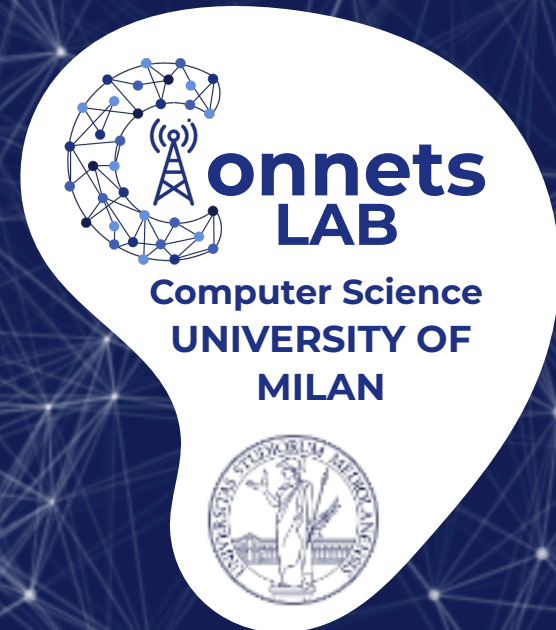
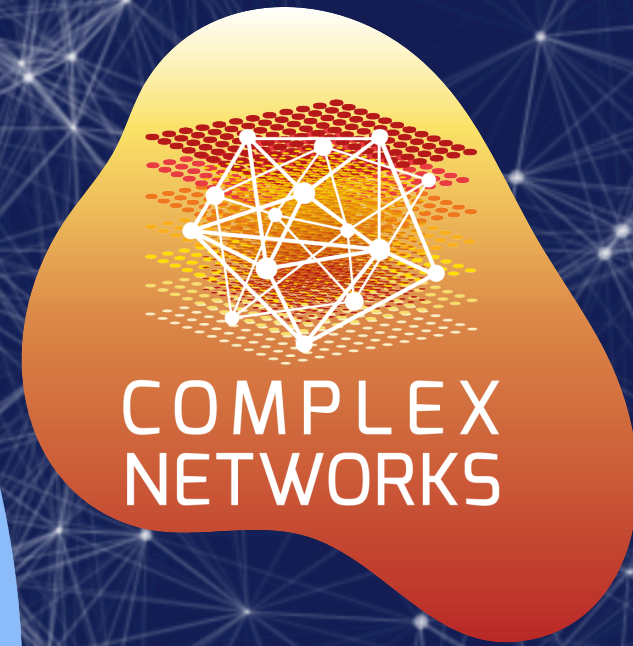


Statistically significant graph evolution rules in temporal networks

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And Sabrina Gaito



Introduction

RESEARCH GROUP WORKS

Network evolution

*Graph evolution rules
Change point/Anomaly detection*

Graph Machine Learning

*Social network analysis using GNN and LLM
Temporal Graph Learning for heterogeneous networks*

User behaviour

*Multilayer community detection
Influence of hubs in a user migration context
User strategies in a reward-based platform*

Web3 platform behavioral and network analysis

*Blockchain-based online social network
NFT networks
Cryptocurrency networks (Luna, Steem, Ethereum, Sarafu)*

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works visit



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The goal



● NETWORK EVOLUTION

● GRAPH EVOLUTION RULES

● STATISTICALLY SIGNIFICANT
GRAPH EVOLUTION RULES

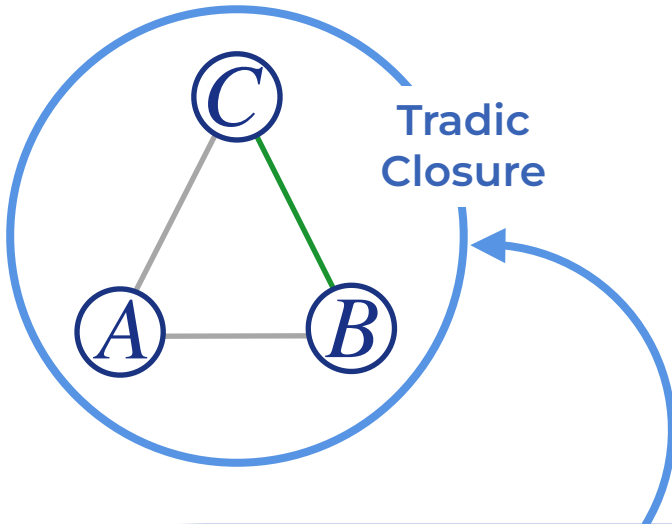
**GRAPH
EVOLUTION RULES**

Background

**MICROCANONICAL
RANDOMIZED
REFERENCE MODELS**

Background

GRAPH EVOLUTION RULES (GER)



Several models, mechanisms and measures have been proposed to describe the network growth

BUT

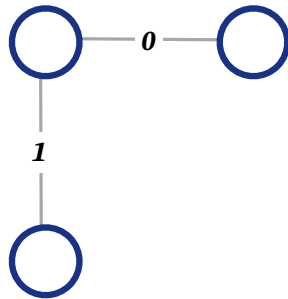
- They assume that the growth is guided by a single parameterized mechanism
- Identifying which mechanism plays a more important role is challenging

