

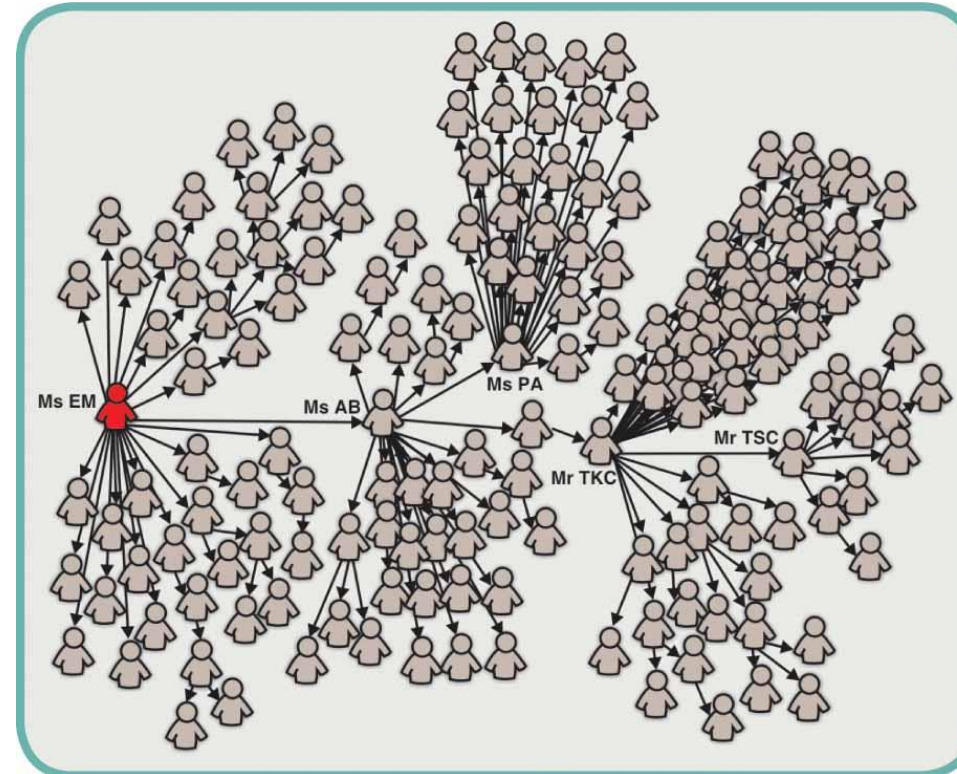
# Learning Networks from Dynamics: Detecting Abrupt Changes in Point Processes

Anna Brandenberger, Elchanan Mossel, Ani Sridhar

Massachusetts Institute of Technology

## Learning a network from diffusions

Motivating example: network epidemics



Contact network of the first 144 cases of a SARS outbreak in Singapore (src: Normile'13)

- Network structure impacts the spread of epidemics in significant and complex ways
- If the network is known, then we can design effective mitigation measures

- Networks are typically learned through contact tracing  $\Rightarrow$  time-consuming
- Can *key features* of the network be learned in a data-driven manner?

## Prior work in learning from dynamics

Exact estimation of networks from diffusions

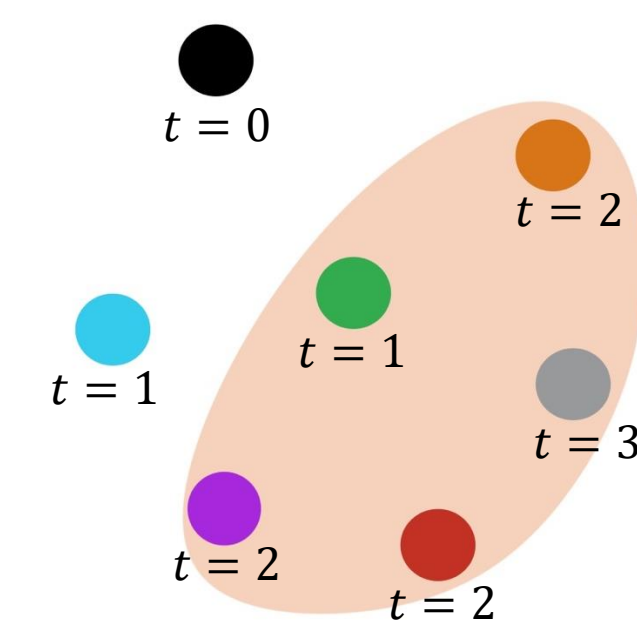
- Early empirical work in epidemiology [Wallinga-Teunis'04] and information flow in blogs [Adar-Adamic'05]
- Scalable and principled methods: the NetInf algorithm [Gomez-Rodriguez et. al.'12]
- Sample complexity of learning networks from cascades
- Convex optimization, message passing, etc.
- Related: learning linear dynamical systems from time series

