Parcours Intelligence Artificielle

Machine Learning for Network Modeling

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Why are we here?

Short course on Machine Learning for Network Modeling

Planning: 4 dense sessions, 2.5 hours each

- 1. Introduction to Graph Theory and Network Science
- Network models Static and dynamic graphs*
- 3. Structure and topology inference
- 4. Processes and signals over graphs

^{*} Session 2 is going to be given by Fabian Tarissan, CNRS, ENS Paris-Saclay

.:: In this lecture

- 1. Diffusion processes on networks
- 2. Control of diffusion processes
- 3. Control of competitive diffusion processes
- 4. Adding restrictions to the control problem
- 5. Application: Delay in transportation networks
- 6. Graph signals
- 7. Application: Monitoring information cascades in Twitter

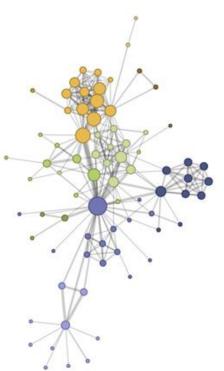
Basics

DPs arise in systems with interconnected agents (real or electronic networks)

- each agent has a variable state
- agent behavior depends on, and propagates to, its close environment
- the propagation causes changes in agents' state according to some "rules"



- Epidemiology: diseases/viruses
- Computer systems: computer viruses, fault cascade, computational errors (e.g. sensor networks)
- Social and information networks: information, ideas, rumors, social behaviors...



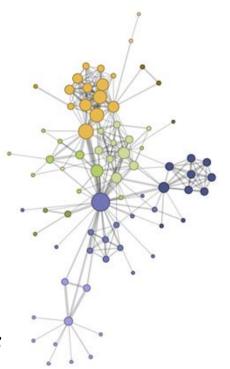
Basics

DPs arise in systems with interconnected agents (real or electronic networks)

- agent behavior dep This is what we will talk about
- the propagation cal



- Epidemiology: diseases/viruses
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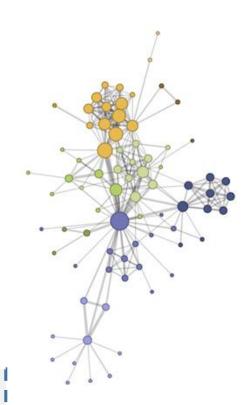


Directions of research

Depending on the situation, a DP can be desired or undesired

Roughly three directions of research

- Network assessment: worst case analysis, risk/vulnerability assessment
- **DP engineering:** influence maximization, (viral) marketing
- **DP suppression and control:** containment of viruses, rumors, social behaviors, etc., using *control actions*



Diffusion Models

Multitude of diffusion models

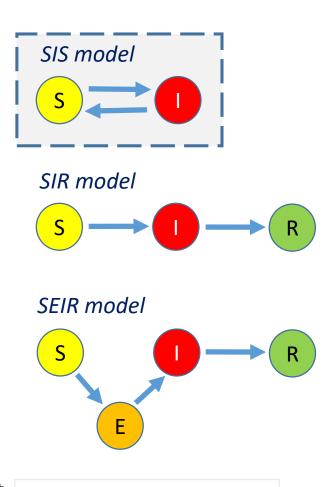
no single model describes all possible complex diffusion phenomena

Well-studied models

- compartmental models from epidemiology (SIS, SIR, SEIR, ...)
- other models from statistical physics (e.g. Percolation)
- common characteristic: constant propagation rates

Modern information-oriented models

- Information Cascades, Hawks Processes, ...
- Common direction: propagation rates variable in time to model user interest



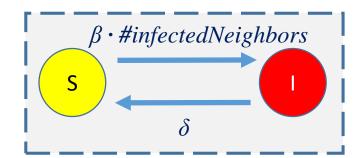
S: susceptible | E: exposed | R: recovered

Diffusion Models – SIS demo

Example

uncontrolled SIS process on contact network

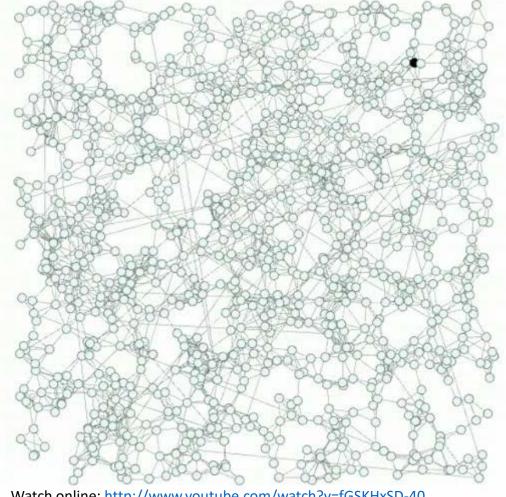
Homogeneous continuous-time SIS model for one node



 $X_i(t): 0 \to 1$ at rate $\beta \sum_i A_{ji} X_j(t)$ $X_i(t) \colon 1 \to 0$ at rate δ

- spreading rate β
- node self-recovery rate δ
- adjacency matrix A
- network state X
- two possible events each time: infection or recovery

SIS diffusion process in a contact nework

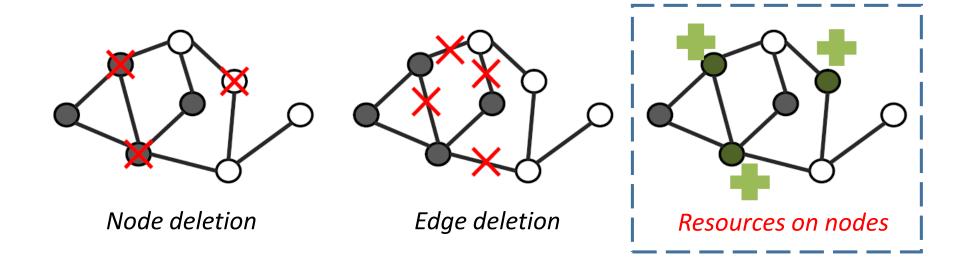


Watch online: http://www.youtube.com/watch?v=fGSKHxSD-40

Diffusion Suppression and control

Possible control actions

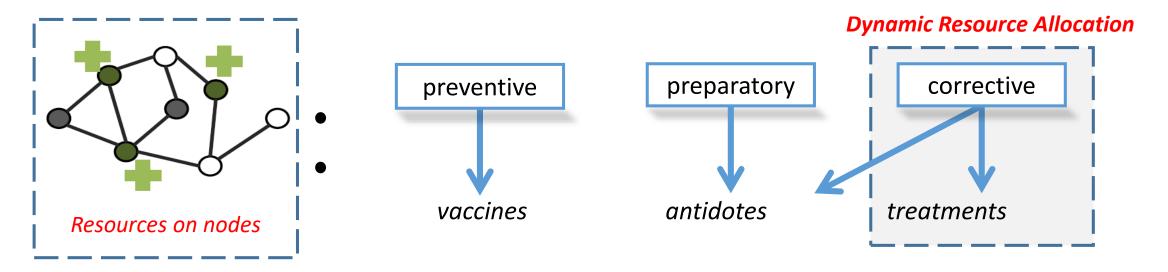
DP suppression and control using *control actions* on <u>nodes</u> or <u>edges</u>



