

Maximum Likelihood Estimation and Graph Matching in Errorfully Observed Networks

Jesús Arroyo

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Joint work with:



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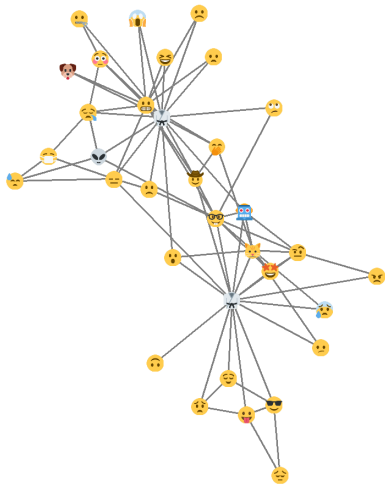


Carey E. Priebe
(Johns Hopkins University)

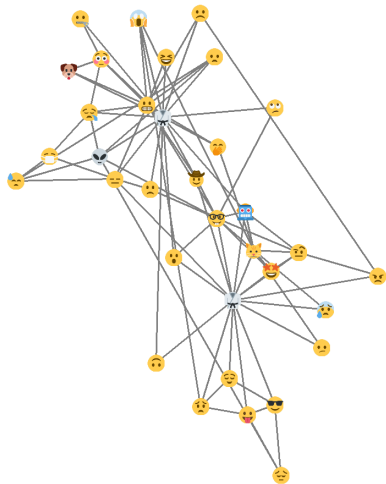
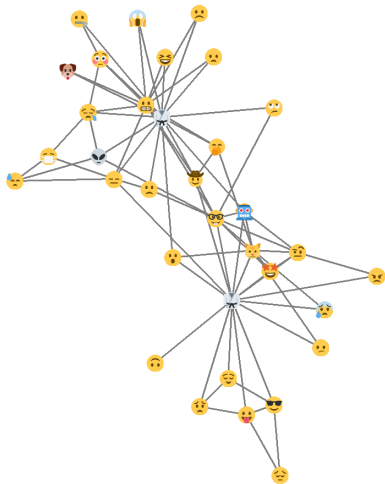


Vince Lyzinski
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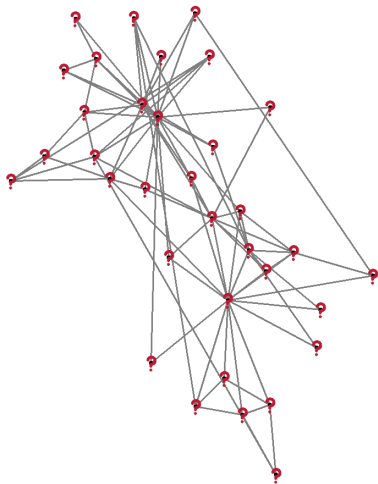
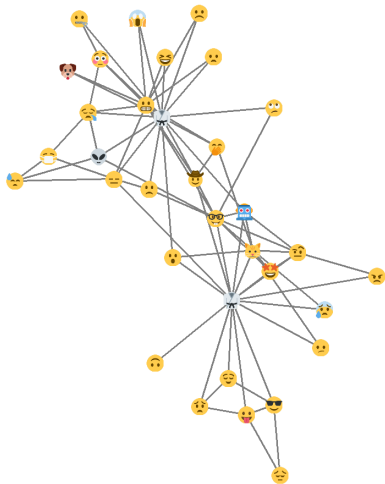
Graph matching



