

Link Reliability

Missing and spurious interactions in complex networks

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DI PADOVA

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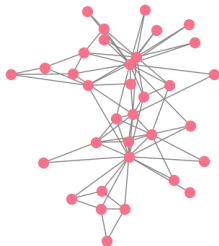
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- Network science aims to unfold the functional needs of a system by looking at the **interactions between units**.
- Unfortunately, the **reliability of network** data is often a source of concern.
- In this presentation, we will examine a framework to assess the reliability of complex networks, based on **Bayesian inference**.

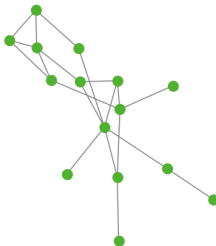
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We compare three **different networks** with different numbers of nodes, structures, and complexities:

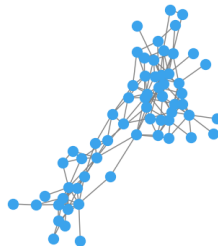
Karate Club



Florentine Families



Dolphins



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- Is analytically and computationally **tractable**.

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Stochastic block model

A block model $M = (P, \mathbf{Q})$ is completely defined by the partition P of nodes into groups and the matrix \mathbf{Q} of probabilities of connections between groups.

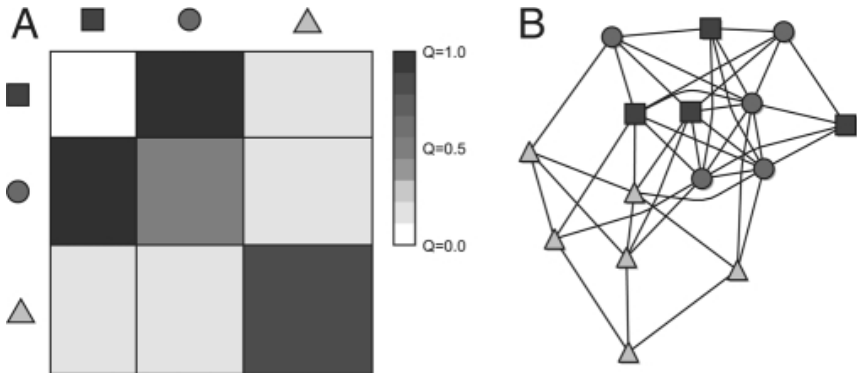


Figure: (A) Probability matrix Q of a SBM $M = (P, Q)$, being $P = (4, 5, 6)$.
(B) A realization of the model described in A

Considering an **observed network** A^O and a set of generative models \mathcal{M} , we can compute the probability $p(X = x|A^O)$ for an **arbitrary network property** X as:

$$p(X = x|A^O) = \int_{\mathcal{M}} dM p(X = x|M) p(M|A^O)$$

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Using the **Bayes theorem** we can rewrite

$$p(M|A^O) = \frac{p(A^O|M)p(M)}{\int_{\mathcal{M}} dM' p(A^O|M')p(M')}$$

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The likelihood of each model M can be written as:

$$p(A^O|P, Q) = \prod_{\alpha \leq \beta} Q_{\alpha\beta}^{I_{\alpha\beta}^O} (1 - Q_{\alpha\beta})^{r_{\alpha\beta} - I_{\alpha\beta}^O}$$

where $I_{\alpha\beta}^O$ is the number of links in A^O between nodes in groups α and β of P , and $r_{\alpha\beta}$ is the maximum number of such links.

