

From a static to a dynamic analysis of complex networks

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Soutenance de HDR



Outline

- 1 Context & Overview
- 2 Describing the spreading dynamics
- 3 Modeling the interaction structure and dynamics
- 4 Conclusion

About complex networks analysis

What are complex networks?

Extracted from real-world data

ex: computer networks, social networks, biological networks, ...

- agents interacting, no “central brain”
- \Rightarrow emerges from local interactions

Studied as graphs

- system common features visible with graphs
ex: heterogeneous degree distribution, high clustering, ...
- similar methods and algorithms

Accounting for temporality

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Scientific interest

Attractive because ...

- Variety of objects
- Create bridges between different topics

Getting the “big picture”

... and in retrospect

- Variety of communities to learn about
- Highly active research fields

⇒ Stimulating but demanding

Focus on three main axes of research

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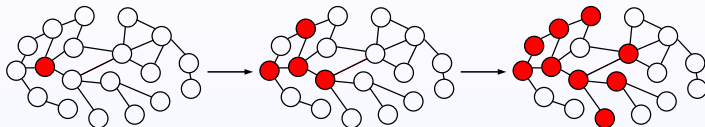
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Focus on three main axes of research

Describing spreading processes

Dynamical processes on complex networks

Synchronization, opinion dynamics, routing, **spreading**, ...



Models which represent:

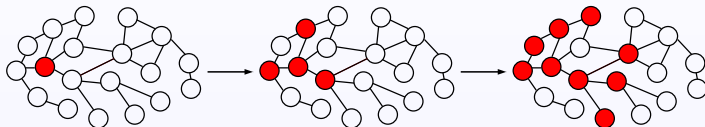
- an infection propagating in a population
- a packet spreading in a computer network
- ...

Also a means to investigate the dynamical structure

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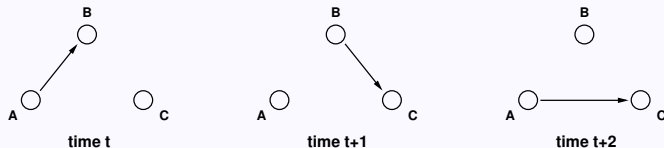
Also a means to investigate the dynamical structure

Predicting and recovering links

Understanding the microdynamics

Unknown mechanisms \Rightarrow learn from data

ex: *triadic closure*



Specificities of the problem

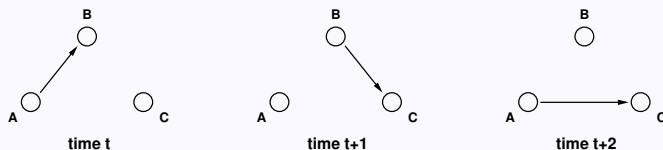
- tuning number of predictions
- on large graphs
- how to incorporate temporal information?

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Model the interaction structure and dynamics

What is modeling here?

Generating artificial structures resembling real data



⇒ allows to uncover possible explanations

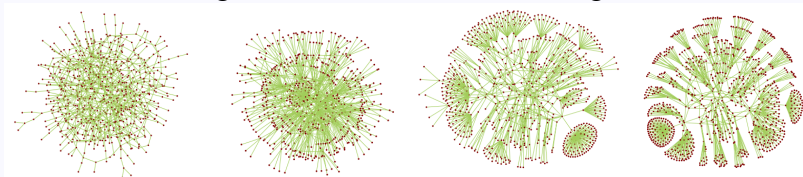
Challenges

- realistic structural constraints on graphs
- towards modeling temporal networks

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Presentation outline

- **Describing spreading dynamics**
- Predicting links
- **Modeling the structure and dynamics**

Highlight the connexion between these topics

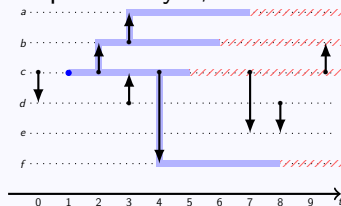
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Spreading cascade in a temporal network

Model analyzed

SIR model: infection probability 1, deterministic recovery after τ



Spreading cascade (but many other names)

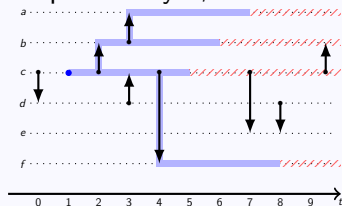
Advantages:

- simplicity of the model
- allows to tune the time scale
- probe of the temporal network (motif)

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Role of in/out-degree correlations

Peruani, Tabourier, 2011

Phonecall dataset

~14 millions phone calls between ~1 million users during 1 month
European mobile phone provider

Strongly asymmetric roles (super-spreaders, super-receivers)

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Critical recovery time τ_c

τ such that cascade size diverges

$$\tau_c = \frac{\langle k_{out} \rangle}{\langle \rho \rangle \langle k_{out} \cdot k_{in} \rangle}$$

with some assumptions (no cycles)

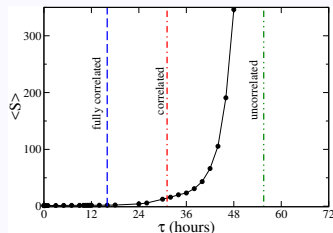
Role of in/out-degree correlations

Peruani, Tabourier, 2011

Experimental observations

- real (k_{in}, k_{out}) distribution: 32 hours
- fully-correlated data $k_{in} = k_{out}$: 14 hours
- uncorrelated data: 55 hours

⇒ significant differences



underestimation of the threshold

Role of individual temporal patterns

Measuring the characteristics of the cascades

- cascade = tree \Rightarrow size (σ), depth (δ)
- abundance of each type of cascades

To what can we compare these measures?