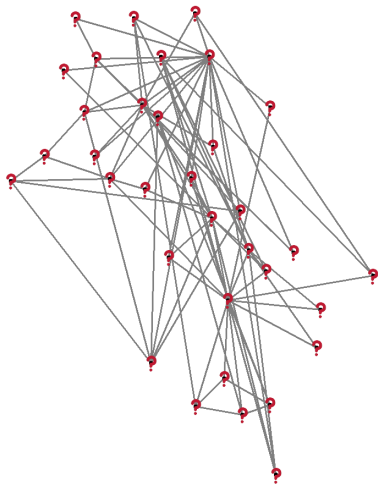
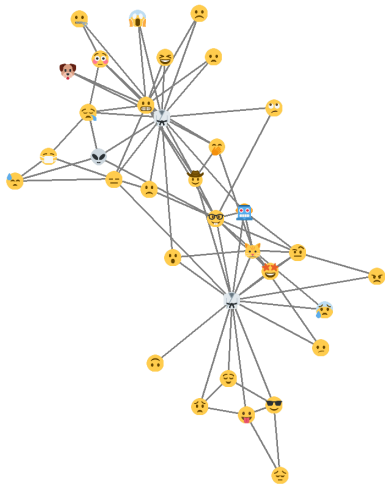
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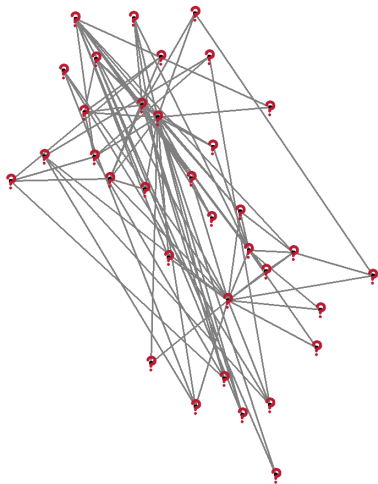
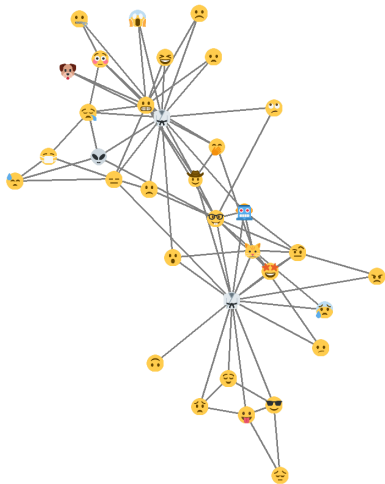
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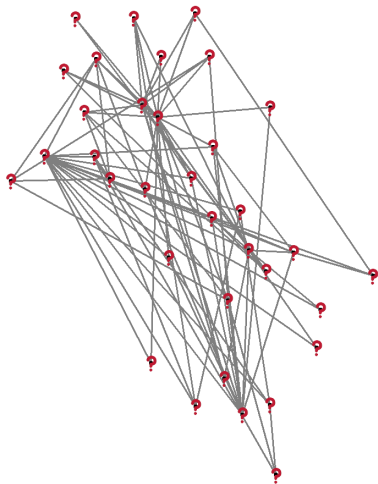
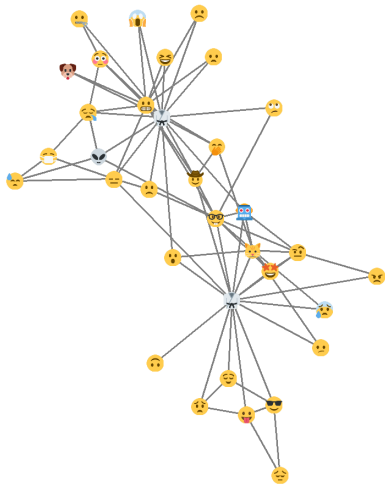
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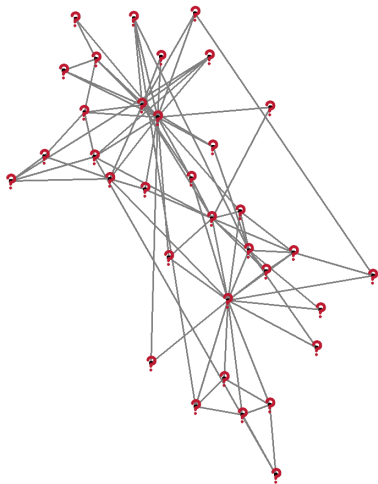
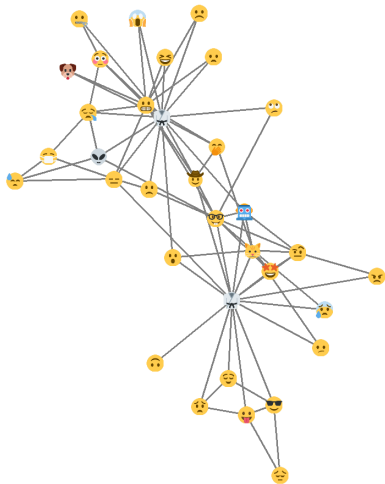
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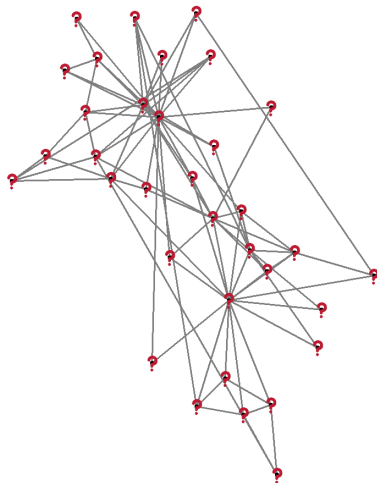
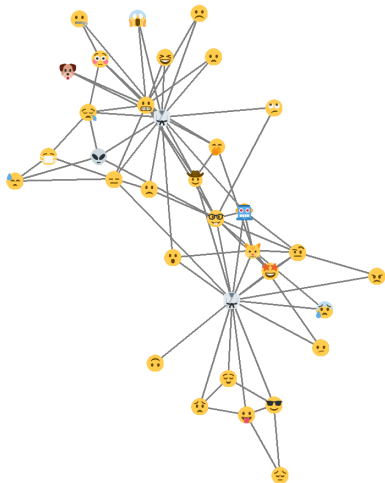
Graph matching