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We can assume an uninformative prior, i.e.:

$$p(P, Q) \sim \text{const}$$

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Then, we can study the property of having a link between nodes i and j , given their groups σ_i and σ_j in P :

$$p(A_{ij} = 1|P, Q) = Q_{\sigma_i \sigma_j}$$

We can write the **link reliability** $R_{ij}^L = p(A_{ij} = 1|A^O)$ as:

$$R_{ij}^L = \frac{1}{Z} \sum_{P \in \mathcal{P}} \left(\frac{l_{\sigma_i \sigma_j} + 1}{r_{\sigma_i \sigma_j} + 2} \right) \exp[-\mathcal{H}(P)]$$

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where the hamiltonian $\mathcal{H}(P)$ is

$$\mathcal{H}(P) = \sum_{\alpha \leq \beta} \left[\ln(r_{\alpha\beta} + 1) + \ln \left(\frac{r_{\alpha\beta}}{l_{\alpha\beta}^O} \right) \right]$$

and Z is the normalization.

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In practice, it's impossible to sum over all the possible partitions.

We can treat R_{ij}^L as an **ensemble average** and use the **Metropolis algorithm** to sample the relevant contributions.

Sampling procedure (1)



We start initializing the N nodes into N groups, uniformly at random:

```
# Groups data structure: list of lists
N = A.shape[0]
groups = [[] for _ in range(N)]

# Uniformly random initialization
for i in range(N):
    g = np.random.randint(0, N)
    groups[g].append(i)

# Compute the hamiltonian
H = hamiltonian(A, groups)
```

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At each step, we select a random node and attempt to move it into a new random group:

```
# Randomly select a node and a group
i = np.random.randint(0, N)
g_prop = np.random.randint(0, N)

# Move the node to another group
groups_prop = swap(groups, i, g_prop)
```

Then, we compute $\Delta\mathcal{H}$:

- If $\Delta\mathcal{H} \leq 0$ we accept the change
- Otherwise, the change is accepted with probability $\exp(-\Delta\mathcal{H})$

```
# Compute the Hamiltonian of the new configuration
H_prop = hamiltonian(A, groups_prop)

# Acceptance probability
if H_prop <= H:
    groups = groups_prop
    H = H_prop
else:
    r = np.random.rand()
    if r < np.exp(H - H_prop):
        groups = groups_prop
        H = H_prop

return groups, H
```

Sampling procedure (3)



The sampling procedure starts after an **equilibration period**.

```
# Transient
for _ in range(transient):
    groups, H = singleStep(groups, H, A)
```

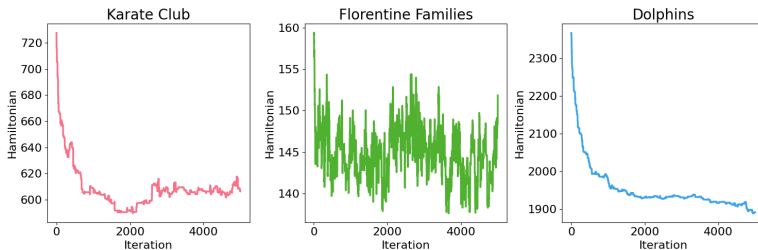


Figure: Transient time

Sampling procedure (4)



The sampling procedure must consider only **uncorrelated partitions**.

```
for k in range(n_samples):  
    for _ in range(delay):  
        groups, H = singleStep(groups, H, A)  
        partitions_set.append(groups)  
        hamiltonians_list.append(H)
```

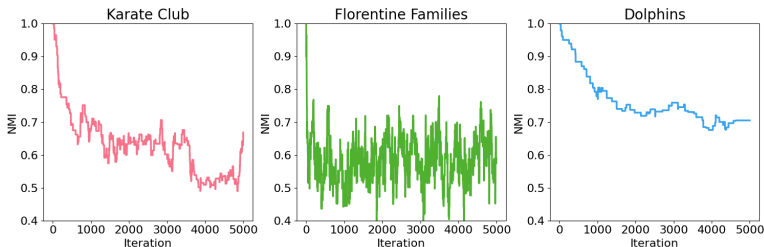


Figure: Normalized Mutual Information

Offset

You can always **rescale the hamiltonian** by choosing an offset.