Graph evolution rules mining can detect evolutionary behaviors, while avoiding any a-priori mechanism

Graph evolution rules

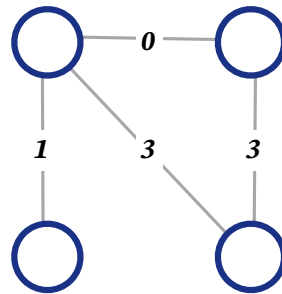
Precondition

Body

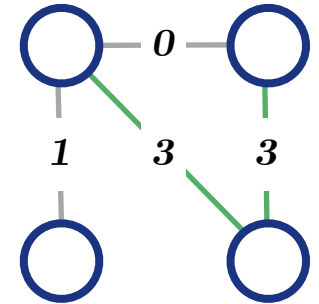


Postcondition

Head



Alternative visualization:



- Similar to association rules in data mining, a GER consists of a precondition (referred to as the *body*) and a postcondition (referred to as the *head*)
- A subgraph matching the body is likely to evolve into one matching the head



Explore structural evolutionary properties

Lack of a null model to measure the significance of rules

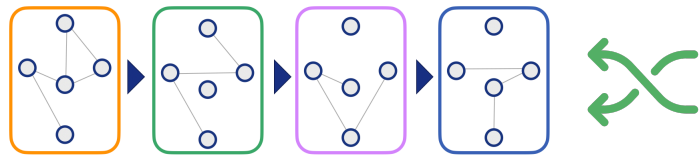


Microcanonical Randomized Reference Models¹

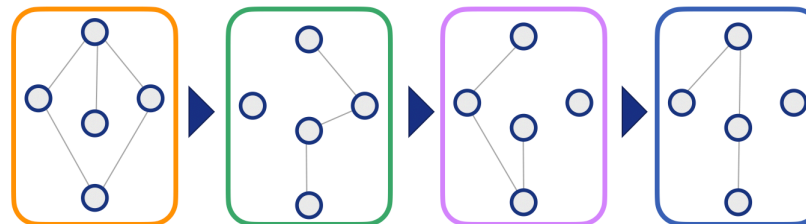


| MRRMs categories | | Timeline representation | Snapshot representation |
|------------------|-----------------------|-------------------------|---------------------------|
| | | What's preserving | Topology |
| | Temporal distribution | <i>Link shuffling</i> | <i>Snapshot shuffling</i> |