Graph matching



Graph matching



- How can we formulate a statistical model for graph matching?
- When is statistically feasible to solve the problem?

Graph matching problem

Goal: find a meaningful correspondence between the vertices of the graphs

- Typically defined as the permutation that minimizes the number of edge disagreements:

$$\begin{aligned}\hat{Q}_{\text{GM}} &= \operatorname{argmin}_{Q \in \Pi_n} \|A - QBQ^T\|_F^2 \\ &= \operatorname{argmin}_{Q \in \Pi_n} \sum_{i \neq j} (A_{ij} - B_{\sigma(i), \sigma(j)})^2\end{aligned}$$

- ▶ $A, B \in \{0, 1\}^{n \times n}$ adjacency matrices
- ▶ Q is a permutation matrix, σ is the associated permutation function

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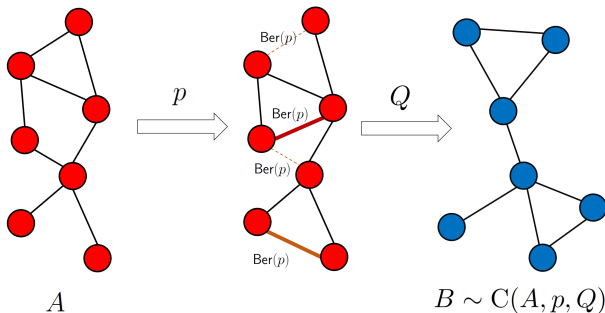
Some applications:

- Aligning biological networks
- De-anonymization in social networks
- Unsupervised word translation

Corrupting channel model

Model: B is an edge and vertex-label corrupted version of A

- 1 Edges and non-edges of A are independently flipped with probability p .
- 2 Vertices are shuffled with a permutation Q .



Maximum likelihood estimation

- Consider the **maximum likelihood estimator** for the parameters

$$(\hat{p}_{\text{MLE}}, \hat{Q}_{\text{MLE}}) := \operatorname{argmax}_{p, Q} \sum_{u > v} \log \mathbb{P}_p (A_{uv} = (QBQ^T)_{uv}) .$$

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- Extensions of the model to non-uniform corrupting probabilities
 - This result also holds with non-uniform corrupting probabilities.

When is the MLE correct?

- The difficulty of the problem depends on how different is A from QAQ^T for any $Q \neq I$, measured by

$$\|A - QAQ^T\|_F^2 = \sum_{i \neq j} (A_{ij} - A_{\sigma(i), \sigma(j)})^2.$$

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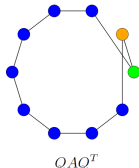
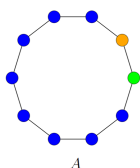
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- $\Pi_{n,k}$ is the set of permutations that permute exactly k nodes.

Examples:

- $k = 2$
- $\|A - QAQ^T\|_F^2 = 4.$



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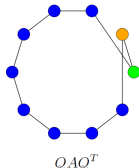
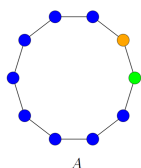
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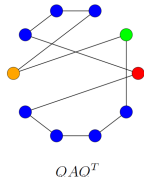
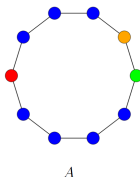
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- $k = 3$

- $\|A - QAQ^T\|_F^2 = 10$.