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- This is meant to avoid roundoff errors and vanishing information.
- It allows you to work with bigger networks.
- Working with huge networks would require studying the fluctuations around the mean.

```
np.exp(-np.array(hamiltonian_list, dtype=np.float128) + offset)
```

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Sampling procedure (6)



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```
with multiprocessing.Pool(processes=n_cores) as pool:  
    results = pool.starmap(samplingBranch, input)
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Sampling procedure (6)



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with multiprocessing.Pool(processes=n_cores) as pool:  
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```
0[|||||||||||||||||||||100.0%] 4[|||||||||||||||||||||100.0%]  
1[|||||||||||||||||||||100.0%] 5[|||||||||||||||||||||100.0%]  
2[|||||||||||||||||||||100.0%] 6[|||||||||||||||||||||100.0%]  
3[|||||||||||||||||||||100.0%] 7[|||||||||||||||||||||100.0%]  
Mem[||||||| 1.66G/15.6G Tasks: 54, 127 thr; 8 running  
Swp[| 32.1M/8.00G Load average: 1.83 0.42 0.17  
Uptime: 50 days, 22:56:49
```

Figure: A snapshot of the system displayed by `htop` in the terminal.

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Spurious interactions: A^O is obtained by adding $\lceil f E \rceil$ edges to A^T

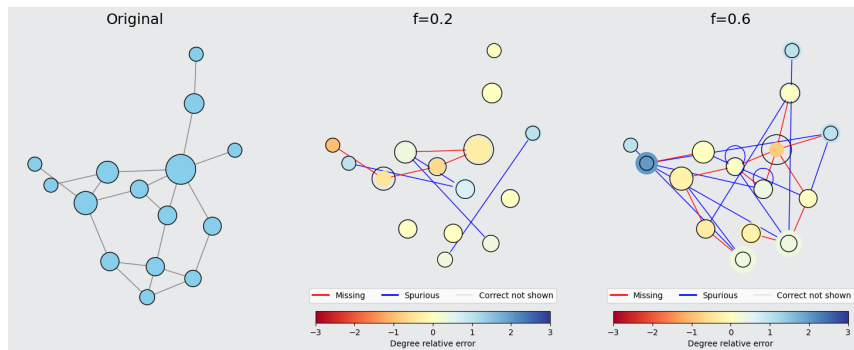
Corrupt a graph



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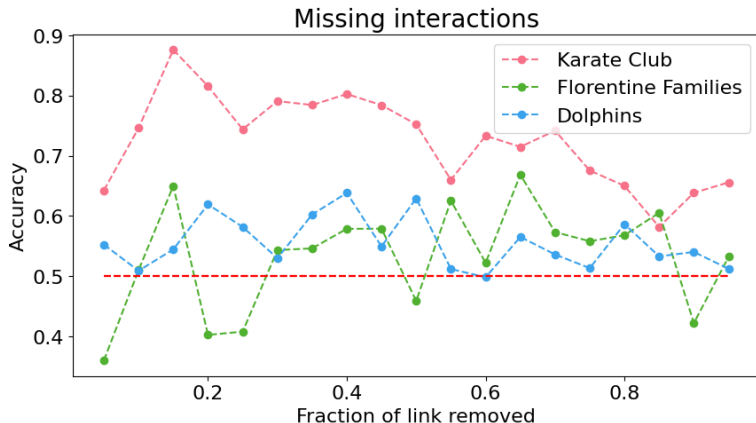


To test the ability to identify **missing interactions**, we compute the probability that *a false negative has a higher reliability than a true negative*.

Test link reliability (1)



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Test link reliability (2)

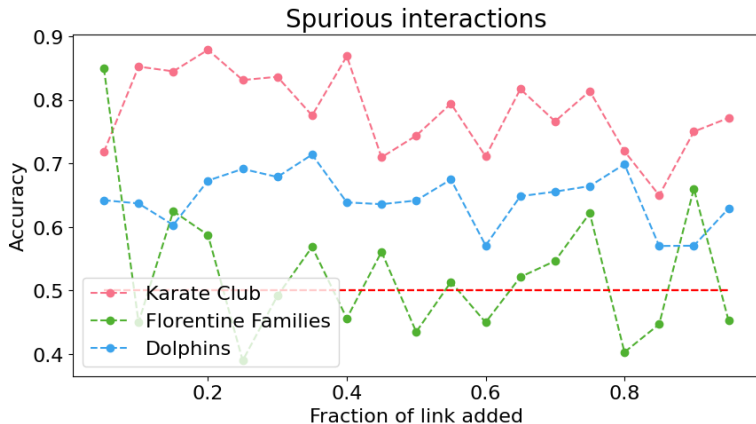


To test the ability to identify **spurious interactions**, we compute the probability that *a false positive has a lower reliability than a true positive*.

Test link reliability (2)



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Similarly, one can test the **network reliability** $R_A^N = p(A|A^O)$ that can be written as:

$$R_A^N = \frac{1}{Z} \sum_{P \in \mathcal{P}} h(A; A^O, P) \exp(-\mathcal{H}(P))$$

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where

$$h(A; A^O, P) = \exp \left\{ \sum_{\alpha \leq \beta} \left[\ln \left(\frac{r_{\alpha\beta} + 1}{2r_{\alpha\beta} + 1} \right) + \ln \left(\frac{\binom{r_{\alpha\beta}}{l_{\alpha\beta}^O}}{\binom{2r_{\alpha\beta}}{l_{\alpha\beta} + l_{\alpha\beta}^O}} \right) \right] \right\}$$

and

$$\mathcal{H}(P) = \sum_{\alpha \leq \beta} \left[\ln(r_{\alpha\beta} + 1) + \ln \left(\binom{r_{\alpha\beta}}{l_{\alpha\beta}^O} \right) \right]$$

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Optimal network

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Unfortunately, a full scan in the configuration space is unfeasible. We need to define an **heuristic maximization method**.