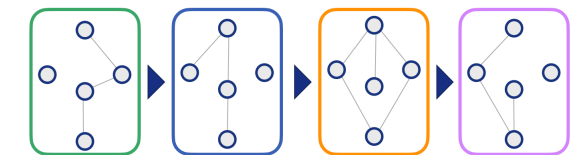
Preserving Temp. Distribution



Original graph



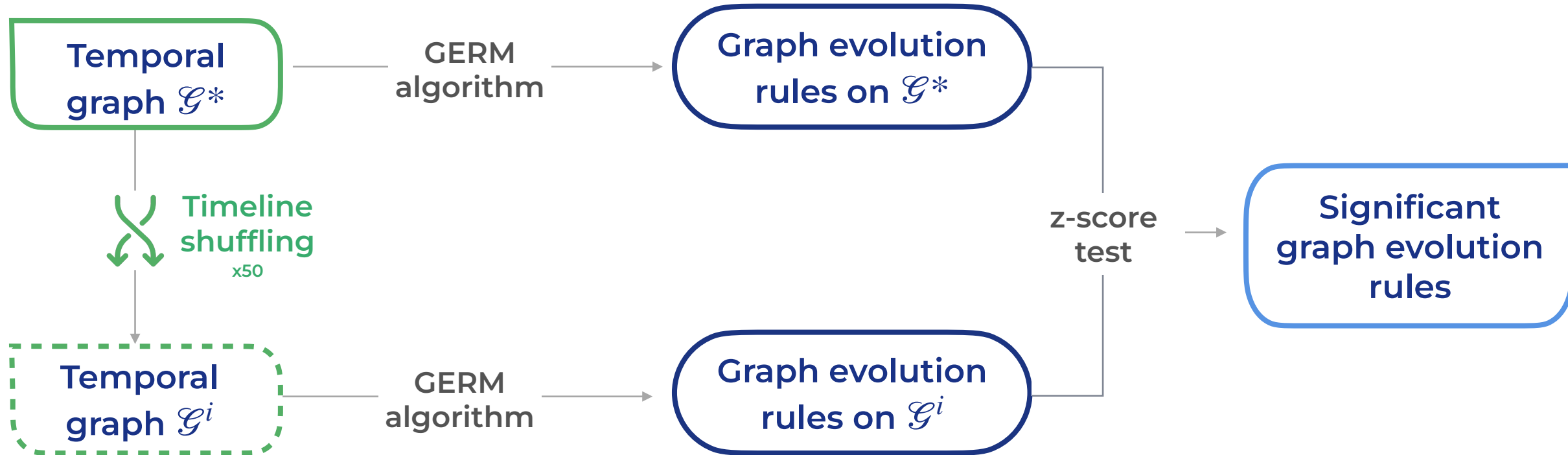
Preserving Topology



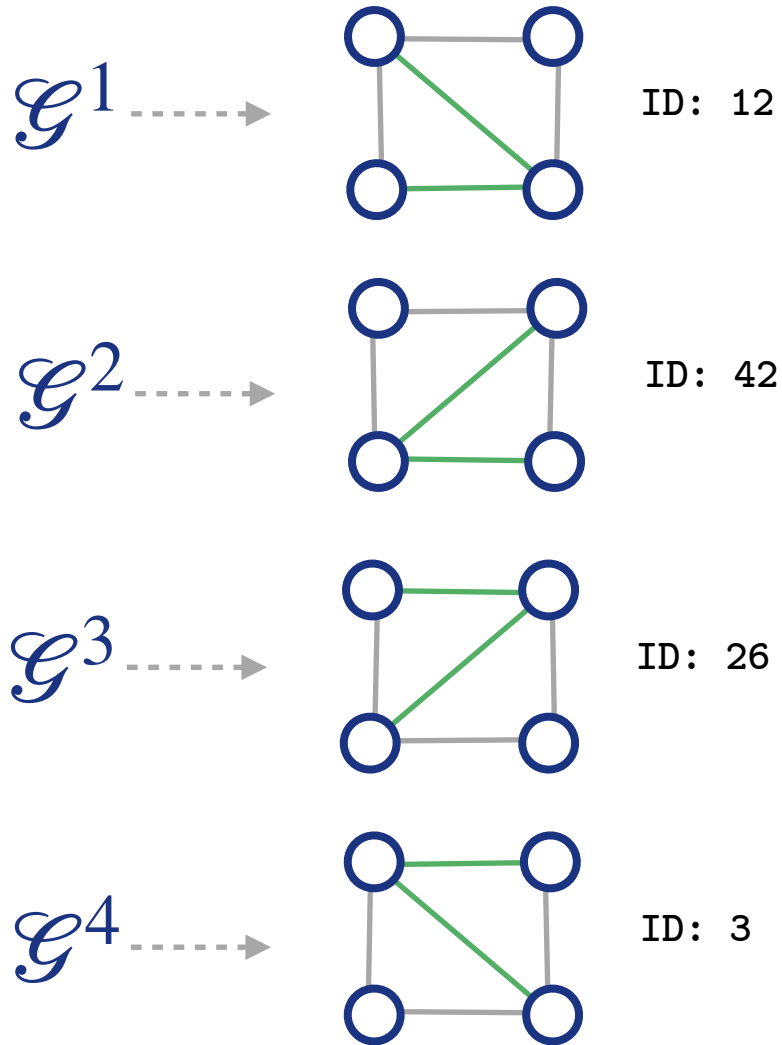
[1] Gauvin, Laetitia, et al. "Randomized reference models for temporal networks." *SIAM Review* 64.4 (2022): 763-830.

Methodology

PIPELINE



General mapping



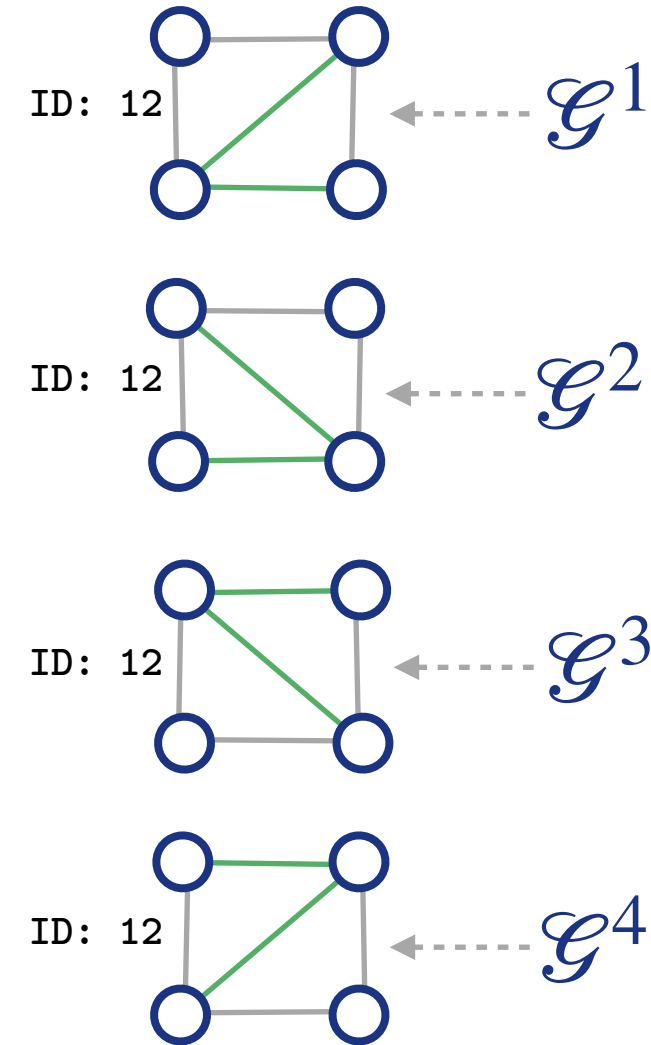
The application of GERM on each realization generates patterns with their respective edge lists and supports, identified by independent incremental IDs.