Common thread: theoretical works are based on optimal estimators (likelihood based) and used to recover the *entire network*

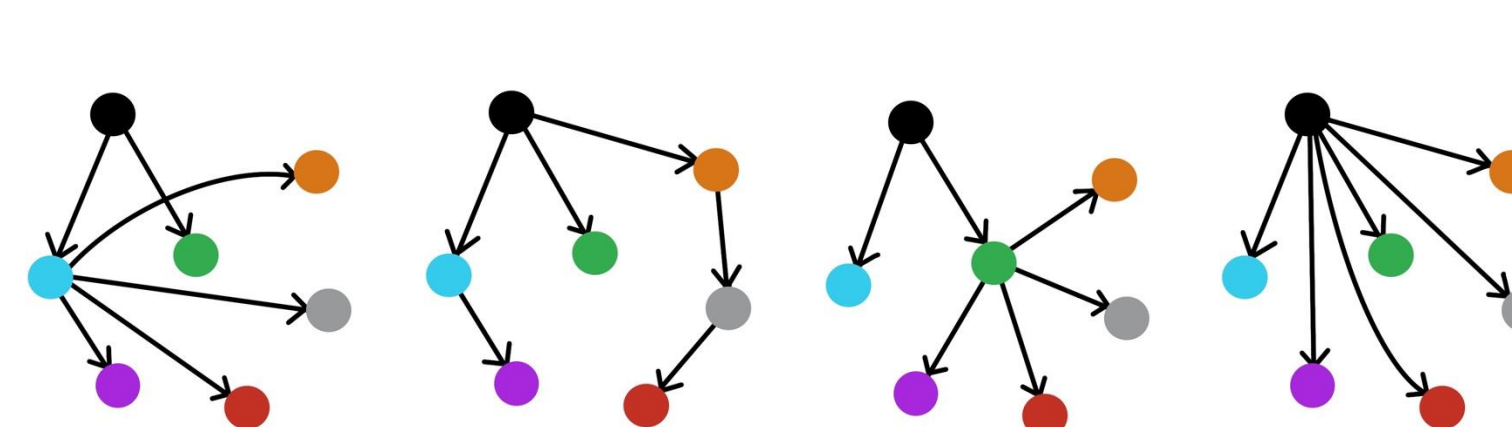
## This work: inference of high-degree vertices

**Mathematical abstraction:** a diffusion (e.g., cascade or epidemic) spreads on an unknown graph. Given the "infection times" for each vertex, learn the network.

Observations:



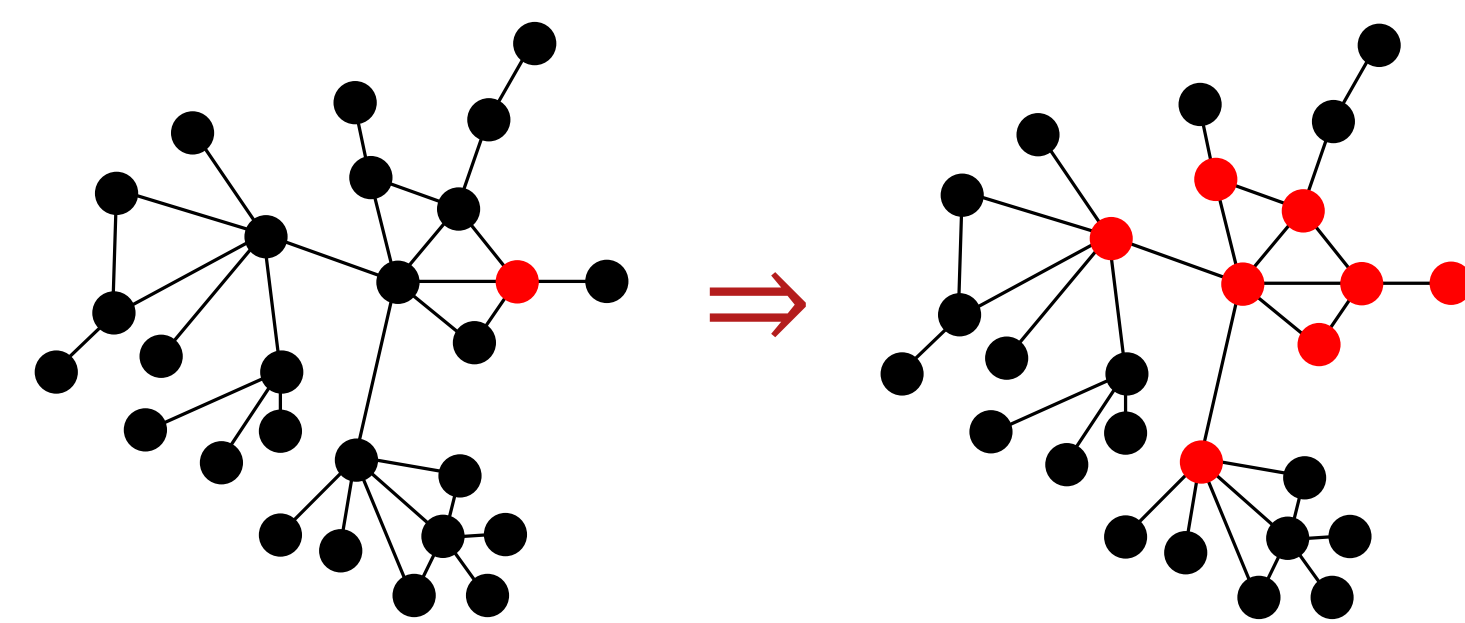
Many possible explanations:



To determine individual edges, need to observe many diffusions on the graph

Instead: attempt to infer only high degree vertices  
 $\Rightarrow$  can this be done in a single diffusion?

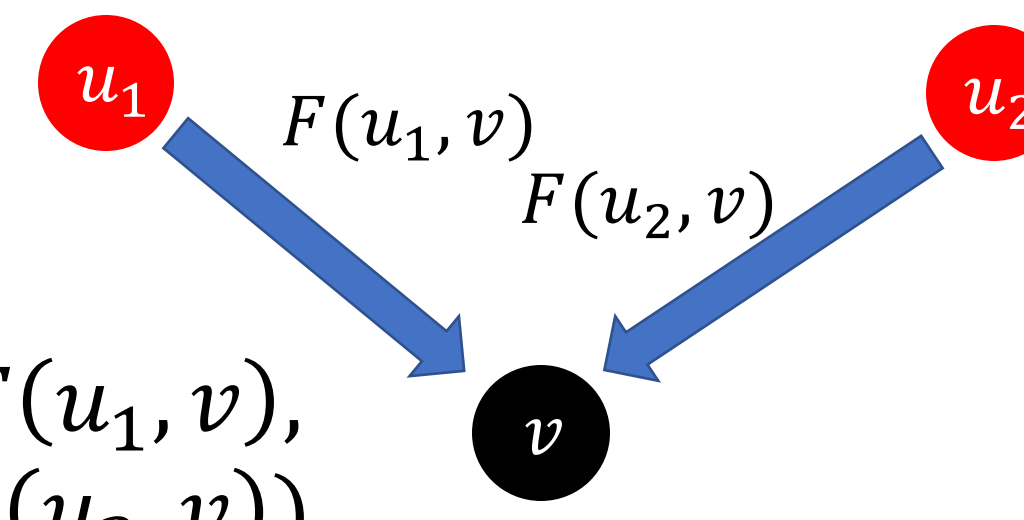
**The model of continuous time diffusion:** Susceptible-Infected model, first passage percolation



