19th Workshop on Modelling and Mining Networks Network Diffusion — Framework to Simulate Spreading Processes in Complex Networks

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Agenda

In this talk, I will introduce a computational library — network-diffusion. Here is the agenda:

- Motivation
- 2 Key Features
- 3 Example I a Predefined Model
- 4 Example II a Custom Model
- 5 Resources and References
- 6 Limitations of network-diffusion



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Spreading phenomena are one of the issues considered by a network science. They can be obeserved in various areas like: dynamics of political opinions, marketing campaigns, spread of epidemics, computer viruses, etc.



Figure: Artistic representation of a social network.¹



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¹Source: www.uniroma3.it/articoli/

Analytical approaches are often insufficient for large graphs, prompting researchers to use computational methods, i.e. simulations.



Thus, like in other branches of computer science, there have been developed tools which addres that issue, allowing to avoid starting from scratch and enhancing the reproducibility of results.²



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²Although network science still ingloriously stands out in this matter. • • • •

There is a bunch of tools that helps in sumulating diffusion processes in networks:

- **NDlib**[RMR⁺18],
- GLEaMviz[BGG⁺11],
- SimInf[WBEE19],
- STEM[DBE+19],
- EpiModel[JGM18],
- Sispread[ACBV07],
- ..

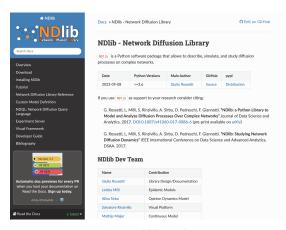


Figure: NDlib's website.



However, if we consider...

...more complex network models...

...spreading multiple processes at the same time...

...a gap among the available toolkits emerges.



A focus of our research group is oriented i.a. to: multilayer networks, temporal networks, spreading phenomena, data streams.



As a result of our recent activities, we decided to merge and wrap up a code we developed into a reusable library. We also decided to to share it with the community in an attempt of filling the gap in.



The main operating principles that we determined were:

- compatibility with other tools commonly used in data science,
- development of a tool as a framework with open interfaces,
- supporting both multilayer and temporal networks,
- supporting spreading models with discrete states.



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Key Features

Functionalities of network-diffusion:

- end-to-end simulation workflow,
- predefined spreading models,
- an interface for implementing custom spreading models,
- support for the temporal network models (CogSNet + discrete windows),
- support for the multilayer networks,
- centrality measures for multilayer networks.



Key Features

Environmental requirements for network-diffusion:

- support for Linux, macOS, and Windows³,
- Python (preferred 3.10) compatibility,
- C snippets in the CogSNet module to speed-up computations,
- NetworkX compatibility.



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Key Features

To prepare the experiment we have to provide a network, a spreading model and auxiliary parameters. Then, the simulation unfolds as follows:

```
1: procedure PERFORM_PROPAGATION(network, model, epochs)
2:
      states_0 ← model.determine_initial_states()
3:
      model.update_network(states_0)
      for e in [1, ..., epochs] do
4:
5:
              states_e ← model.network_evaluation_step(network)
6:
              model.update_network(network, states_e)
7:
      end for
      logs ← generate logs from experiment
8:
   return logs
10: end procedure
```

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Example I - a Predefined Model

In this example, we will see how to trigger spread under the Linear Threshold Model within a simple, multilayer network.

Linear Threshold Model

Each node:

- can fall in two states: active and inactive,
- becomes active if the fraction of its active neighbors to all neighbours exceeds certain threshold.

In case of multilayer networks the actors not the nodes⁴ are a subject of the diffusion. Thus, we have to define how to aggregate impulses from the layers. In this example we will consider "OR" strategy, which says that the actor can be activated if any of nodes representing it in the network gets activated.

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Lab

⁴which can be considered as avatars of the actors on the network's layers

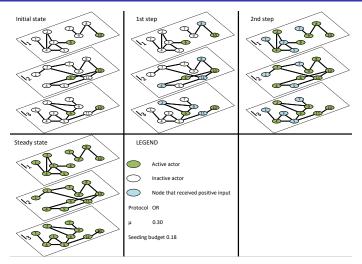


Figure: Propagation according to LTM with the OR protocol in a multilayer network - active actors: seeds $\{6, 10\}$, in a stable state: all of them.

Example I - a Predefined Model

Let's model this problem with network-diffusion!



Example II - a Custom Model

In this example we will consider a joint disease-awareness model (SIR-UA) that can be used e.g. to assess the effectiveness of various countermeasures against the spread of COVID-19 (see the next presentation for details):

Table: Transition weights with explanation.

Symbol	Formula / Value	Description
α	0.19	probability of infection
		for unaware agents
α'	$0.35\alpha = 0.07$	probability of infection
		for aware agents
β	0.10	probability of recovery
$\overline{\gamma}$	0.01	probability of awareness
		for uninfected agents
δ	$\gamma + 1 - 0.3$	probability of awareness
		for infected agents

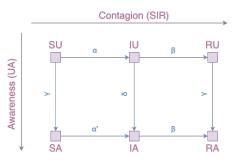


Figure: State and transition graph for SIR-UA.



Example II - a Custom Model

Let's model this problem with network-diffusion!



Resources and References

The library can be installed via:

pip install network-diffusion

Other useful resources have been also published:

- PyPI website: pypi.org/project/network-diffusion
- GitHub page: github.com/anty-filidor/network_diffusion
- Reference guide: network-diffusion.readthedocs.io
- A preprint of the paper: arxiv.org/abs/2405.18085



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Python Implementation

• Advantage and drawback due to slower performance.

Partial C Code

• Some code in C with Python bindings, limited by small team.

Support for Discrete Spreading

• Limited to discrete spreading processes.

Framework Design

- Set of interfaces for custom experiments.
- Few pre-defined spreading models compared to NDlib.

No User Interface

- Not user-friendly for non-programmers.
- Alternatives like NetLogo required.



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Thank you for your attention!