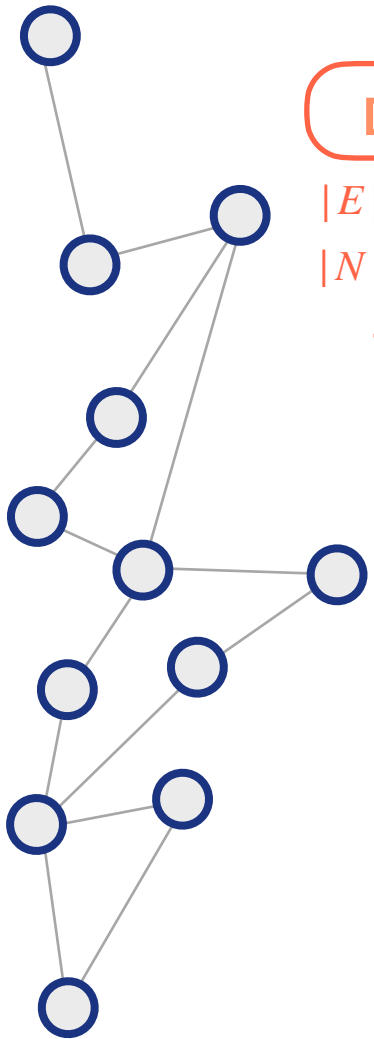
This can create isomorphic rules with different IDs, making impossible the comparison among different random realizations



Create a general mapping, starting from the set of edge list (rules are consistent in number among realizations, so there's a change they're equals), and then checking the isomorphism for each realization



Datasets



DBLP

$|E| = 277081$

$|N| = 129073$

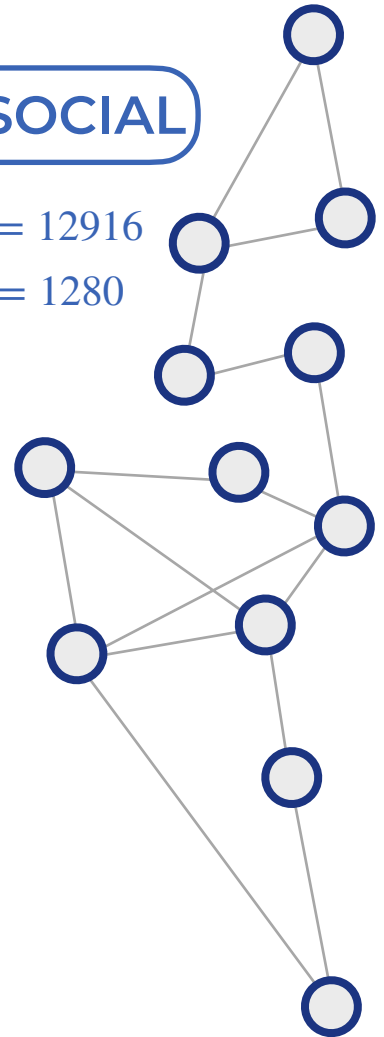
11-years
period

Each operation is a tuple
 (u, v, t)
That records a relationship (co-authorship
or online interaction) between users u and
 v at timestamp t

UC-SOCIAL

$|E| = 12916$

$|N| = 1280$



UC-WEEKLY

28 timestamps

UC-MONTHLY

7 timestamps

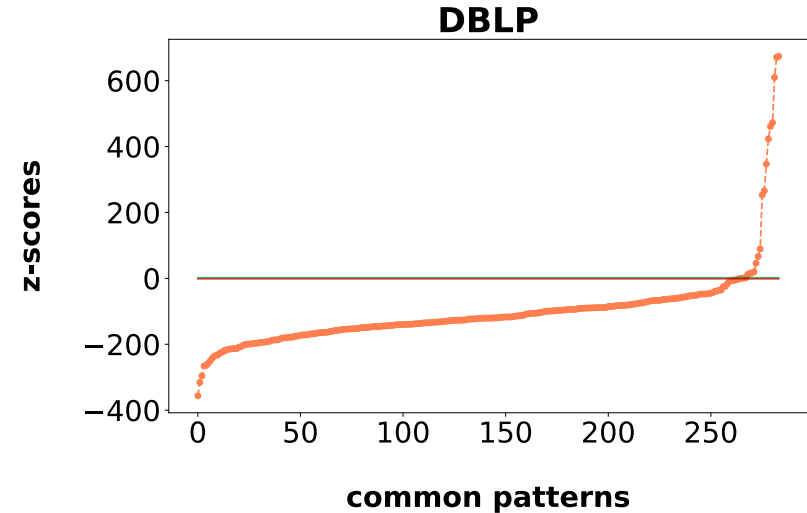
Timestamps
(originally in
microseconds)
are aggregated
at different
granularities

Findings

Outcomes on real and randomize graphs

Graph representation

| Number of frequent rules | Graph representation | | |
|--------------------------|----------------------|---------------------------------------|--|
| | \mathcal{G}^* | $\bar{\mathcal{G}}^i$ Mean shuffle | $\bigcup \mathcal{G}^i$ Union shuffle |
| DBLP | 296 | 3795 | 3871 |
| UC-monthly | 266 | 1269 | 1378 |
| UC-weekly | 999 | 1039 | 1235 |