Diffusion Suppression and control

Healing resources on nodes

DP suppression and control using *control actions* on <u>nodes</u>

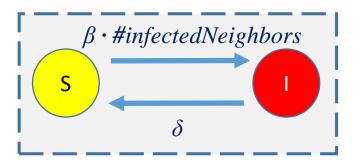


Diffusion Suppression and control

Introducing resources

Homogeneous continuous-time SIS model for one node

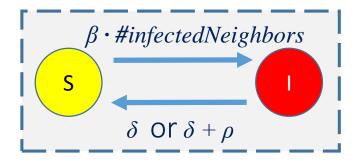
without control



$$X_i(t): 0 \to 1$$
 at rate $\beta \sum_j A_{ji} X_j(t)$
 $X_i(t): 1 \to 0$ at rate δ

- two possible events each time: infection or recovery
- spreading rate β
- node self-recovery rate δ
- adjacency matrix A
- network state X

with control



$$X_i(t): 0 \to 1$$
 at rate $\beta \sum_j A_{ji} X_j(t)$
 $X_i(t): 1 \to 0$ at rate $\delta + \rho R_i(t)$

- treatment efficiency ho
- resource allocation R

Dynamic Resource Allocation (DRA)

A modelling and control framework

DRA objective

$$\min_{R} C_{\gamma}(R) = \int_{t=0}^{+\infty} e^{-\gamma t} \mathbb{E}[N_{I}(t)] dt$$

$$\min_{R} C_{\gamma}(R) = \int_{t=0}^{+\infty} e^{-\gamma t} \mathbb{E}[N_{I}(t+u)|X(t) = X] du$$

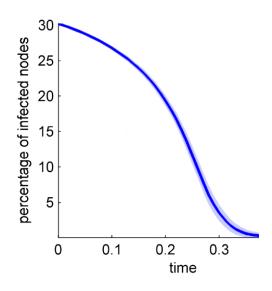
Formally a DRA strategy

$$R: \mathbb{R}_+ \to \{0,1\}^N$$

s.t. $\forall t \in \mathbb{R}_+, \sum_i R_i(t) \leq b(t)$

Constraints

- unlimited resources, disposed at limited constant rate
- limited accumulation of resources on single node
- inability to store resources



Dynamic Resource Allocation (DRA)

Score-based strategies

Score-based DRA strategies

$$R_i(t) = \begin{cases} 1 & \text{if } S_i(X(t)) \ge \theta_t \\ 0 & \text{otherwise} \end{cases}$$

where
$$\sum_{i} R_i(t) = b_{tot}$$

Heal the nodes with the top- b_{tot} ranked scores...

Baseline heuristics and LRIE

Strategy	Scoring function $S^i(X)$ for node i		
RAND	$\sigma(X_i)+R_i$, where R_i is i.i.d. uniform in [0, 1]		
MN	$\sigma(X_i) + \sum_j A_{ij}$		
PRC	$\sigma(X_i) + \overline{P_i}$, where P_i is the PageRank score for		
	node i		
LRSR	$\sigma(X_i) + (\lambda_1 - \lambda_1^{G \setminus i})$, where λ_1 is the largest eigen-		
	value of A , and $\lambda_1^{G\setminus i}$ the largest eigenvalue of		
	the matrix $A^{G\setminus i}$ for the network without node i		
MSN	$\sigma(X_i) + \sum_j A_{ij} \overline{X}_j$		
LIN	$\sigma(X_i) - \sum_{j=1}^{3} A_{ji} X_j$		
LRIE	$\sigma(X_i) + \sum_j [A_{ij}X_j - A_{ji}X_j]$, sums MSN and LIN		

•
$$\sigma(1) = 0$$
 and $\sigma(0) = -\infty$

OPTIMAL Greedy DRA

LRIE - Largest Reduction of Infectious Edges

Derivation

rewrite the DRA objective according to the Markovian property

$$\min_{R} C_{\gamma}(R) = \int_{t=0}^{+\infty} e^{-\gamma t} \mathbb{E}[N_{I}(t)] dt$$

$$\min_{R} C_{\gamma}(R, t, X) = \int_{u=0}^{+\infty} e^{-\gamma u} \mathbb{E}[N_{I}(t+u)|X(t) = X] du$$

then, a second order approximation

$$C_{\gamma}(R, t, X) = \frac{1}{\gamma} \sum_{i} X_{i} + \frac{1}{\gamma^{2}} \Phi'_{t, X}(0) + \frac{1}{\gamma^{3}} \Phi''_{t, X}(0) + O(\frac{1}{\gamma^{4}})$$

$$S_{\text{LRIE}}(X(t)) = A\overline{X}(t) - A^{\top}X(t)$$

$$= \left[\sum_{j} [A_{ij}\overline{X}_{j}(t) - A_{ji}X_{j}(t)]\right]_{i=1}^{N}$$

For an infected node *i*

$$\sum_{j}[A_{ij}\overline{X}_{j}(t)-A_{ji}X_{j}(t)]$$

virality

vulnerability

infectious edge

OPTIMAL Greedy DRA

LRIE - Largest Reduction of Infectious Edges

Toy example



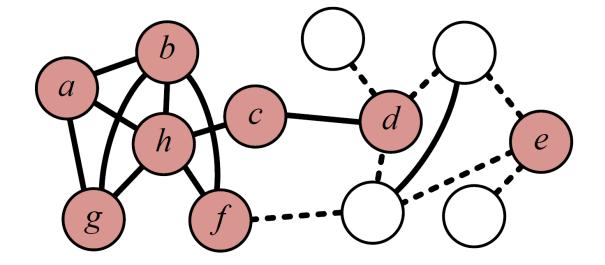
infected node



healthy node

edge between nodes of the same state

possible to transmit the disease



- Node **h** is the most central
- Node **e** and **d** are the most viral
- Node e is the least vulnerable (safest)

