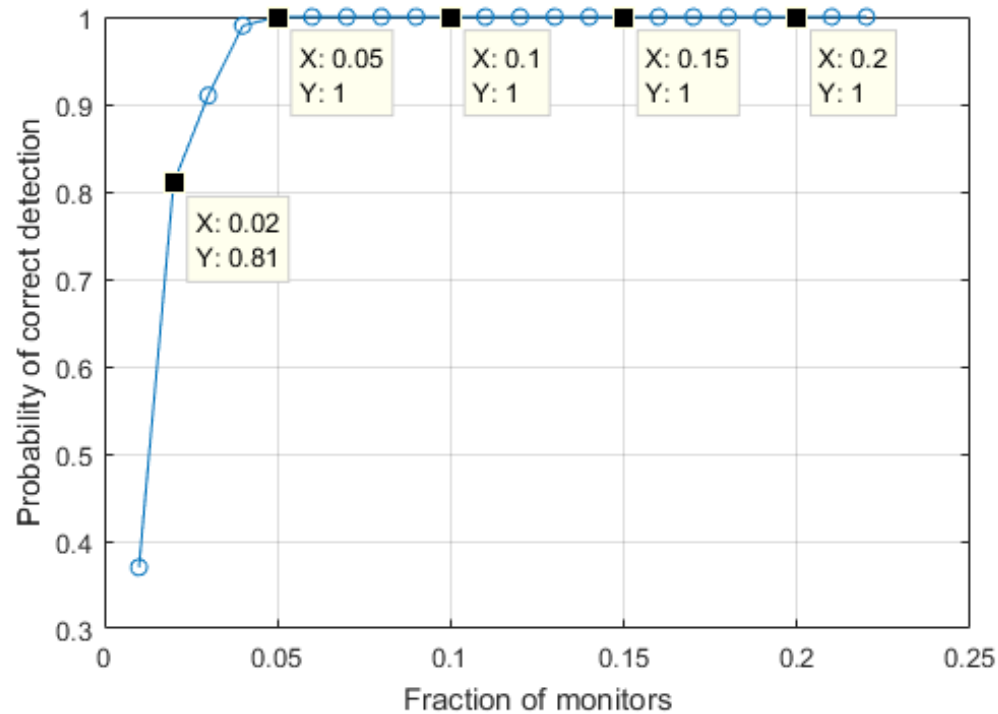
- **Estimate the distances** between each monitor i and the potential source, by computing the dissimilarity between the observed $\tilde{F}_i(t)$ and the theoretical $F_d(t)$.
- Create a set of potential sources using **triangulation**.
- Select the most likely rumour origin, using the approximate model of infection, given a rumour source s :

$$F(x_i(\tau) = 1|s) \approx \sum_{t=1}^{\tau} [1 - \prod_{j \in N_i} (1 - \mu \times F(x_j(t-1) = 1))] \times \prod_{\theta=1}^{t-1} (1 - f_i(\theta))$$

- For each potential source s , compute the dissimilarity between empirical $\tilde{F}_i(t)$ and analytical $F(x_i(T) = 1|s)$. The **most likely rumour origin** is the node with the **lowest dissimilarity**.