16

over-represented
(significant) rules

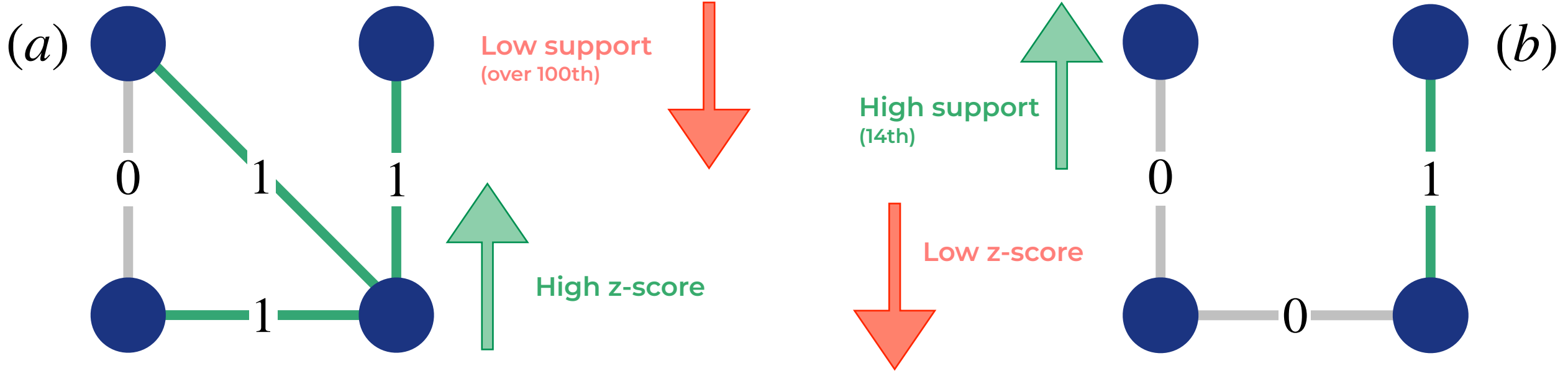
264

under-represented
rules

The rules obtained in the 50 realizations of the null model is stable as the mean and union shuffle count are similar while being very distant from the real graph count

Discussion

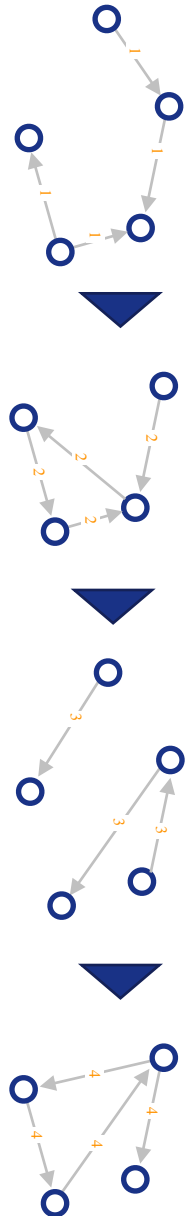
WHAT IF WE DO NOT IMPLEMENT THE NULL MODEL?



