Influence Prediction on Networks

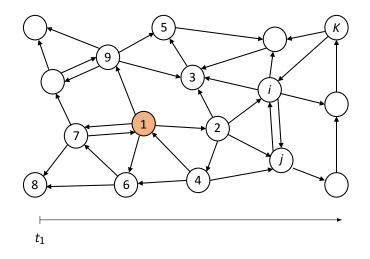
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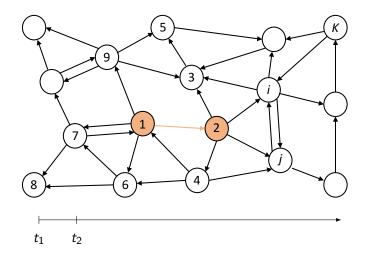
Joint work with Shui-Nee Chow (GT Math), Hongyuan Zha (GT CSE), Haomin Zhou (GT Math)

SIAM CSE'17, 02/27/17

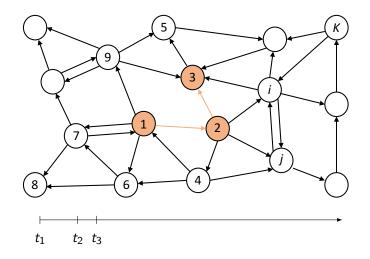
Research supported in part by NSF grant DMS-1620342



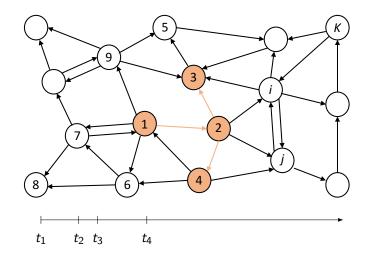
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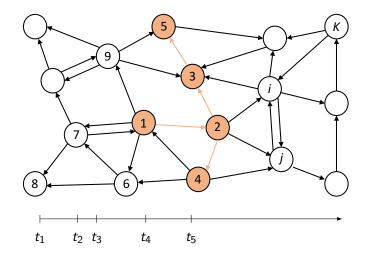
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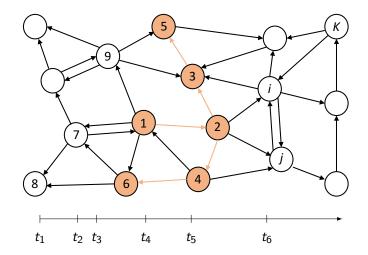
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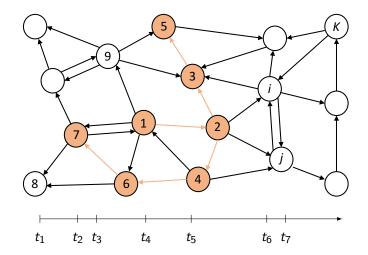
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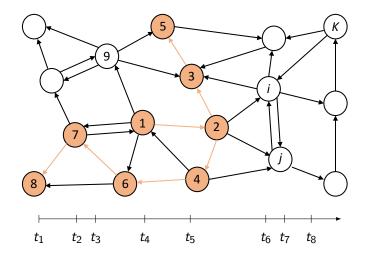
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Problem description

Propagation network:

- G = (V, E) network (directed graph)
- $S \subset V$ source set
- { α_{ij} : (*i*, *j*) \in *E*}: $t_{ij} = t_j t_i \sim \text{Exp}(\alpha_{ij})$

Then information propagates by gradually activating more nodes.

Definition (Influence)

Given S, the expected number of activated nodes at time t is called the influence of S, denoted by $\mu(t; S)$.

Influence prediction

Question:

Given S, how to compute influence $\mu(t; S)$ for all t?



Influence prediction has many applications

▶ Influence maximization: fix t and $n \in \mathbb{N}$, solve

$$\underset{S \subset V}{\text{maximize }} \mu(t; S) \quad \text{ s.t. } |S| \le n$$

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- Outbreak detection
- Propagation control

Exact solution? Not tractable.

Exact solution requires working in a state space of size $O(2^{K})$.