LRIE node ranking

Priority 1: $e / S_e = 3-0$

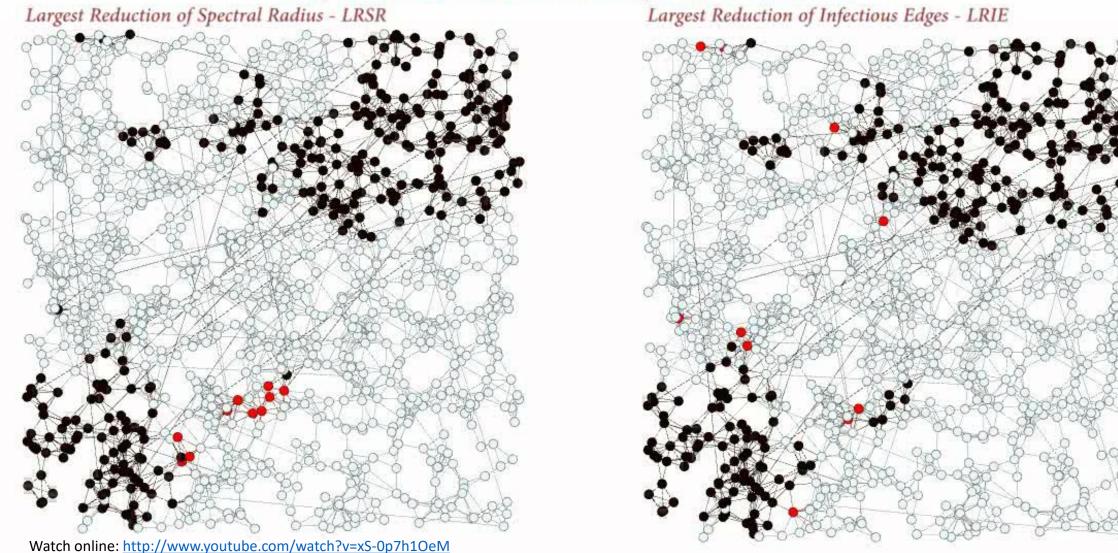
Priority 2: $d / S_d = 3-1$

Priority 3: $f / S_f = 1-2$

OPTIMAL Greedy DRA

Demonstration on an artificial contact network

Comparison of Resource Allocation strategies for diffusion control



LRIE: pros & Cons



Advantages

- brings the intuitive idea of reduction of infectious edges (front)
- optimal greedy, fast and quite efficient
- can adapt to network and/or budget changes
- not difficult to imagine a distributed version

Disadvantages

- ignores macroscopic network properties (e.g. clusters)
- cannot apply coordinated actions

Question to answer

LRIE is particularly elegant but greedy?

Can we do better?

(Global) Priority Planning

Definitions

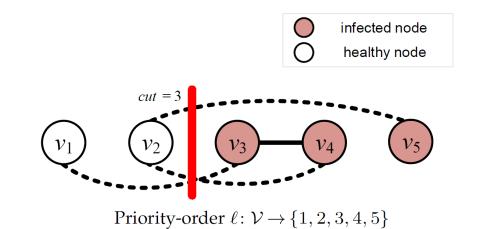
Priority-order: a bijection ℓ : $\mathcal{V} \rightarrow \{1, ..., N\}$ s.t. $\ell(v)$ the position of node v in the order

Priority planning: DRA strategies that are based on a priority-order

• limited budget r, max resource per node ρ , healing top-q(t) nodes (i.e. left-most)

$$q(t) = \min \left\{ \lceil \frac{r}{\rho} \rceil, \sum_{i} X_i(t) \right\}$$

$$\rho_i(t) = \begin{cases} \frac{r}{\lceil \frac{r}{\rho} \rceil} & \text{if } X_i(t) = 1 \text{ and } \ell(v_i) \leq \theta(t); \\ 0 & \text{otherwise} \end{cases}$$



Global Priority Planning

Graph theoretic properties of a priority-order

Cut at position
$$c$$
: $C_c(\ell) = \sum_{i,j} A_{ij} \mathbb{1}_{\{\ell(v_i) < c \le \ell(v_j)\}}$

MaxCut of
$$\ell$$
 $\mathcal{C}^*(\ell) = \max_{c=1,\ldots,N} C_c(\ell)$

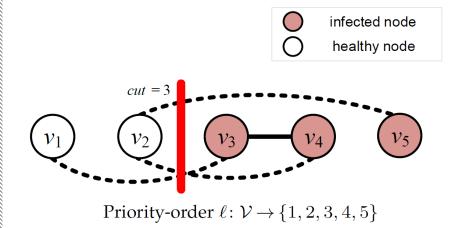
Cutwidth of
$$G$$
: $\mathcal{W} = \min_{\ell} \mathcal{C}^*(\ell)$

Extinction time:
$$\tau_x = \min\{t \in \mathbb{R}_+ | X(0) = x, X(t) = \mathbf{0}\}$$

- non-inf random quantity depending on the DRA strategy
- *sub-critical* behavior: $\mathbb{E}[\tau_x] \leq \text{polynomial function}$
- super-critical behavior: $\mathbb{E}[au_x]$ > exponential function

Requirement for designing a strategy:

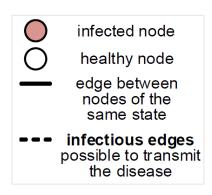
lacktriangle connect the properties of the order ℓ to $\mathbb{E}[au_x]$

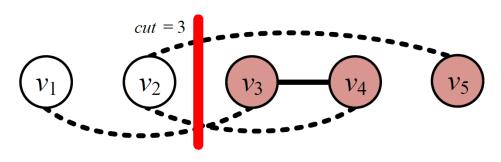


Priority Planning

MaxCut Minimization strategy (MCM)

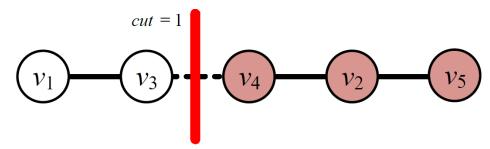
Toy example





Priority-order with MaxCut = 3

(a) Priority-order $\ell \colon \mathcal{V} \to \{1, 2, 3, 4, 5\}$



Priority-order with minimal MaxCut = 1

- (b) Priority-order $\ell' : \mathcal{V} \to \{1, 3, 4, 2, 5\}$
- Red vertical line: the *front* separating the healthy (left) from the infected part (right) of the network
- The MaxCut indicates highest vulnerability for the healthy part and is the most difficult step of the priority plan

Theoretical results

How good priority-orders are?

UPPER BOUND

Let d the maximum number of neighbors, $q=\lceil\frac{r}{\rho}\rceil$ the number of treated nodes, and $\epsilon=\frac{d(3+2\ln N+4q)}{\mathcal{C}^*(\ell)}$. Assume that:

$$r + \delta q > eta \mathcal{C}^*(\ell) \left(1 + 2\sqrt{\epsilon} + \epsilon
ight)$$

Then the following upper bound holds for the expected extinction time $\mathbb{E}[\tau_1]$:

$$\mathbb{E}[au_1] \leq \frac{6N}{eta}.$$

Theoretical results

How good priority-orders are?