Rule apparently **NOT** important but actually **worthy** of attention

Rule apparently **important** but actually **NOT worthy** of attention

Conclusions



Extract GERs
from temporal
graph



Extract GERs
from null model
realizations



Obtain
statistically
significant rules

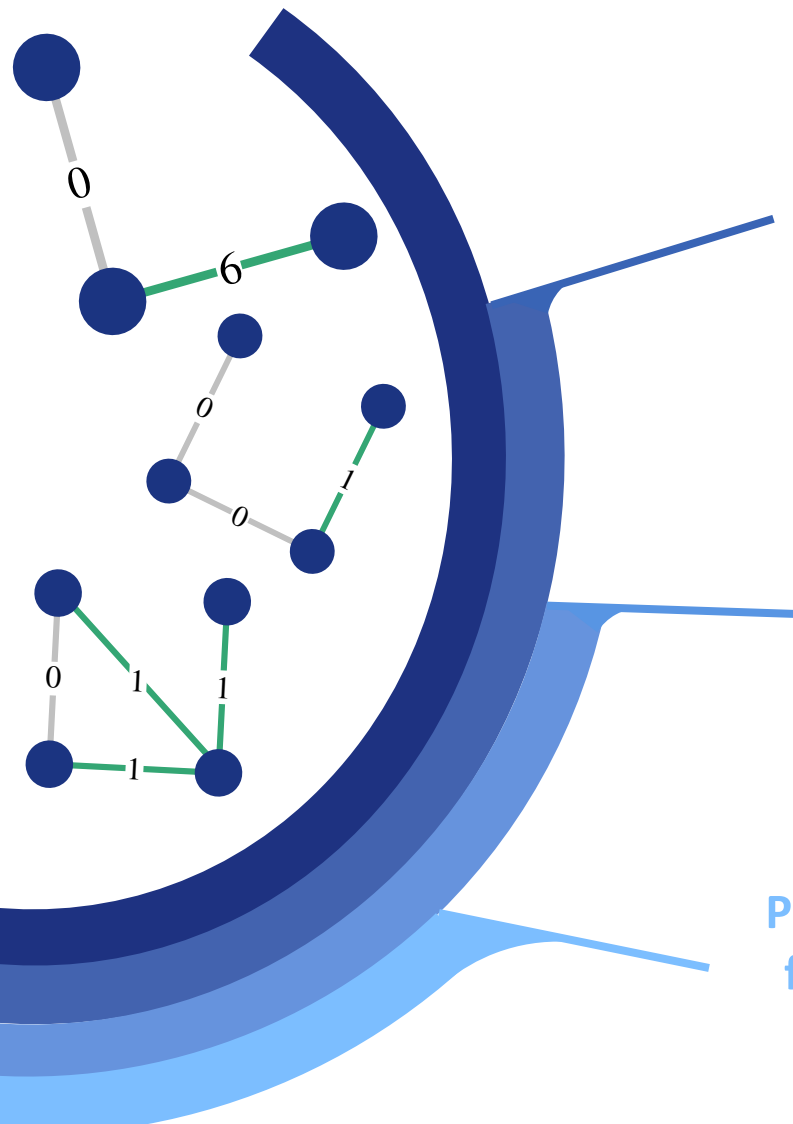


Comprehensive
analysis of
results

We saw that:

- By shuffling timestamps, we weaken some temporal constraints on the original graph
- Some rules are frequent but not significative and viceversa

Future and ongoing works



Factors of under/over representation

Identifying the mechanisms and the factors acting on the over and under-representation of the GERs

Extension to other GER algorithms

Apply the same methodology to other state-of-the-art GER algorithms

Python toolkit for GER easy usage:
GERANIO

Recently released a python toolkit that helps applying, drawing, and analysing graph evolution rules.



Graph Evolution Rules ANalytics vIsualization tOolkit





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<https://alessiaatunimi.github.io/>

Thanks for your
attention

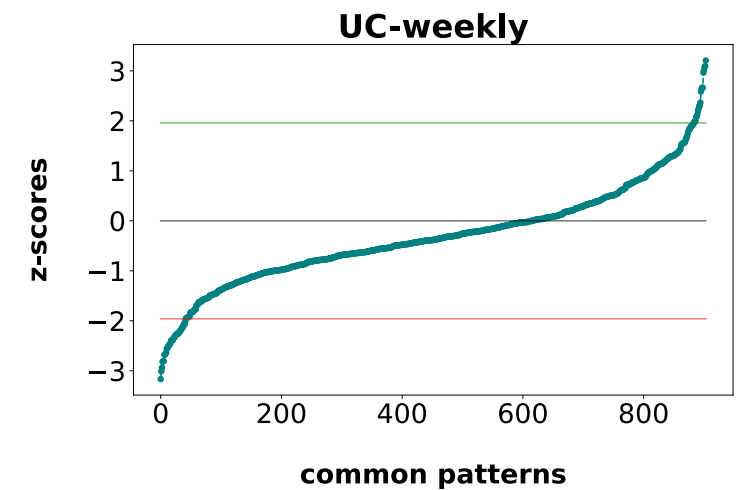
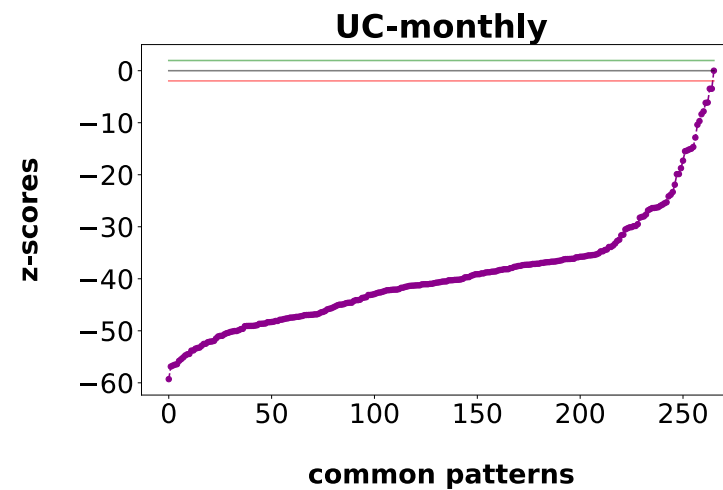
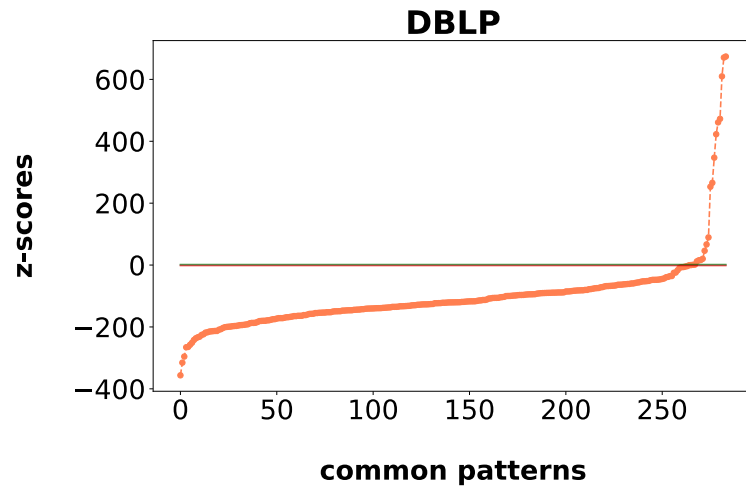
Research Lab website:
<https://connets.di.unimi.it/>