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N(t) and its transition states

From now on, since S is arbitrary and fixed, we drop it for notation simplicity.

Let N(t) be the (random) number of activated nodes in G, and M_k be the state that N(t) = k. Then

$$\underline{M_0} \rightleftharpoons \cdots \rightleftharpoons \boxed{M_{k-1}} \stackrel{q_{k-1}(t)}{\underset{r_k(t)}{\rightleftharpoons}} \underbrace{M_k} \stackrel{q_k(t)}{\underset{r_{k+1}(t)}{\boxtimes}} \underbrace{M_{k+1}}_{r_{k+1}(t)} \rightleftharpoons \cdots \rightleftharpoons \underbrace{M_K}$$

where $q_k(t)$ is the transition rate from M_k to M_{k+1} , and $r_k(t)$ is the transition rate from M_k to M_{k-1} at time t.

Key quantities

Number of activated nodes:

N(t)

Probability that N(t) is in state M_k :

$$\rho_k(t) = \Pr(N(t) = k)$$

Influence (i.e., expected number of activated nodes):

$$\mu(t) = \mathbb{E}[N(t)] = \sum_{k=0}^{K} k \rho_k(t)$$

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Note the key is to compute $\{\rho_k(t)\}!$

Fokker-Planck equation

Recall the state transition graph:

$$\boxed{M_0} \rightleftharpoons \cdots \rightleftharpoons \boxed{M_{k-1}} \stackrel{q_{k-1}(t)}{\rightleftharpoons} \underbrace{M_k}_{r_k(t)} \stackrel{q_k(t)}{\boxtimes} \underbrace{M_{k+1}}_{r_{k+1}(t)} \boxtimes \cdots \rightleftharpoons \underbrace{M_K}$$

The Fokker-Planck equation is a system of deterministic differential equations that governs the time evolution of $\rho_k(t)$:

$$\begin{split} \rho_0'(t) &= -q_0(t)\rho_0(t) + r_1(t)\rho_1(t), \\ \rho_k'(t) &= q_{k-1}(t)\rho_{k-1}(t) - [q_k(t) + r_k(t)]\rho_k(t) \\ &+ r_{k+1}(t)\rho_{k+1}(t), \quad \text{for } 1 \le k \le K - 1, \\ \rho_K'(t) &= q_{K-1}(t)\rho_{K-1}(t) - r_K(t)\rho_K(t). \end{split}$$

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Matrix formulation

The matrix form of the Fokker-Planck equation above is

$$\rho'(t) = \rho(t)[Q(t) + R(t)]$$

where $\rho(t) = (\rho_0(t), \rho_1(t), \dots, \rho_K(t)) \in \mathbb{R}^{K+1}$ is a row vector, and Q(t) is a *bidiagonal* matrix:

and R(t) is a (lower) bidiagonal matrix with $r_k(t)$'s.

Composition of Q and R

Theorem Let $S_k := \{U \subset V : |U| = k\}$ and Pr(t; U) be the probability that $U \in S_k$ is activated first. Define

$$\alpha(U) = \sum_{i \in U} \sum_{j \in N_i^{out} \cap U^c} \alpha_{ij}, \quad \beta(U) = \sum_{i \in U} \beta_i, \quad \gamma(U) = \sum_{i \in U} \gamma_i$$

Similarly $\beta(U) = \sum_{i \in U} \beta_i$ and $\gamma(U) = \sum_{i \in U} \gamma_i$. Then there are

$$egin{aligned} q_k(t) &= \sum_{U \in \mathcal{S}_k} \left[lpha(U) + eta(U^c)
ight] \Pr(t; U) \ r_k(t) &= \sum_{U \in \mathcal{S}_k} \gamma(U) \Pr(t; U) \end{aligned}$$

for k = 0, 1, ..., K.