LOWER BOUND

Let $\delta=0$ (no self- healing), $\eta\in[0,\frac12]$, and d, q and ϵ defined as before. Assume that $q<\frac{\mathcal C^*(\ell)}{d}$ and

$$r < (1-\eta)eta \mathcal{C}^*(\ell)(1-rac{dq}{\mathcal{C}^*(\ell)})$$

Then the following lower bound holds for the expected extinction time $\mathbb{E}[\tau_1]$:

$$\mathbb{E}[au_{\mathbf{1}}] \geq rac{1}{r} \mathrm{exp}\left(rac{\eta^2}{12} \Big(rac{\mathcal{C}^*(\ell)}{d} - q\Big)
ight).$$

Maxcut Minimization (MCM)

MCM Strategy

MCM strategy

- seeks for the priority-order ℓ with the *minimum* $\mathit{MaxCut}\ C^*(\ell)$ of edges
- heals the q(t) leftmost infected nodes in ℓ
- uses a relaxation of $\ell_{_{MCM}}(\mathcal{G}) = \operatorname*{argmin}_{\ell} \mathcal{C}^*(\ell)$ by

```
MpLA: \phi(\mathcal{G}, \ell) = \left(\sum_{i,j} A_{ij} |\ell(v_i) - \ell(v_j)|^p\right)^{1/p}
```

Algorithm 1 MCM strategy *▶ Prior to he diffusion process*: Compute the priority-order $\ell = \ell_{MCM}(\mathcal{G})$ by minimizing the maxcut $\mathcal{C}^*(\ell)$ Order the nodes of \mathcal{G} according to ℓ , i.e. compute the node list $(v_1, ..., v_N)$ s.t. $\forall i \in \{1, ..., N\}, \ell(v_i) = i$ During the diffusion process: **Input:** network \mathcal{G} , state vector X(t), resource budget r, resource threshold ρ **Output:** the resource allocation vector $\rho(t)$ $q \leftarrow \left\lceil \frac{r}{\rho} \right\rceil$ if $\sum_{i} X_{i}(t) < q$ then return $\frac{r}{a}X(t)$ end if // a zero vector in \mathbb{R}^N $\rho(t) \leftarrow \mathbf{0}$ $budget \leftarrow q$ $i \leftarrow 1$ while budget > 0 do if $X_{v_i}(t) = 1$ then $\rho_{v_i}(t) \leftarrow \frac{r}{a}$ $budget \leftarrow budget - 1$ end if $i \leftarrow i + 1$ end while return $\rho(t)$

Maxcut MinimiZation (MCM)

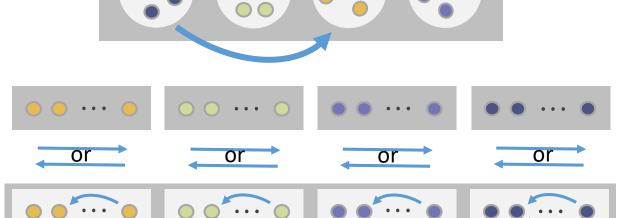
Solving the MLA problem

Learning an ordering for a network

1. find communities in G and order them (high-level nodes) with spectral sequencing

2. order nodes inside each cluster with spectral sequencing, orient to each other, and then optimize with node swaps internally to clusters

3. apply the swap-based approach again to the overall node ordering



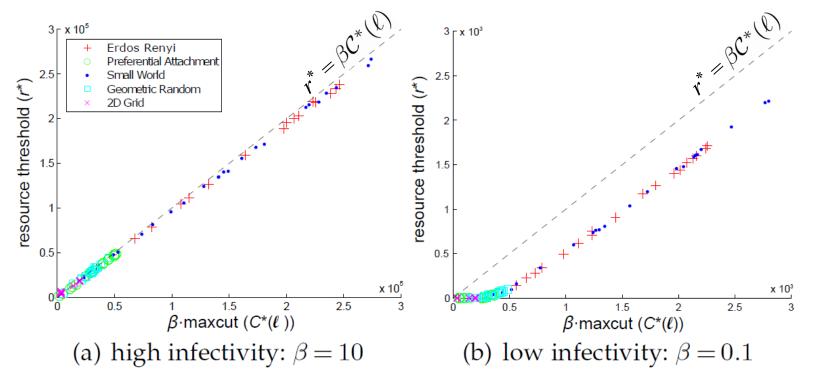
1b.

Results

Quality of the theoretical bound

Verifying

$$r^* \approx \beta C^*(\ell)$$



- picks orderings at random out of MCM, RAND, MN, LN, LRSR
- various random network models, N = 1,000, $q = \{1,...100\}$
- $lacktriangleright r^*$ was estimated empirically with simulations

Results

Experiments on real-networks

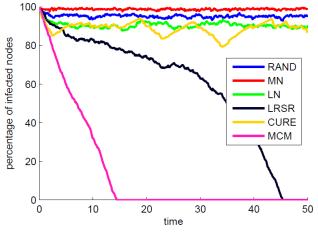
GermanSpeedway

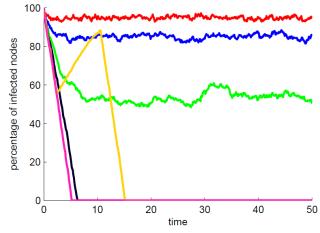
 $N = 1,168 \text{ nodes}, E = 1,243 \text{ edges}, \quad max(d) = 12, \beta = 1, \delta = 0, q = 1$

MaxCut: 650+/-50 RAND, 379 MN and LN, 104 LRSR, 29 CURE and MCM

OpenFlights

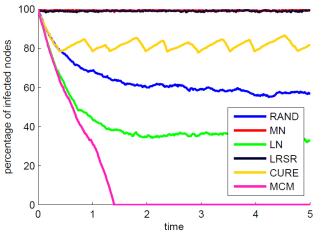
N=2,939 nodes, E=30,501 edges, $max(d)=242,\,\beta=1,\,\delta=0,\,q=1$ $MaxCut:\,7,800+/-100$ RAND, 7,504 MN and LN, 6,223 LRSR, 2,231 CURE and MCM



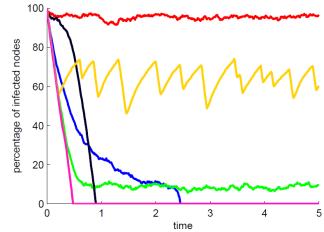


(a) low resource budget: r = 100

(b) high resource budget: r = 250



(a) low resource budget: r = 3000



(b) high resource budget: r = 7000

Global Priority Planning

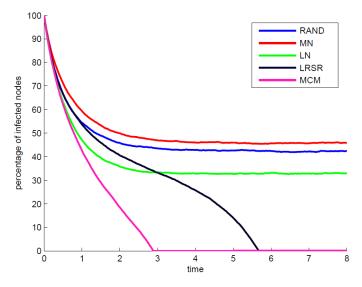
Experiments on real-networks

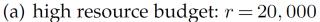
Subset of Twitter network with 81.306 nodes

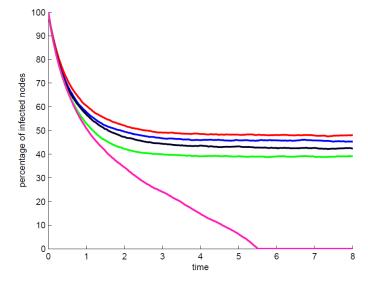


MCM can remove the contagion with ~5 times less resources than its best competitor!!