Consistency of the MLE

Consider a sequence of networks $\{A_n\}$ with n vertices, and parameters $\{p_n, Q_n\}$

Definition

$\{\hat{Q}_n\}$ is **consistent** if $\|\hat{Q}_n - Q_n\|_F \xrightarrow{a.s.} 0$ as $n \rightarrow \infty$.

Consistency: all vertices are correctly matched in the limit.

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Theorem

- \hat{Q}_{MLE} is **consistent** if

$$\min_{Q \in \Pi_{n,k}} \|A_n - Q A_n Q^T\|_F^2 \geq \frac{6k \log n}{(1/2 - p_n)^2}, \quad \forall k \geq 2.$$

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- \hat{Q}_{MLE} is **not consistent** if there exists $m = \Omega(n)$ disjoint permutations Q_1, \dots, Q_m

$$\max_{i \in [m]} \|A_n - Q_i A_n Q_i^T\|_F^2 = o\left(\frac{\log n}{(1/2 - p_n)^2}\right).$$

Relaxed consistency of the MLE

Matching all vertices correctly can be hard or impossible, for example:

- low-degree vertices
- automorphisms in the graph

Definition (Relaxed consistency)

$\{\hat{Q}_n\}_{n=n_0}^{\infty}$ is $\{k_n\}$ -consistent if $\frac{\|\hat{Q}_n - Q_n\|_F}{k_n} \xrightarrow{a.s.} 0$ as $n \rightarrow \infty$

Relaxed consistency: at most $o(k_n)$ misaligned vertices.

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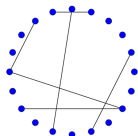
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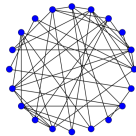
$$\min_{Q \in \Pi_{n,k}} \|A_n - QA_nQ^T\|_F^2 \geq \frac{6k \log n}{(1/2 - p_n)^2}, \quad \forall k \geq k_n.$$

Consistency in random graph models

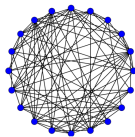
Erdős-Rényi graph model: $A_n \sim G(n, \alpha_n)$



$\alpha = 0.05$



$\alpha = 0.2$



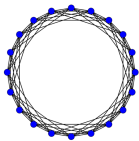
$\alpha = 0.4$

Corollary

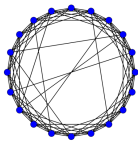
\hat{Q}_{MLE} is **consistent** if

$$\alpha_n \geq \frac{c_1 \log n}{n(1/2 - p_n)^2}.$$

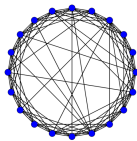
Newman-Watts small world graph model: $A_n \sim \text{NW}(n, d_n, \beta_n)$,



$d = 4,$
 $\beta = 0$

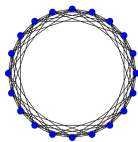


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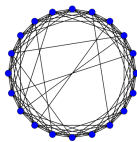


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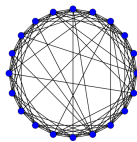
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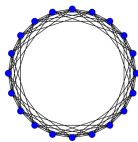
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Corollary

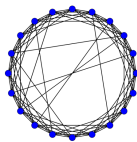
- For any $\{d_n\}$, \hat{Q}_{MLE} is **not consistent** if $(1/2 - p_n)^2 = o(\sqrt{\log n/n})$ and

$$\beta_n = o\left(\frac{\log n}{(1/2 - p_n)^2 n}\right).$$

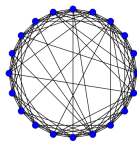
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Corollary

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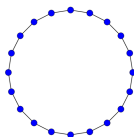
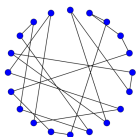
$$\beta_n = o\left(\frac{\log n}{(1/2 - p_n)^2 n}\right).$$

- \hat{Q}_{MLE} is **consistent** if $d_n = o(\beta_n^2 n)$ and

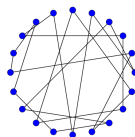
$$\beta_n \geq C \sqrt{\frac{\log n}{n(1/2 - p_n)^2}}.$$