$\lambda$  = (pairwise) spreading rate

$T(v)$  = infection time of vertex  $v$

$$T(v) = \min(T(u_1) + F(u_1, v), T(u_2) + F(u_2, v))$$



Specifically, let  $\mathcal{I}(t)$  be the set of infected vertices at time  $t$ . Then, for an uninfected vertex  $v$ ,

$$P\{v \in \mathcal{I}(t + \epsilon) \mid \mathcal{I}(t)\} = \epsilon |\mathcal{N}(v) \cap \mathcal{I}(t)| + o(\epsilon)$$

where  $\mathcal{N}(v)$  is the set of neighbors of  $v$ .

**Model assumptions:** given a graph  $G$  with  $n$  vertices

- At most  $m$  (fixed) are *high-degree* (degree  $D \geq n^\alpha$ ),
- The rest are low degree (degree  $d \leq n^{o(1)}$ ),
- Two high degree vertices  $u$  and  $v$  satisfy  $\text{dist}(u, v) = \omega(1)$  as  $n \rightarrow \infty$ .

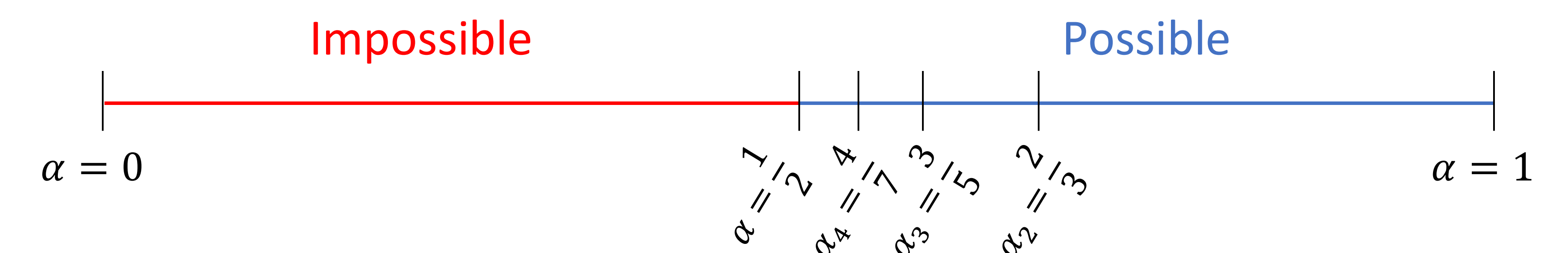
## Main result: phase transition in $\alpha$

**Theorem (Mossel-Sridhar'24).** Let  $\alpha < 1/2$ . There exists a distribution  $\mu$  over admissible graphs such that if  $G \sim \mu$  then it is impossible to tell whether there exists a high-degree vertex with probability greater than  $o(1)$  as  $n \rightarrow \infty$ .