Strategy	Maxcut	Maxcut	Expected resource threshold
		% w.r.t. RAND	$\delta = 1, \beta = 0.1, q = 100$
RAND	$670,000 \pm 1000$	100.0 %	67,000
MN	628,571	93.8 %	62,957
LN	628,571	93.8 %	62,957
LRSR	349,440	52.2 %	34,944
MCM	71,956	10.7 %	7,196



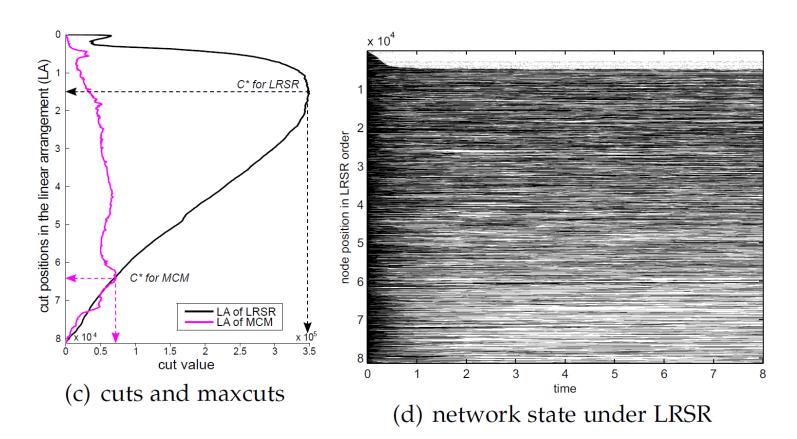




(b) low resource budget: r = 12,000

Global Priority Planning

Experiments on real network (TwitterNet)



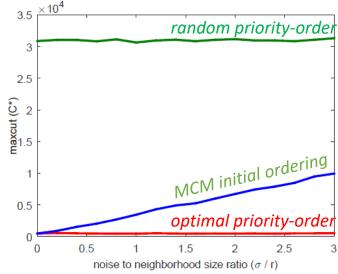
node position in MCM order time (e) network state under MCM

Robustness analysis

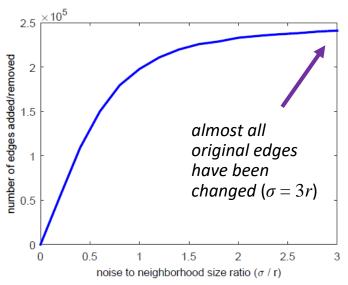
Experiments on an increasingly perturbed contact network

Contact network in $[0,1]^2$ where each node is connected with all nodes in radius r

The priority ordering remains valid after local modifications of the network connectivity



(a) $C^*(\ell)$ value as a function of noise



(b) number of edges added or removed as a function of noise

Question to answer

From disease epidemics to... digital and social epidemics

Are the strategies we have at hand efficient in the presence of **competition**?

Can we do better?

Competitive social Diffusion Processes

Motivation

Various studies have identified diffusion and competition in lifestyle and social behavior

- Obesity [Wing et al. 2009, Christakis et al. 2007, Hill et al. 2010]
- Smoking [Poulsen et. al. 2002, Christakis et al. 2009]
- Alcohol consumption [Rosenquist et al. 2010]
- Emotions in social networks [Fowler et al. 2009, Hill et al. 2010]

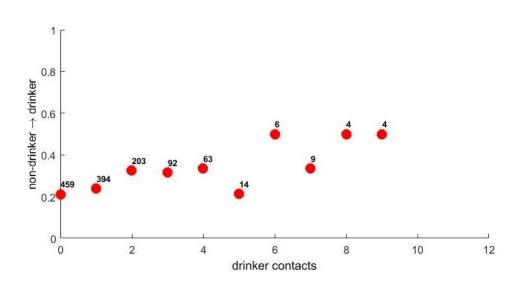
Features / challenging properties

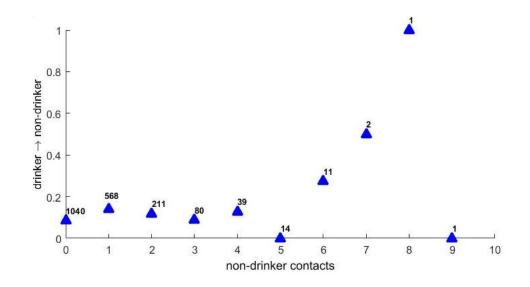
- SIS-like processes with evidence of competition
- "evidently" complex propagation functions
 - non-linear
 - with saturation points

Competitive social Diffusion Processes

Motivation

Example: human behavior for alcohol consumption





Modeling competition with SIS

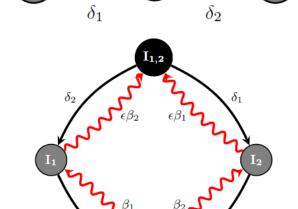
Competitive models from literature

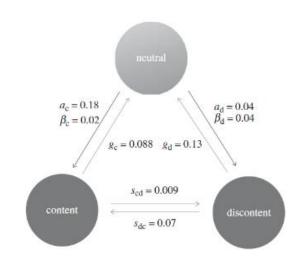
Related epidemic models

- SISa: includes spontaneous infection [Hill et. al. 2010]
- S I₁I₂S:

[Prakash et. al. 2012]

■ S I_{1|2}S
[Beutel et. al. 2012]