Relaxed consistency

Random regular graphs: $A \sim G_{n,d_n}$



$d = 2$



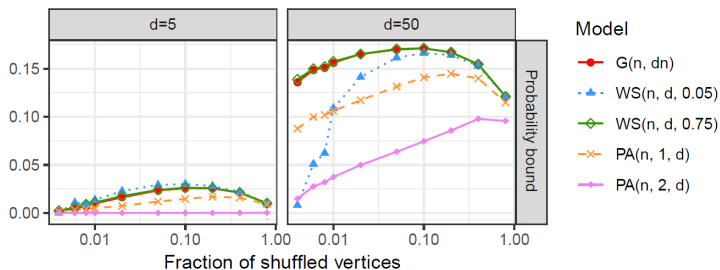
Corollary

If $d_n = c_1 n^{2/3+\epsilon}$ and $(\frac{1}{2} - p_n)^2 \geq c_2 \frac{\log n}{d_n}$, then \hat{Q}_{MLE} is $\{n^{2/3+\epsilon}\}$ -consistent.

A negligible fraction of the vertices ($o(n^{-(1/3-\epsilon)})$) might be incorrectly matched.

Matchability on random graphs

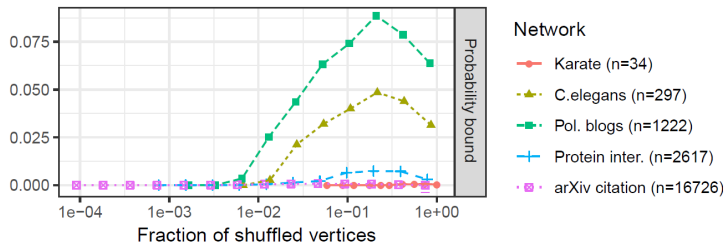
Measure matching feasibility: Upper bound for the noise probability tolerated by a graph based on the theory



Models considered

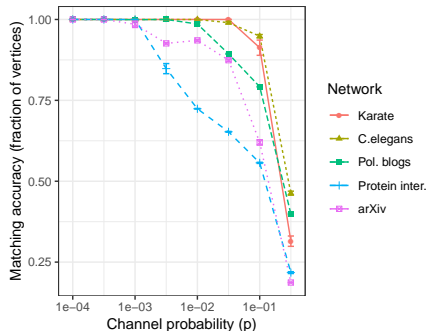
- ▶ $G(n, dn)$ Erdős-Rényi
- ▶ $WS(n, d, \beta)$: Watts-Strogatz
- ▶ $PA(n, \gamma, d)$: preferential attachment

Matchability on real networks



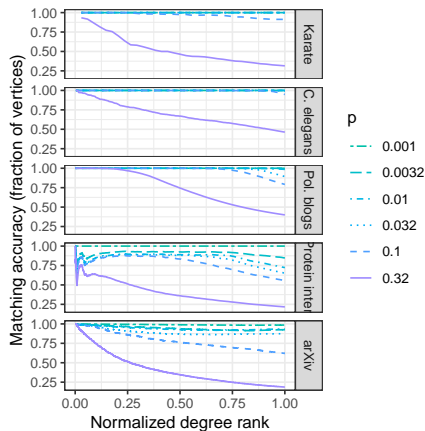
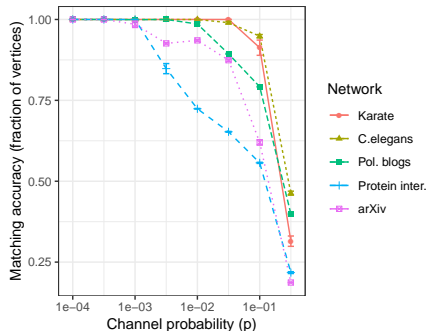
Matchability on real networks

- Sample B from corrupting channel, and approximately compute the MLE
- Evaluate percentage of correctly matched vertices (left) and large degree vertices that are correctly matched (right).



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Conclusion

- Graph matching on errorfully observed networks is *statistically* feasible when:
 - ▶ The topology of the network is sufficiently irregular
 - ▶ The noise is small enough
- What are the computational limits?
- Matching on other relational data structures? multilayer networks, vertex or edge attributes, bipartite graphs, etc.
 - ▶ *Graph matching between bipartite and unipartite networks:* [arXiv:2002.01648](#).

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Thank you!

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