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- Overview of the SIS Model
- Ganesh-Massoulié-Towsley: Upper Bound on Epidemic Die-Out

1 Overview of the SIS Model

We studied SIR Model/Branching Process in the previous lecture, today we will introduce SIS (Susceptible-Infective-Susceptible) model. There are three components.

2 SIS Model

- Contact network through which the infection spreads.
- Infection Rate along each edge is β .
- Recovery Rate δ .

Extent & Length of epidemic

We represent the system by a connected graph G(V, E). Let |V| = n, and let the state at time t be represented by a vector $X(t) = (X_1(t), X_2(t), \dots, X_n(t))^T$. X_i is defined as follows:

$$X_i(t) = \begin{cases} 1 & \text{if node } i \text{ is infected at time } t \\ 0 & \text{otherwise} \end{cases}$$

Assume that infected nodes X_i contaminate neighbors as a Poisson process with rate β and recover with a Poisson process with rate δ . This defines a continuous-time Markov process with transition rates:

$$X_i: 0 \to 1 \text{ at rate } \beta \sum_{(i,j) \in E} X_j,$$
$$X_i: 1 \to 0 \text{ at rate } \delta.$$

Without loss of generality, we can assume $\delta = 1$, since it essentially corresponds to a normalization factor for the die-out time.

In other words, given $X_i(t) = 0$ we have $X_i(t + dt) = 1$ with probability $[\beta \sum_{(i,j) \in E} X_j(t)]dt$ for $dt \to 0$.

Note. X_i changes from 1 to 0 at rate r means if it takes time y to go from 1 to 0, then $Pr(y \ge t) = e^{-tr}$ (an exponential distribution with rate r).

DIE-OUT of epidemic

Let τ be the time takes for epidemic to die. Now the question we are interested in is: What is $E(\tau)$, as a function of β and G?

3 Ganesh-Massoulié-Towsley: Upper Bound on Die-Out Time

Let A be the adjacency matrix of G, $\lambda_1(A)$ be the largest eigenvalue of A.

Theorem 1. If $\beta < \frac{1}{\lambda_1(A)}$, then $E(\tau) = O(\log n)$.

Proof: Given $\beta \lambda_1(A) < 1$, at what t does $\sum_i X_i(t) = 0$? To approach this, we consider the function $Pr(\sum X_i(t) > 0)$ and see when this function goes to 0. Recall that,

If $X_i(t) = 1$, then

$$\begin{aligned} X_i(t+dt) &= 0 & \text{with probability } dt \\ X_i(t+dt) &= 1 & \text{with probability } 1 - dt \end{aligned}$$

If $X_i(t) = 0$, then

$$\begin{split} X_i(t+dt) &= 1 \qquad \text{with probability } dt \cdot \beta \cdot \sum_{(i,j) \in E} X_j(t) \\ X_i(t+dt) &= 0 \qquad \text{with probability } 1 - dt \cdot \beta \cdot \sum_{(i,j) \in E} X_j(t) \end{split}$$

The transition of X_i depends on the value of X_i , it is very hard to handle. Now we consider the continuous-time Markov process $Y = \{Y_i\}_{i \in V}$, $Y_i(0) = X_i(0)$. For k > 0, the transition rates

$$Y_i: k \to k+1$$
 at rate $\beta \sum_{(i,j) \in E} Y_j$

 $Y_i: k \to k-1$ at rate Y_i

It is easy to see that $Y_i \in \{0, 1, 2, ...\}$ (compared to $X_i \in \{0, 1\}$) and, when starting from the same initial conditions, Y_i stochastically dominates X_i . This stochastic dominance is obtained by a coupling argument in which we couple the elementary events in the probability space of X_i with those in the probability space of Y_i . We omit the details here.

By stochastic dominance we have $Pr(Y_i(t) \ge y) \ge Pr(X_i(t) \ge y)$. Therefore, we have $Pr(\text{Epidemic not die-out at time t}) = Pr(\sum X_i(t) > 0) \le Pr(\sum Y_i(t) > 0)$.