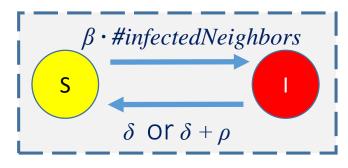




A NOVEL Competitive SIS model

Introducing arbitrary propagation functions and competition

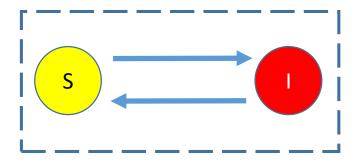
SIS with control



 $X_i(t) \colon 0 \to 1$ at rate $\beta \sum_j A_{ji} X_j(t)$

 $X_i(t): 1 \to 0$ at rate $\delta + \rho R_i(t)$

Generalized SIS with competition and control



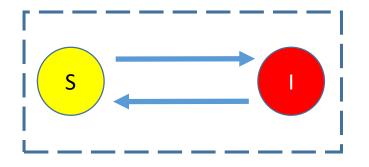
$$X_i(t): 0 \to 1$$
 at rate $\mathcal{I}_i(X(t))$
 $X_i(t): 1 \to 0$ at rate $\mathcal{H}_i(X(t)) + \rho R_i(t)$

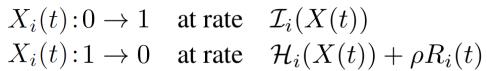
- with \mathcal{I}_i and \mathcal{H}_i two node-specific memoryless propagation functions
- ... that represent the competing positive and negative diffusions

Our Competitive SIS model

Relaxation

Generalized SIS





Relaxation

Assumptions for the propagation functions:

- locality
- exchangeability
- Invariance

 \rightarrow They depend only on the node degree d_i and on the number infected neighbors n_i :

$$\mathcal{H}_i = \mathcal{H}(n_i, d_i)$$

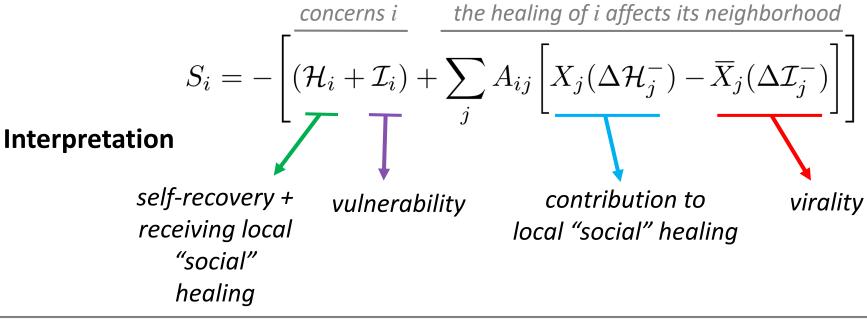
$$\mathcal{I}_i = \mathcal{I}(n_i, d_i)$$

Optimal greedy strategy for competition

Generalized LRIE (gLRIE)

Derivation: By minimizing $C_{\gamma}(R,t,X)$ we obtain the **gLRIE scoring function**

For an infected node i:



$$\mathcal{H}_i = \delta$$

Recovering LRIE:
$$\mathcal{H}_i = \delta$$
 $\mathcal{I}_i = \beta \sum_j A_{ji} X_j$ $\Delta \mathcal{H}_i^- = 0$

$$\Delta \mathcal{H}_i^- = 0$$

$$\Delta \mathcal{I}_i^- = \beta$$

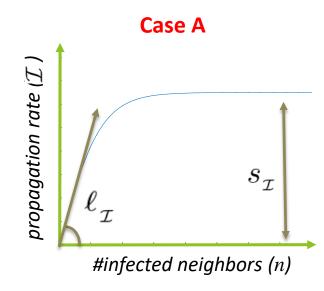
Examples of arbitrary propagation functions

Sigmoid propagation functions for experimentation :

- adjustable non-linearity
- adjustable saturation level

Two variations:

- Case A: Dependency only on #infected neighbors (n)
- **Case B:** Dependency only on the infection ratio (n/d)



$$\begin{cases} \mathcal{I}(n,d) = s_{\scriptscriptstyle \mathcal{I}} \big[1 - \frac{2}{1 + \exp\left(4\ell_{\scriptscriptstyle \mathcal{I}} *\right)} \big] \\ \mathcal{H}(n,d) = s_{\scriptscriptstyle \mathcal{H}} \big[1 - \frac{2}{1 + \exp\left(4\ell_{\scriptscriptstyle \mathcal{H}} (d - *)\right)} \big] + \delta \end{cases}$$

Evolution plots for random graphs

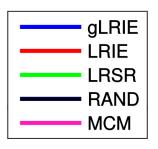
Case A: ~ to #infected neighbors + no competition

Erdös-Rényi networks: N = 1000 nodes, p = 0.001, mean degree 8, 10^4 simulations

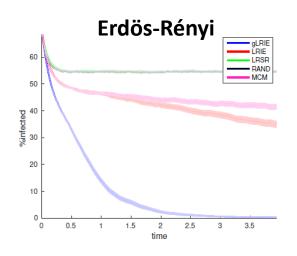
 $^{ullet}\mathcal{H}=0$, $s_{\scriptscriptstyle\mathcal{I}}=13$ and $b_{tot}=10$

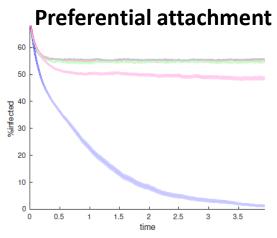
aLRIE RIE RSR from 'linear' to non-linear ${\mathcal I}$ RAND **MCM** (b) $\ell_{\mathcal{I}} = 0.1, \, \rho = 8$ (c) $\ell_{\tau} = 1, \, \rho = 63$ (d) $\ell_{\tau} = 3, \, \rho = 120$ (e) $\ell_{\tau} = 5, \, \rho = 140$ (a) $\ell_{\tau} = 0.01, \rho = 1.6$