Estimate q_k

We assume no self-activation and recovery, and provide two ways to estimate q_k :

▶ Based on shortest distance (FPE-dist): Define the distance from i to j by 1/α_{ij}, let U^{*}_k ∈ S_k pick the k nodes with shortest distance to S, and set

$$\hat{q}_k = lpha(U_k^*)$$

Based on overall probability (FPE-tree): For k = 1, 2, ..., recursively find {U_k¹, ..., U_k^{m_k}} ⊂ S_k with large probabilities in S_k, which essentially constructs a tree of nodes {U_k^l} with relative probabilities in each layer k. Set

$$\hat{q}_k = \sum_{l=1}^{m_k} lpha(U_k^l) \Pr(U_k^l)$$

Experiment setup

Generating propagation networks:

- Various types of networks (directed graphs): Erdős-Rényi's random, small-world, scale-free, Kronecker, etc.
- ▶ Various sizes *K* and densities (average node out-degree).
- For each edge $(i, j) \in E$, draw $\alpha_{ij} \stackrel{i.i.d.}{\sim} Unif(0, 1)$.

Ground truth by MCMC:

Obtained by simulating 5000 cascades and calculating average number of activated nodes. (expensive!)

Experimental results: small networks

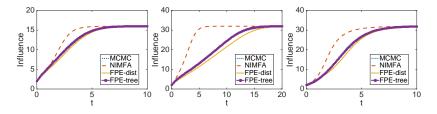


Figure: Left: Erdős-Rényi's network (K = 16, $d^{avg} = 4$). Middle: Erdős-Rényi's network (K = 32, $d^{avg} = 4$). Right: Small-world network (K = 32, $d^{avg} = 4$). Here $d^{avg} = (1/K) \sum_{i} |N_i^{out}|$.

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Experimental results: large networks

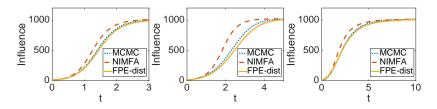


Figure: Left: Erdős-Rényi's network (K = 1024, $d^{avg} = 8$). Middle: Small-world network (K = 1024, $d^{avg} = 6$). Right: Scale-free network (K = 1024, $d^{avg} = 6$).

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More experimental results

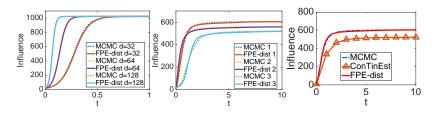


Figure: Left: Dense Erdős-Rényi's random network (K = 1024 and $d^{avg} = 32, 64, 128$ respectively). Middle: Influence prediction on the same Kronecker network of size 1024 using three different choices of source set S_1, S_2, S_3 ($|S_i| = 10$). Right: Comparison with ConTinEst, a state-of-the-art method that learns coverage function using sample cascades.

The estimation of Fokker-Planck equation coefficients q_k seems crude, but why the performance is so good?

We answer this question by building relationship between error in $q_k(t)$ and error in $\mu(t)$.

Error analysis

Lemma

Let $\epsilon \in (0, 1)$, and ρ and $\hat{\rho}$ solve $\rho'(t) = \rho(t)Q_{k+1}(t)$ and $\hat{\rho}'(t) = \hat{\rho}(t)Q_k(t)$ respectively, where Q_k has q_j in Q replaced by \hat{q}_j for $j \ge k$. If every \hat{q}_k satisfies

$$\frac{|\hat{q}_k(t) - q_k(t)|}{q_k(t)} \le \min\left\{\frac{\log(1 + \frac{\epsilon}{2})}{\bar{\alpha}kt\min(\bar{d}, K - k)}, \frac{\epsilon}{2 + \epsilon}\right\}$$

where $\bar{\alpha} = \max\{\alpha_{ij} : (i,j) \in E\}$, $\bar{d} = \max\{|N_i^{out}| : i \in V\}$, then

$$egin{aligned} \hat{
ho}_j(t) &=
ho_j(t), \ ext{for } j = 0, \dots, k-1 \ &|\hat{
ho}_j(t) -
ho_j(t)| /
ho_j(t) &\leq \epsilon, \ ext{for } j = k, \dots, K-1 \ &|\hat{\mu}(t) - \mu(t)| / \mu(t) \leq \epsilon \end{aligned}$$