```

hyper = (n_samples, delay, transient) # Hyperparameters
A_cur = A_obs.copy() # Initialize the current adjacency matrix

for j in range(5): # This is the maximum number of iterations
    # Compute the network reliability and the link reliability matrix
    R_N = getNetworkReliability(A_cur, A_obs, *generatePartitionsSet(A_obs, *hyper))
    R_L = computeLinkReliabilityMatrix(A_cur, *generatePartitionsSet(A_cur, *hyper))

    # Sort the links (increasing R) and not links (decreasing R)
    sorted_links, sorted_not_links = sortLinkLists(*getLinkLists(A_cur), R_L)

    # Iterate to reconstruct the network
    num_iters = np.min([len(sorted_links), len(sorted_not_links)])
    miss_update = 0
    done_update = False
    for k in range(num_iters):
        # Choose a pair of link and not link in order
        link, not_link = sorted_links[k], sorted_not_links[k]

        # Define the proposed adjacency matrix
        A_temp = A_cur.copy()

        # Swap the links
        A_temp[link[0], link[1]] = 0
        A_temp[link[1], link[0]] = 0
        A_temp[not_link[0], not_link[1]] = 1
        A_temp[not_link[1], not_link[0]] = 1

        # Compute the network reliability with the new adjacency matrix
        R_N_temp = getNetworkReliability(A_temp, A_obs, *generatePartitionsSet(A_obs, *hyper))

        # If the network reliability increases, update the adjacency matrix
        if R_N_temp > R_N:
            A_cur = A_temp
            R_N = R_N_temp
            miss_update = 0
            done_update = True
        else:
            miss_update += 1
            # Break condition for the iteration
            if miss_update > 4:
                break