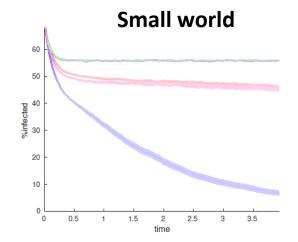
Evolution plots for random graphs

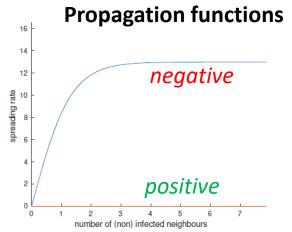


Case A: ~ to #infected neighbors

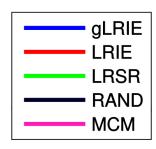




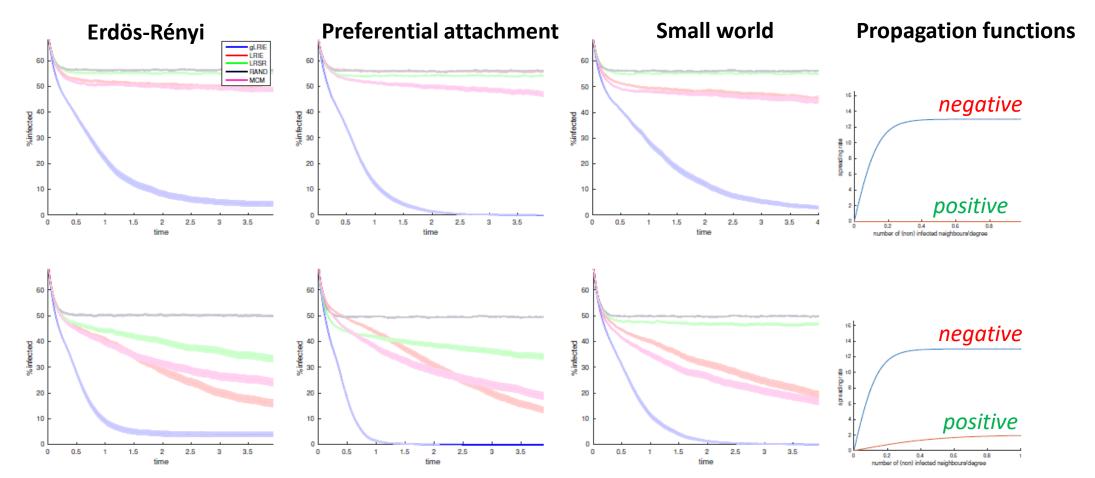




Evolution plots for random graphs



Case B: ~ to %infected neighbors



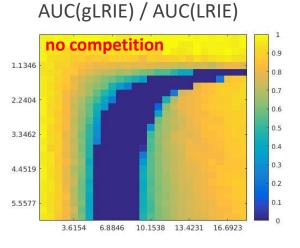
Results

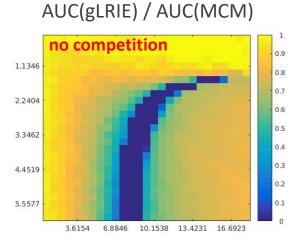
Real-world networks



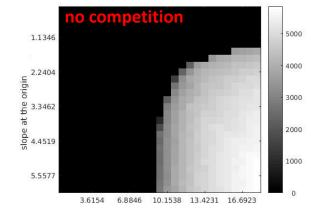
Gnutella

8846 nodes31839 edges









Results

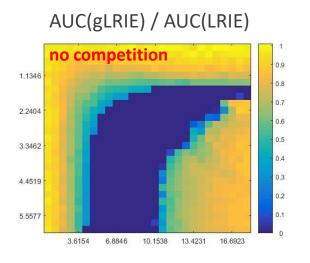
Real-world networks

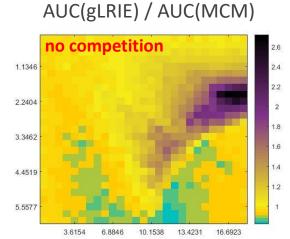
Heatmaps in the $\frac{saturation}{(s_I, \ell_I)}$ -space

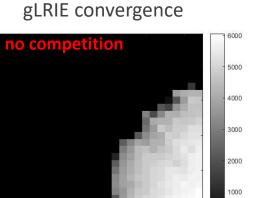
arXiv H.E.Physics

8637 nodes

24803 edges







10.1538 13.4231

saturation level (a / a)

1.1346

₫ 3.3462

ed ools 4.4519

5.5577

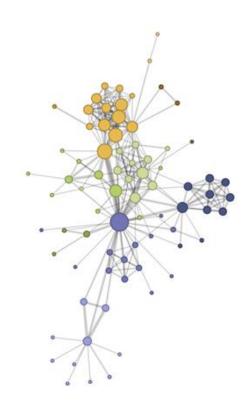
gLRIE pros and cons

Pros

- motivated by social contagions scenarios
- takes into account competition
- arbitrary propagation functions
- Inherits the adaptivity and elegance of LRIE

Cons

inherits the greediness and lack of co-ordination of LRIE



Question to answer

What about relaxing the requirements of the DRA class of resource allocation strategies

Standard DRA

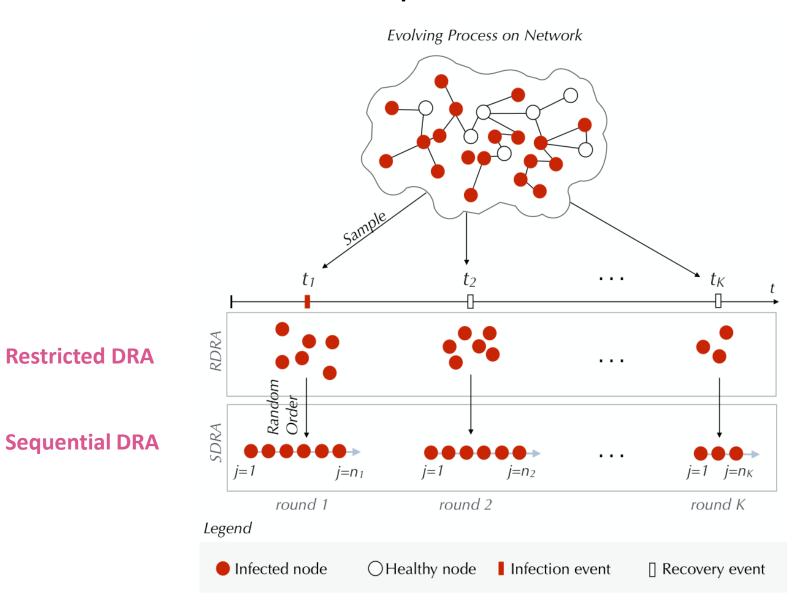
Motivations

- Unrealistic 'power' of the administrator
- Play with access and information

Assumption

access and information are inextricable

Restricted and Sequential DRA



The sequential selection problem (SSP)

BASIC VERSION

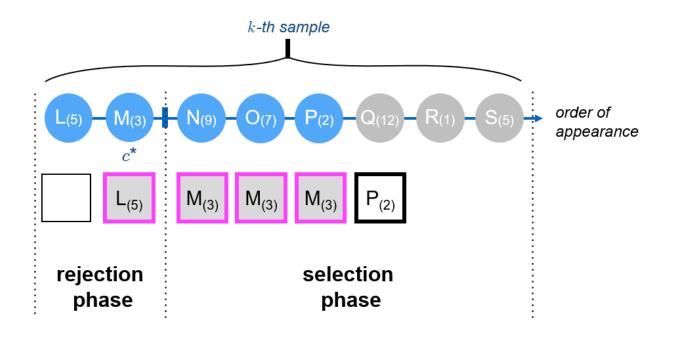
Constraints

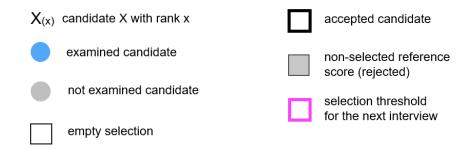
Immediate and **irrevocable** decision after each interview

No info about, or control to, the input

Limitations of the classical SSP setting

- Cold start: zero prior knowledge
- Single-shot problem...