We have

$$Y_i(t+dt) = \begin{cases} Y_i(t) + 1 & \text{with probability } \beta \cdot \sum_{(i,j) \in E} Y_j(t) \\ Y_i(t) - 1 & \text{with probability } Y_i(t)dt \\ Y_i(t) & \text{otherwise} \end{cases}$$

Note. In the above calculation we are ignoring the simultaneous occurrence of more than one event. This is because for $dt \rightarrow 0$, the associated probabilities are lower order terms (they are super-linear in dt).

Through continuous time $\{Y_i(t+dt) = Y_i(t)+1\}$ and $\{Y_i(t+dt) = Y_i(t)-1\}$ can never happen at the same time.

Since for all i we have

$$E[Y_{i}(t+dt) - Y_{i}(t)] = (\beta \sum_{(i,j)\in E} E[Y_{j}] - E[Y_{i}])dt$$

The transition rates for process Y(t) are such that

$$\frac{d(E(Y(t)))}{dt} = (\beta A - I)E(Y(t))$$

where I denotes the identity matrix. Hence,

$$E(Y(t)) = e^{Mt} \cdot Y(0)$$
 where $M = \beta A - I$.

Note. $e^M = I + M + \frac{M^2}{2!} + \cdots$

Consider the following fact that if A has eigenvalue λ then $M = \beta A - I$ has eigenvalue $\beta \lambda - 1$ and e^M has eigenvalue $e^{\beta \lambda - 1}$. We obtain

$$||E(Y(t))||_2 \le e^{(\beta\lambda_1 - 1)t} \cdot ||E(Y(0))||_2.$$

Note. Cauchy-Schwarz inequality says that

$$|\langle x, y \rangle| \le \|x\|_2 \cdot \|y\|_2$$

We obtain:

$$\sum_{i \in V} E(Y_i(t)) \le \|E(Y(t))\|_2 \cdot \|\mathbf{1}\|_2$$

where **1** denotes the vector of ones, so $\|\mathbf{1}\|_2 = \sqrt{n}$.

Note. $Pr(Y_i > 0) \ge Pr(X_i > 0)$, which implies that $Pr(\sum Y_i > 0) \ge Pr(\sum X_i > 0)$. Moreover, by Markov Inequality it holds that:

$$Pr(\sum Y_i > 0) = Pr(\sum Y_i \ge 1)$$
$$\leq \sum E(Y_i).$$

We have

$$Pr(\sum X_i(t) > 0) \le Pr(\sum Y_i(t) > 0)$$

$$\le \sum E(Y_i(t))$$

$$\le \|E(Y(t))\|_2 \cdot \sqrt{n}$$

$$\le \sqrt{n}e^{(\beta\lambda_1 - 1)t} \cdot \|E(Y(0))\|_2$$

$$< n \cdot e^{(\beta\lambda_1 - 1)t}$$

Note. $Y_i(0) = X_i(0)$, which takes values in $\{0,1\}$. So, $||E(Y(0))||_2 \le \sqrt{n}$. Say $t_1 = \frac{100 \ln n}{1-\beta\lambda_1}$, then we have

 $Pr(\text{epidemic not die-out at time } t_1) = Pr(\sum X_i(t_1) > 0) \le \frac{1}{n^{99}}$ We have for any $t \ge t_1$,

$$Pr(\text{epidemic not die-out at time } t) \leq \frac{1}{n^{99}},$$

since the epidemic die-out is an absorbing state in the Markov process. Therefore,

$$E(\tau) = \int_0^\infty Pr(\tau > t)dt$$

= $\int_0^\infty Pr(\sum X_i(t) > 0)dt$
 $\leq t_1 + \int_{t_1}^\infty Pr(\sum X_i(t) > 0)dt$
 $\leq \frac{100 \ln n}{1 - \beta \lambda_1} + \int_{t_1}^\infty \frac{1}{n^{99}}dt$
= $O(\ln n).$

Theorem 2. If

 $\beta > \frac{c}{\lambda_1(A) - \lambda_2(A)} \quad (where \ c > 0 \ is \ a \ sufficiently \ large \ constant)$ then $E(\tau) = \Omega(\exp(n)).$