This work is a short version of the paper
*"Unfolding temporal networks
through statistically significant
graph evolution rules"*

accepted at DSAA2023

z-score results



General mapping

Algorithm 2 General mapping: set of edges

Input: results of GERM on each realization of the null model
germ

Output: *edges_set*

```
1: edges_set = dict()
2: m = 0
3: for model ∈ germ do
4:   for p, pattern ∈ model do
5:     push(edges_set[patternedges], (p, m))
6:   end for
7:   m ← m + 1
8: end for
```

Algorithm 3 General mapping: isomorphism check

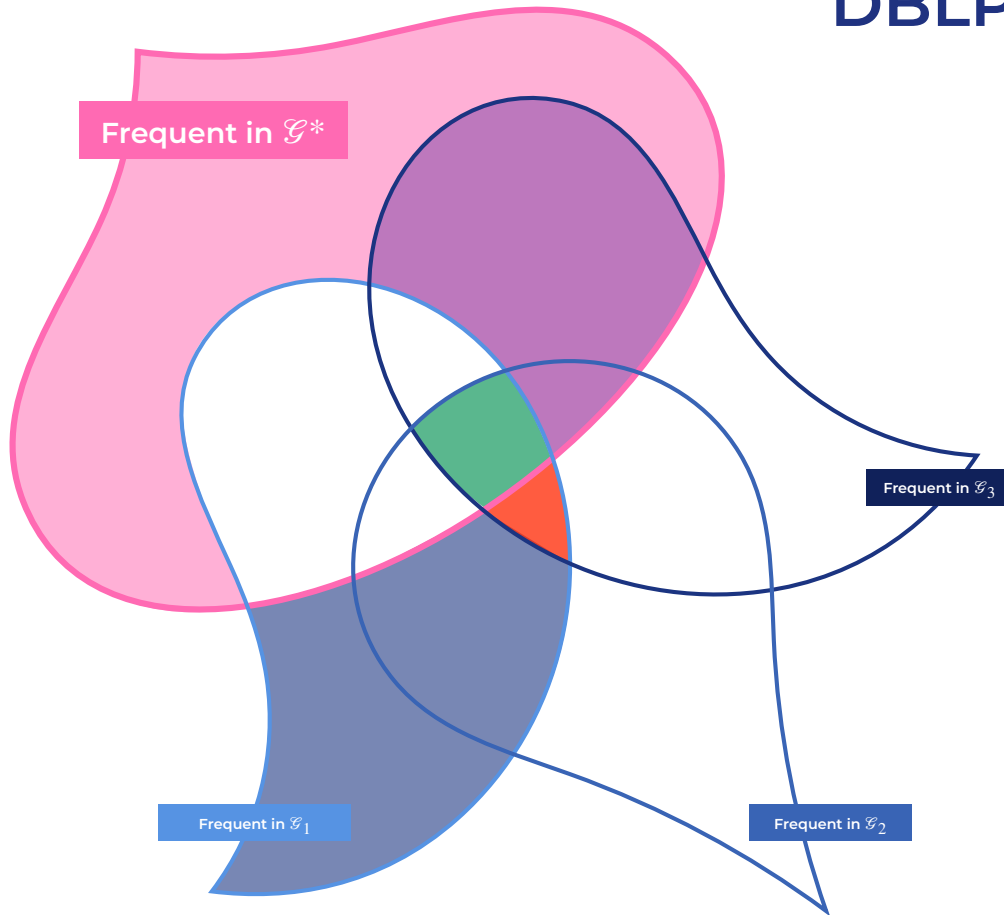
Input: *edges_set*

Output: *new_shuffle_germ*

```
1: mapping = dict()
2: new_shuffle_germ = dict()    ▷ Inspired by Python,
   dict() creates an associative map
3: i = max(mapping.keys)
4: for edges, occurrences ∈ edges_set do
5:   id ← 0
6:   for p, pattern ∈ mapping do
7:     if G(edges) is_isomorphic G(pattern) then
8:       id ← p
9:       continue
10:    end if
11:  end for
12:  if id = 0 then
13:    i ← i + 1
14:    id ← i
15:    mapping[i] = G(edges)
16:  end if
17:  for p, m ∈ occurrences do
18:    new_shuffle_germ[m][id] = germ[m][p]
19:  end for
20: end for
```