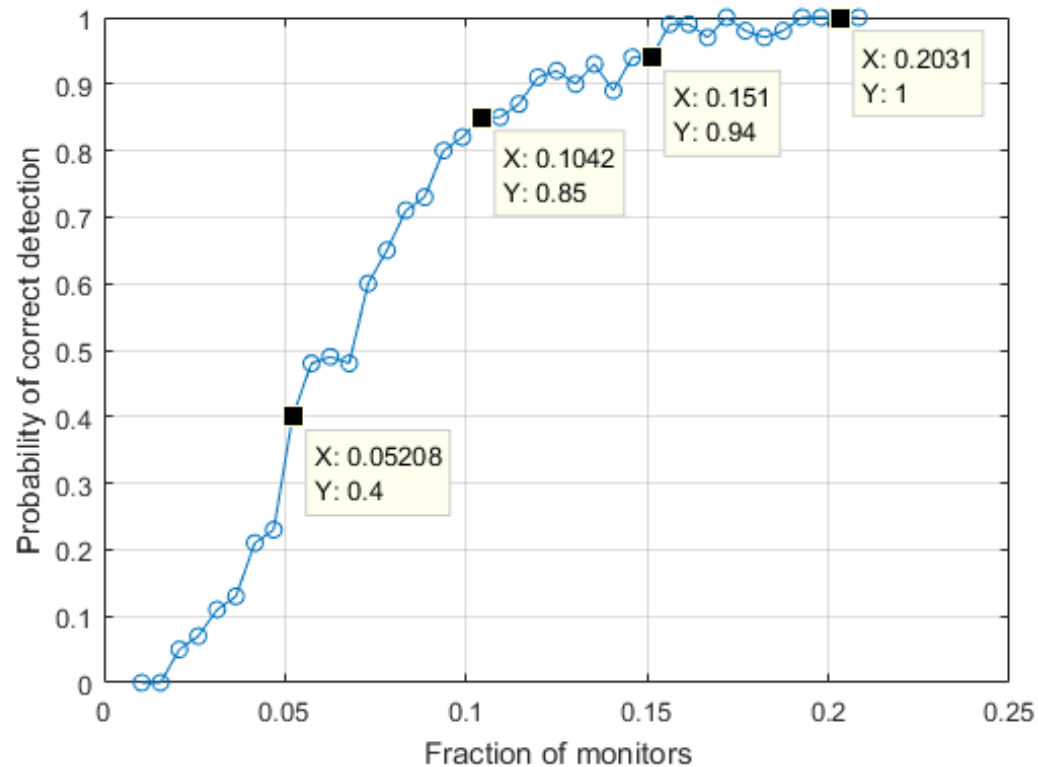
Simulations

- 10 Rumors, small-world network, 1000 Nodes, $\mu = 0.5$, 100 experiments.



Simulations

- 10 Rumors, Facebook network, 192 Nodes, $\mu = 0.5$, 100 experiments.



Conclusion

- Mathematical models of information propagation, which accurately capture the diffusion process.
- Source detection algorithm, which assumes:
 - Single source, which emits multiple rumours.
 - All rumours start at the same time, which is known.
 - A finite set of monitor nodes is observed at discrete times.
- Future extensions:
 - Source detection with unknown start time.
 - Multiple source detection algorithm.

Thank you for listening!

How do we find the optimal parameters α_d in the distance-dependent probabilities?

- The distance-dependent probability of infection for a node at distance d , at time t is:

$$F_d(t) = \sum_{\tau=d}^t (\mu \times \alpha_d)^\tau \times (1 - \mu \times \alpha_d)^{t-\tau} \times \binom{t-1}{\tau-1}$$

- Artificially spread a number of rumours from a random node in the network, and obtain the empirical probabilities $\tilde{F}_i(t)$.
- The optimal parameter α_d minimizes the dissimilarity between $F_d(t)$ and $\tilde{F}_i(t)$ for a particular distance d :

$$\alpha_d^{opt} = \operatorname{argmin}_{\alpha_d} \sum_{i \in N_d} \sum_{t=0}^T \|F_d(t) - \tilde{F}_i(t)\|^2,$$

where N_d is the set of nodes at shortest distance d from the source.

How do we estimate the shortest distances between monitor nodes and the source?

- We find the dissimilarity between the distance-dependent analytical probability of infection $F_d(t)$, and the observed infection probability at a node i , using mean-squared error.

- Then, the optimal distance for a monitor i is:

$$d_{i,s} = \operatorname{argmin}_d \sum_{t=0}^T \|F_d(t) - \tilde{F}_i(t)\|^2$$

- We select as potential sources all the nodes at distance $d_{i,s}$ from node i .

How do we select the most likely rumour origin?

- Select the most likely rumour origin, using the approximate model of infection, given a rumour source s :

$$F(x_i(T) = 1|s) = \sum_{t=1}^T [1 - \prod_{j \in N_i} 1 - \mu \times F(x_j(t-1) = 1)] \times \prod_{\tau=1}^{t-1} (1 - f_i(\tau))$$