    # Break condition for the whole loop
    if not done_update:
        break

```

Rationale of network reconstruction

The crucial assumption is that $R_{AT}^N \gg R_{Ao}^N$.

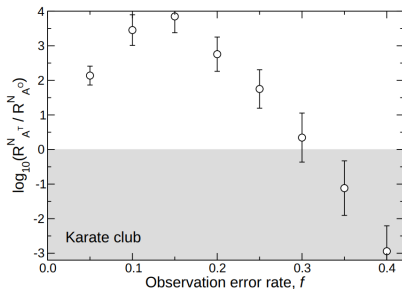


Figure: From the *Supporting Information* of Guimera and Sales-Pardo.

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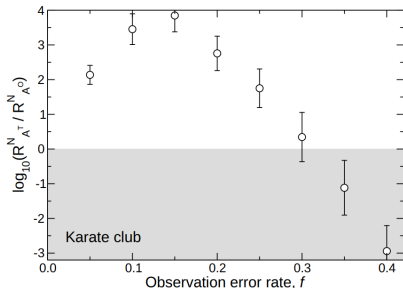


Figure: From the *Supporting Information* of Guimera and Sales-Pardo.

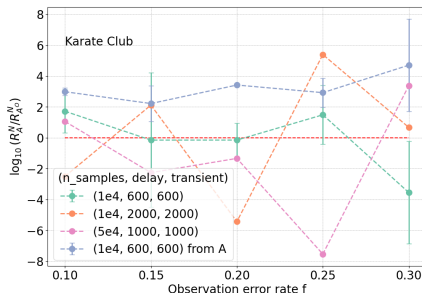
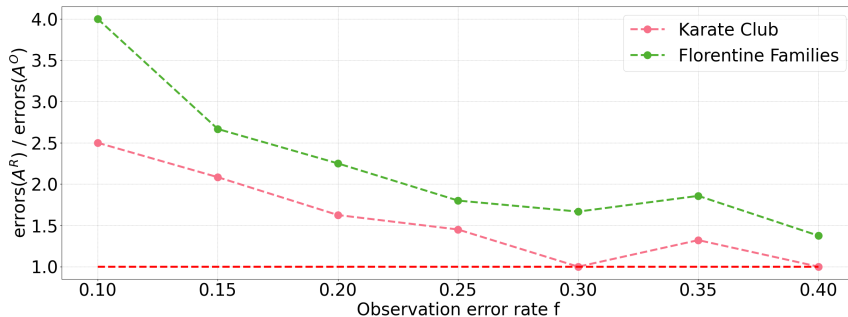


Figure: Preliminary studies on network reconstruction.

Network reconstruction (4)



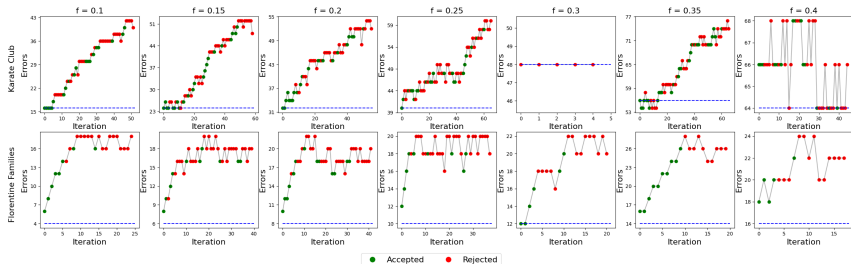
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Network reconstruction (4)



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The **network reconstruction framework** should be debugged by:

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- changing the underlying groups' data structure to speed up the computations and the debugging

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Moreover, one can study **other optimizations**:

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- dynamic decorrelation time based on NMI

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Moreover, one can study **other optimizations**:

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Finally, it would be **interesting to study**:

- the role of the network topology with synthetic data
- the impact of possible node metadata

We review this **bayesian framework** to test **interactions in complex networks**.

- We developed the notion of reliability in the context of the SBM.
- We implemented a sampling procedure to compute the ensemble averages.
- We studied the link reliability for missing and spurious interactions.
- We introduced the network reliability.