The Warm-starting Sequential Selection Problem

Warm-starting Sequential Selection Problem (WSSP)

Background

$$\mathcal{B} = (b, n, s, \mathbf{C}^R)$$
, where

- $n \in \mathbb{N}$: nb. of nodes to evaluate,
- $b \in \mathbb{N}$: nb. of resources,
- $s: \mathcal{V} \to \mathbb{R}_+$: scoring function,
- $\mathbb{C}^R \subset \mathcal{V}$: preselection, i.e. set of currently treated nodes
- Process & Decisions

$$\mathbf{C} = (C_1, ..., C_n) \in \mathcal{P}_n(\mathcal{V} \setminus \mathbf{C}^R) \text{ and } (R_{C_1}, ..., R_{C_n}) \in \{0, 1\}^n$$

Evaluation

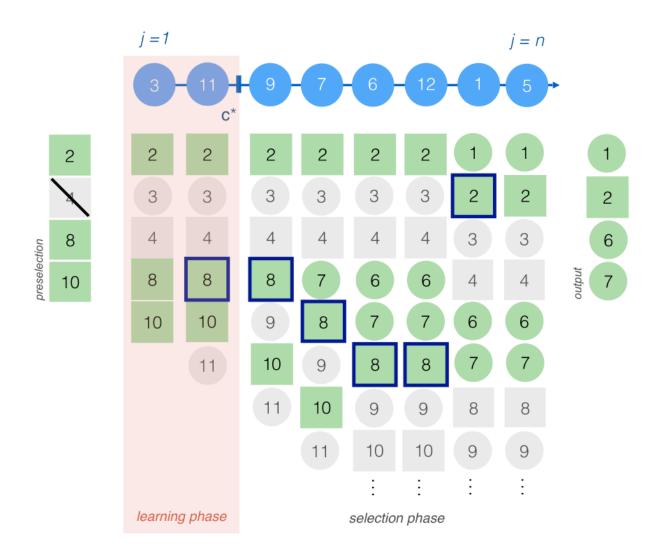
The regret function is defined as:

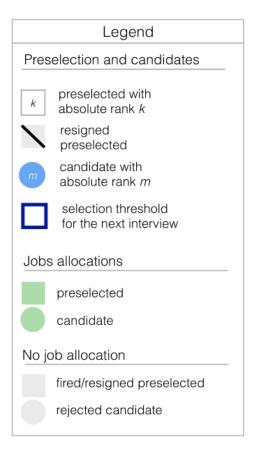
$$\phi_{\mathcal{B}} = \max_{R_i^*, i \in \mathcal{C}} (\mathbf{S} \cdot \mathbf{R}^*) - (\mathbf{S} \cdot \mathbf{R}) \in \mathbb{R}_+,$$

where
$$C = (\mathbf{C}^R, C_1, ..., C_n) \subset \mathcal{V}$$
.

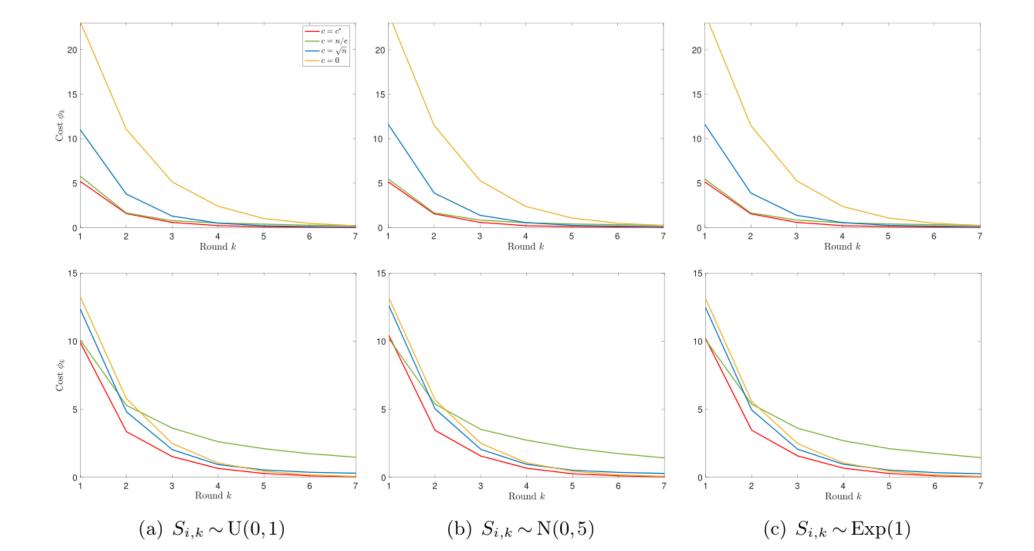
Goal: minimize $\mathbb{E}[\phi_{\mathcal{B}}]$.

Warm-starting & multi-round sequential selection processes





Warm-starting & multi-round sequential selection processes



Pluging warm-starting into Sequential DRA

Various strategies

- Hiring-above-the-mean (MEAN) [Broder et al., 2009]:
 Acceptance threshold is the mean of employees
 Goal: grow the company as much as possible while keeping maximal the average score of the employee
- Cutoff-based Cost Minimization (CCM) [Fekom, Vayatis, Kalogeratos 2019]
 Generic algorithm i.e. works with any scoring function Goal: minimize the expectation of the ranks of the selected
- Warm-starting Dynamic Thresholding (WDT)) [Fekom, Vayatis, Kalogeratos 2019]
 Assumes score distribution is known
 Optimal acceptance threshold
 Goal: maximize the expectation of the scores of the selected

Example on a scale-free network

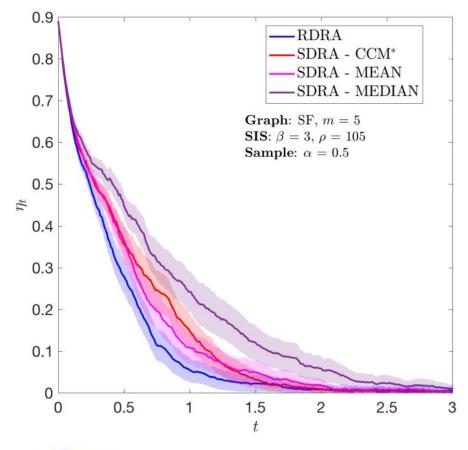


Figure: Percentage of infected nodes w.r.t. time.

Discussion