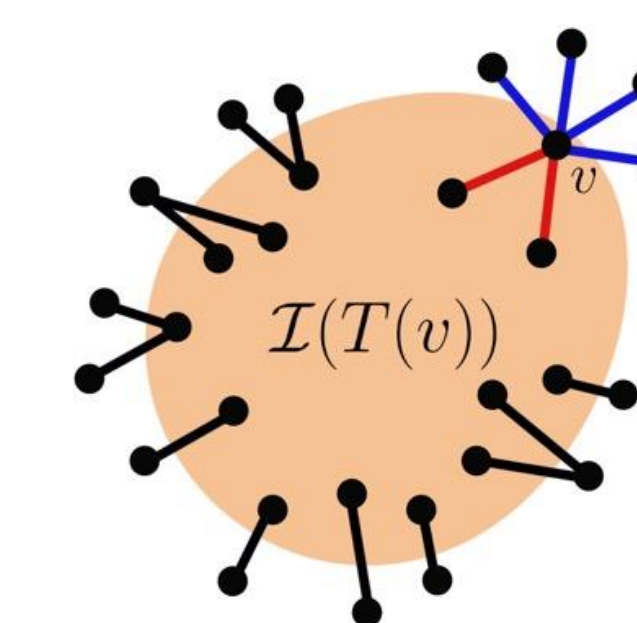
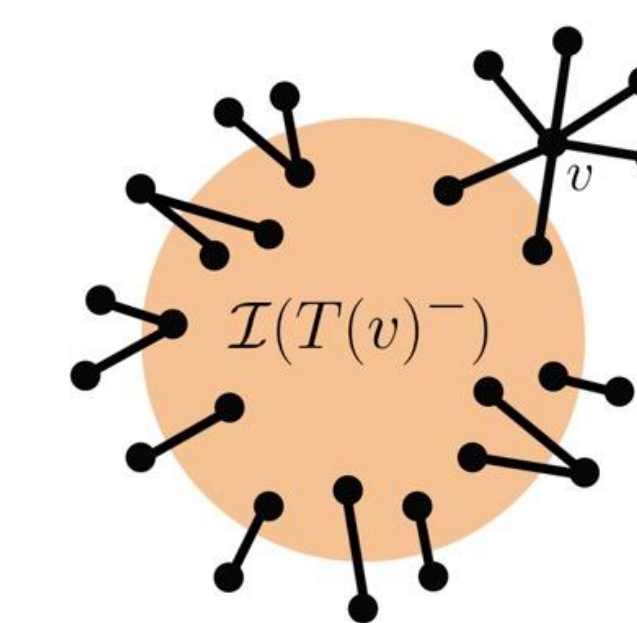
**Theorem (B.-Mossel-Sridhar'24+).** Let  $\alpha > 1/2$ . There is an algorithm (depending on  $\alpha$ ) which outputs a set  $S$  of time indices satisfying

$$\{T(v) : \deg(v) \geq D\} \subseteq S \subseteq \bigcup_{v: \deg(v) \geq D} (T(v) - \delta, T(v) + \delta)$$

with probability  $1 - o(1)$ , where  $\delta = 1/\text{poly}(n)$ .



## Main ideas: 1. Second derivative of infection curve $\mathcal{I}(t)$



$$\mathbb{E} \left[ \frac{d}{dt} I(t) \mid \mathcal{F}_t \right] = \lambda \cdot \text{cut}(\mathcal{I}(t))$$

$$\text{cut}(\mathcal{I}(T(v))) - \text{cut}(\mathcal{I}(T(v)^-)) = \deg(v) - 2|\mathcal{N}(v) \cap \mathcal{I}(t)|$$

- Second derivative of the infection curve is a (nearly) unbiased estimator for vertex degree. This identifies when a high-degree vertex is infected if  $\alpha > 2/3$ .

## Main ideas: 2. Higher derivatives of infection curve

**1. Local polynomial approximation.** Whp, for every  $t \geq 0$  there is a degree  $\ell - 1$  polynomial  $\bar{I}_{\ell-1}(\cdot, t)$  with  $\mathcal{F}_t$ -meas. coefficients such that

$$|I(s) - \bar{I}_{\ell-1}(s, t)| \lesssim n\delta^\ell + \sqrt{n\delta}, \quad s \in (t - \delta, t + \delta),$$

provided no abrupt changes (high-deg infections) in  $(t - \delta, t + \delta)$ .

**2. Analysis of higher-order derivatives.** We show that

$$\Delta_\delta^{(\ell)} I(t) = I(t + \delta) - \bar{I}_{\ell-1}(t + \delta, t^-) \pm O(n\delta^\ell + \sqrt{n\delta})$$

**Residual:** Difference between local polynomial prediction based on info before time  $t$  and actual process behavior

Note: Residual is similar in magnitude to error terms when no large jumps, by Step 1.

**3. Conditions for detection.** If  $\deg(v) \geq D$  then

$$|I(T(v) + \delta) - \bar{I}_{\ell-1}(T(v) + \delta, T(v)^-)| \gtrsim D\delta$$

The jump can be detected if

$$D \gtrsim n\delta^{\ell-1} + \sqrt{\frac{n}{\delta}} \Rightarrow D \gtrsim n^{\ell/(2\ell-1)} \text{ with } \delta = n^{-1/(2\ell-1)}$$