Error analysis

Theorem

Let $\epsilon \in (0, 1)$, and $\rho(t)$ and $\hat{\rho}(t)$ solve $\rho'(t) = \rho(t)Q(t)$ and $\hat{\rho}'(t) = \hat{\rho}(t)\hat{Q}(t)$ respectively, where \hat{Q} has q_k in Q replaced by \hat{q}_k for all k. If every q_k satisifies

$$\frac{|\hat{q}_k(t) - q_k(t)|}{q_k(t)} \le \min\left\{\frac{\log(1 + \frac{\epsilon}{2})}{\bar{\alpha}kt\min(\bar{d}, K - k)}, \frac{\epsilon}{2 + \epsilon}\right\}$$

and let $c_{K}(t) := \frac{1}{K} \sum_{j=0}^{K-1} \frac{K-j}{j!} (\bar{q}t)^{j}$ where $\bar{q} := \max_{k} \{q_k\}$, then

$$\frac{|\hat{\mu}(t)-\mu(t)|}{\mu(t)} \leq \left[(1+\epsilon)^{\mathcal{K}}-1\right]\min\left\{1,c_{\mathcal{K}}(t)e^{-\underline{\alpha}t}\right\}, \quad \forall t\geq 0,$$

where $\underline{\alpha} := \min\{\alpha_{ij} : (i,j) \in E\}.$

Error analysis

Corollary

Suppose $\rho(t), \hat{\rho}(t), \mu(t), \hat{\mu}(t)$ are defined and conditions for $\bar{\alpha}$ and $\underline{\alpha}$ as above. Let $\varepsilon > 0$ and $c \in (0, \underline{\alpha})$, then

 $|\hat{\mu}(t) - \mu(t)|/\mu(t) \le \varepsilon e^{-ct}$

as long as the estimated $\hat{q}_k(t)$ satisfies

$$\frac{|\hat{q}_k(t) - q_k(t)|}{q_k(t)} \leq \frac{\underline{\alpha} - c}{K\bar{q}_k} + \frac{\log \varepsilon - K\log 2 - \log c_K(t)}{K\bar{q}_k t}$$
$$= C_k - O\left(\log t/t\right)$$

for each k = 0, 1, ..., K - 1, where $\bar{q}_k := \bar{\alpha}k \min\{\bar{d}, K - k\}$ and $C_k := (\underline{\alpha} - c)/K\bar{q}_k$.

Experimental results

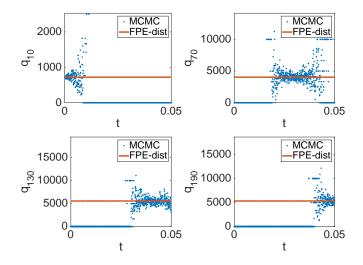


Figure: \hat{q}_k (red) and q_k (blue) for for k = 10, 70, 130, 190 in Erdős-Rényi's network ($K = 300, d^{avg} = 150, \alpha_{ij} \stackrel{i.i.d.}{\sim} \text{Unif}(0, 1)$).

Experimental results

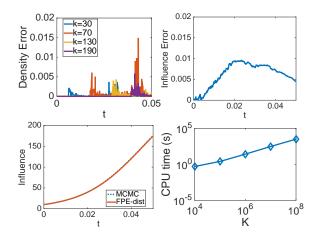


Figure: Upper left: $\frac{|\hat{\rho}_k(t) - \rho_k(t)|}{\rho_k(t)}$ for k = 30, 70, 130, 190. Upper right: $\frac{|\hat{\mu}(t) - \mu(t)|}{\mu(t)}$. Lower left: $\hat{\mu}(t)$ and $\mu(t)$. Lower right: CPU time (in seconds) to solve Fokker-Planck equation for networks with various sizes.

Summary

In this work, we have

- Built a general framework for influence prediction based on time evolutions of ρ_k(t).
- Provided methods to estimate coefficients of the related Fokker-Planck equations.
- Established relationship between coefficient error and prediction error.

Future work

- Non-Markov propagations.
- Prediction directly based on historical cascade data.

and more ...

Thank you

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