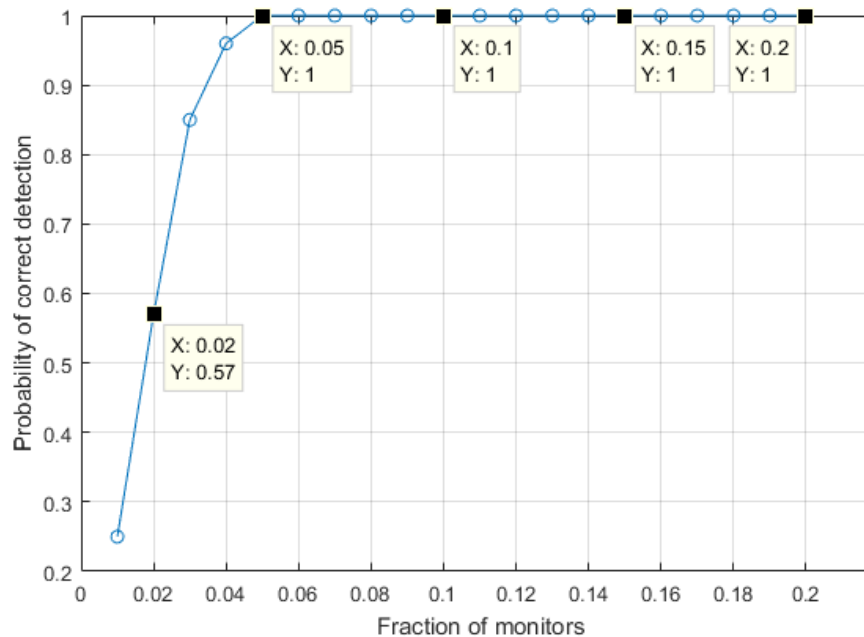
- For each potential source s , compute the dissimilarity between the observed infection probabilities of all monitors, and the theoretic model of infection:

$$\bar{C}(s) = \sum_i \sum_{t=0}^T \|F(x_i(t) = 1|s) - \tilde{F}_i(t)\|^2$$

- The most likely rumour origin is the node with the lowest dissimilarity.

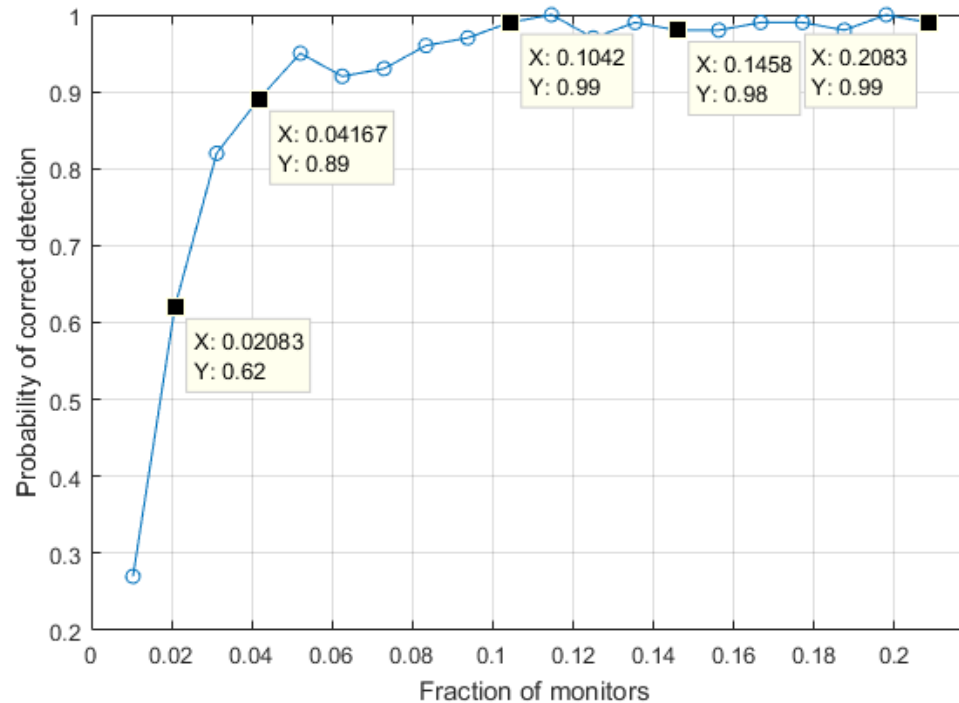
More simulation results

- 10 Rumors, small-world network, 1000 Nodes, *varying* spreading probability



More simulation results

- 10 Rumors, Facebook network, 192 Nodes, *varying* spreading probability



Probability of infection

- A node has the infection at time t if it got initially infected at any of the times before, $\tau = 1, 2, \dots, t$.
- The events of a node getting the initial infection at different times are mutually disjoint.
- Hence, the probability of infection is given by the sum of the likelihoods of first infection at different discrete times:

$$F_i(t) = \sum_{\tau=1}^t f_i(\tau)$$

Probability of being susceptible

- A node is susceptible at time t if it didn't get infected at any of the times before, $\tau = 1, 2, \dots, t$.
- The events of a node not getting the initial infection at different times are mutually disjoint.
- Hence:

$$\bar{F}_i(t) = \prod_{\tau=1}^t 1 - f_i(\tau)$$

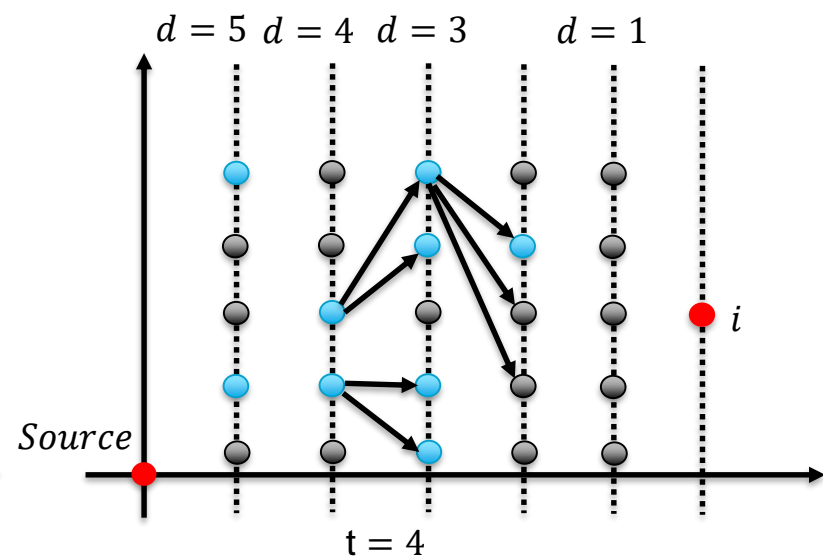
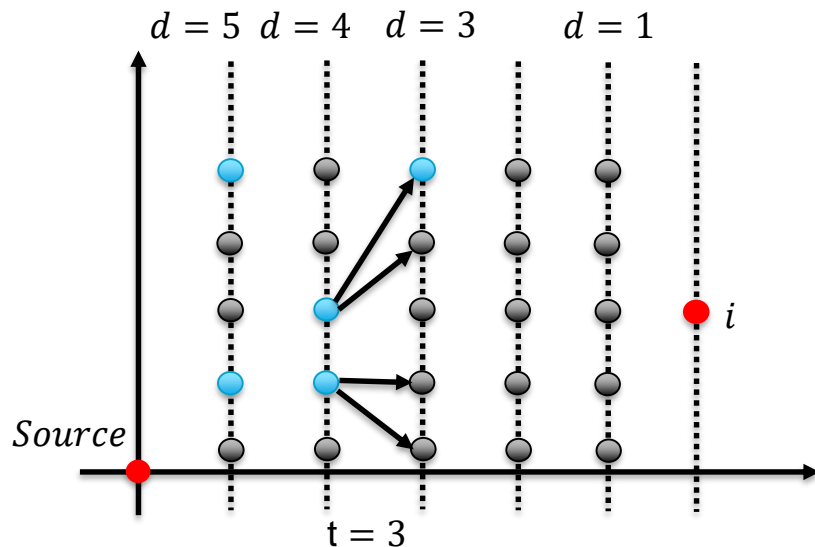
Distance-dependent probability of infection

- Number of paths is: $\binom{t-1}{d-1}$.
- Probability of each path: $p_S^d \times (1 - p_S)^{t-d}$.
- A node at distance d gets infected if *any* succession of d success steps, and $t - d$ failure steps happens.
- Different successions of S and F events are mutually disjoint.
- Hence, the probability of first infection is:

$$f_d(t) = \underbrace{(\alpha_d \mu)^d}_{p_S} \times (1 - \alpha_d \mu)^{t-d} \times \underbrace{\binom{t-1}{d-1}}_{\text{\# of paths from source to destination}}$$

Distance-dependent probability of infection

- A node at distance d gets infected if *any* succession of d success steps, and $t - d$ failure steps happens.
- There can be a success following a failure, at the next time step.



Comparison to existing methods

- The authors in [1] propose a Monte Carlo method for single source estimation, with unknown infection time. In a **random geometric graph**, the probability of the origin to be within the first **10% ranked nodes** is around **0.5** when **observing 5%** of the network, increasing to **0.9** when **observing the full network**.
- In a **small-world network**, our method achieves correct detection probability of **0.75** when **observing 5%** of the network, and **1** when **observing the full network**. The number of **rumours is 2**, and the rumour **start time is known**.