Comparing to null models

- *time mixing model*: random shuffle of all time labels
- *correlation mixing model*: random shuffle of time labels for interactions originating from same source



Role of individual temporal patterns

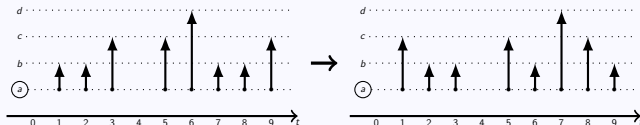
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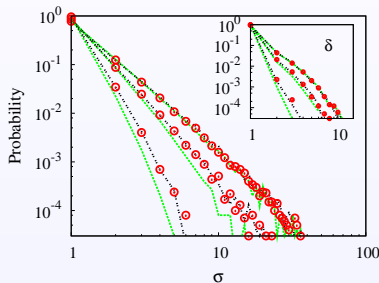


- └ Describing the spreading dynamics
- └ Role of individual temporal patterns

Experimental results on phonecall data

Tabourier, Stoica, Peruani 2012

Distributions of σ (size) and δ (depth) for $\tau = 30\text{mins}$, 3hrs , 12hrs



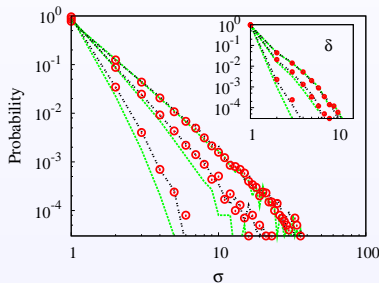
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real data

time mixing model breaks bursty temporal activity patterns
 \Rightarrow Bursty patterns accelerates spreading at short timescales

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About the *correlation mixing model*:

- σ and δ insufficient to distinguish real data from *cmm*
- but comparing cascades and other patterns with *cmm*
→ *cmm* underestimates short cyclic patterns

cmm breaks correlation between temporal activity and structure
⇒ correlations create a trapping effect

Consistent with “while bursts hinder propagation at large scales,
conversations favor local rapid cascades” *Miritello et al., 2011*

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Conclusion and prospect

- Schematic picture of spreading:
rapid cascades at small scales, slower on larger scales
- Due to individual activity patterns, correlations between structure and temporality
- Understanding one step further how the temporal network structure affects spreading phenomena
→ new null models for temporal networks

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Motivation

To explain the observed structure

- generate a structure that resembles real data
- unbiased interpretation assumes **uniform generation**

With graphs

- either biased generation model
ex: Barabási-Albert, Watts-Strogatz, ...
- or relatively simple properties
ex: Erdős-Rényi, configuration model, ...

Versatile method to generate **uniformly** graphs
with complex constraints

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Switching methods principle

Standard switch method

ex: generate a random graph with a fixed degree distribution

- start from a graph of the set
- iterate: select two edges randomly, exchange the ends
 - if new graph \in set, continue
 - if new graph \notin set, exchange back, continue

After enough iterations \rightarrow random element of the set

More complex constraints

Problem: standard method does not guarantee ergodicity
 \rightarrow biased generation process

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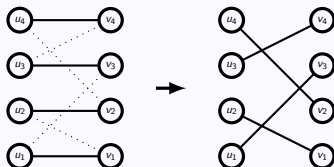
Problem: standard method does not guarantee ergodicity
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Obtaining ergodicity

Tabourier, Roth, Cointet, 2011

***k*-switch method**

- start from a graph of the set
- iterate: select k edges randomly, permute the ends randomly
 - if new graph \in set, continue
 - if new graph \notin set, exchange back, continue



Experimentally

low values of k sufficient with most constraints

Illustration

Tabourier, Cointet, Roth 2017

Scientific coauthoring networks

Scientists publishing papers together
underlying bipartite structure

→ interested in the structure **projection** on authors

- Usual models for social event-based networks:
 - standard configuration model M
 - bipartite configuration model B
- More complex model:
 - bipartite + monopartite configuration model on authors MB

MB demands k -switch generation

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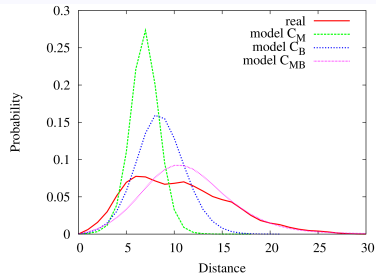
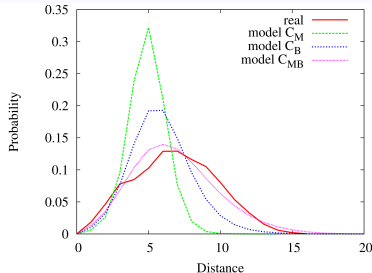
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Experimental results

Measure: distance distribution in the models vs real data

datasets: archeology in Britain, in Europe



⇒ more accurate model for coauthoring network

Generalization to temporal networks: principle

Motivation

Most models focused on aggregated properties *Gauvin et al., 2018*

→ alternate models preserving structuro-temporal properties

ex: fixed number of temporal motifs

Applying the k -switch method on temporal networks

time label used as an attribute constraint

⇒ same randomization technique

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Generalization to temporal networks: challenges

- **technical problem:**
more elaborate the constraint \Rightarrow slower generation process
- **interpretation problem:**
how to have more flexibility on the temporal granularity?
example of strict vs approximate simultaneity

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Conclusion

Common thread

- describing the temporal network
- predicting links
- modeling the structure of interactions
- on specific applications

serve to identify dynamical rules of the network
→ **reconstruct the general evolution of the system**

Thank you for your attention !

My collaborators