Other set of rules to consider

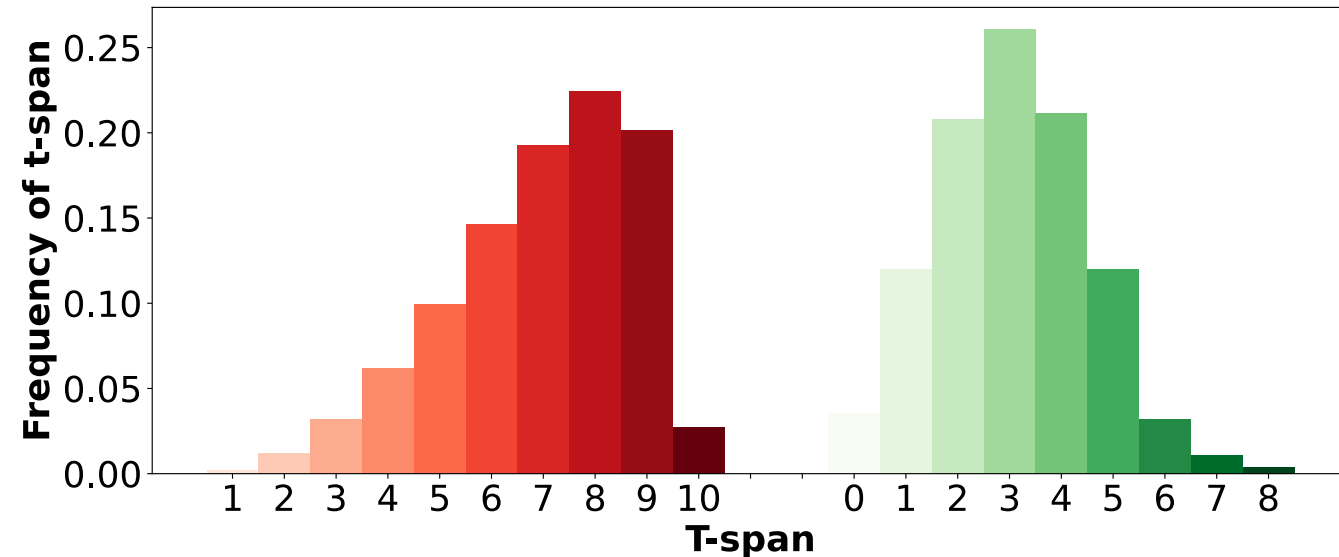
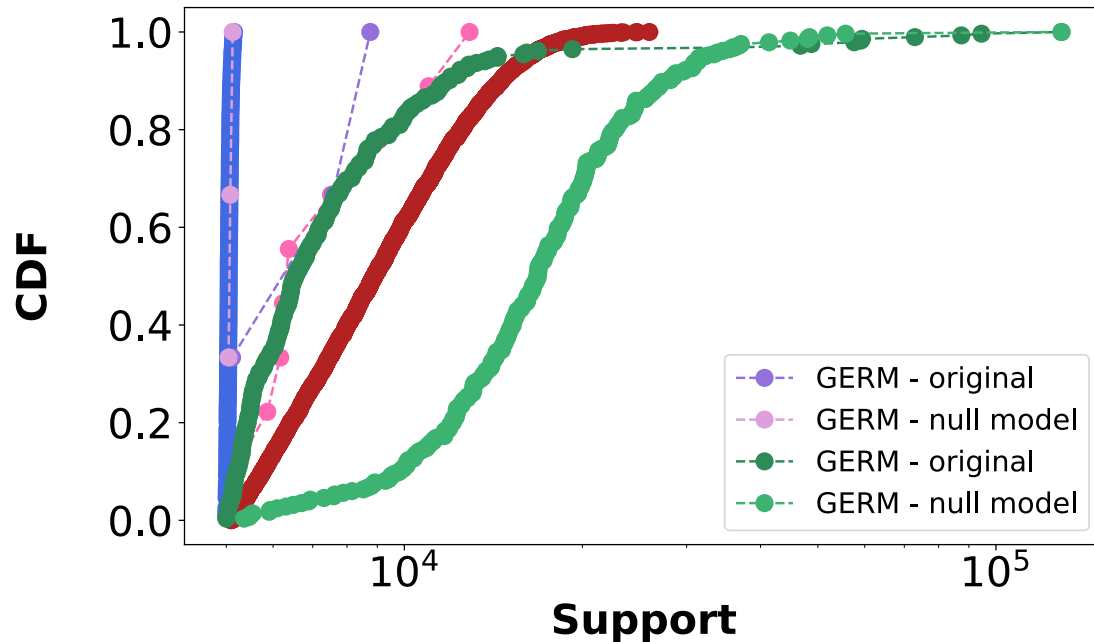
DBLP CASE STUDY



| | | Frequent in GERM | | <i>TOT</i> |
|---|------------------|------------------|--------------|-------------|
| | | Frequent | Not frequent | |
| Frequent in how many realizations of the null model | DBLP | | | |
| | Frequent in all | 284 | 3431 | 3715 |
| | Frequent in some | 3 | 153 | 156 |
| Not frequent | 9 | | | |
| <i>TOT</i> | 3871 | 296 | | |

Do other set tell something about evolution?

DBLP CASE STUDY



References

¹E. Scharwa"chter, E. Mu"ller, J. Donges, M. Hassani, and T. Seidl, "Detecting change processes in dynamic networks by frequent graph evolution rule mining," in *2016 IEEE 16th International Conference on Data Mining (ICDM)*. IEEE, 2016, pp. 1191–1196.