A Polynomial-time Iterative Algorithm for Random Graph Matching with Non-vanishing Correlation

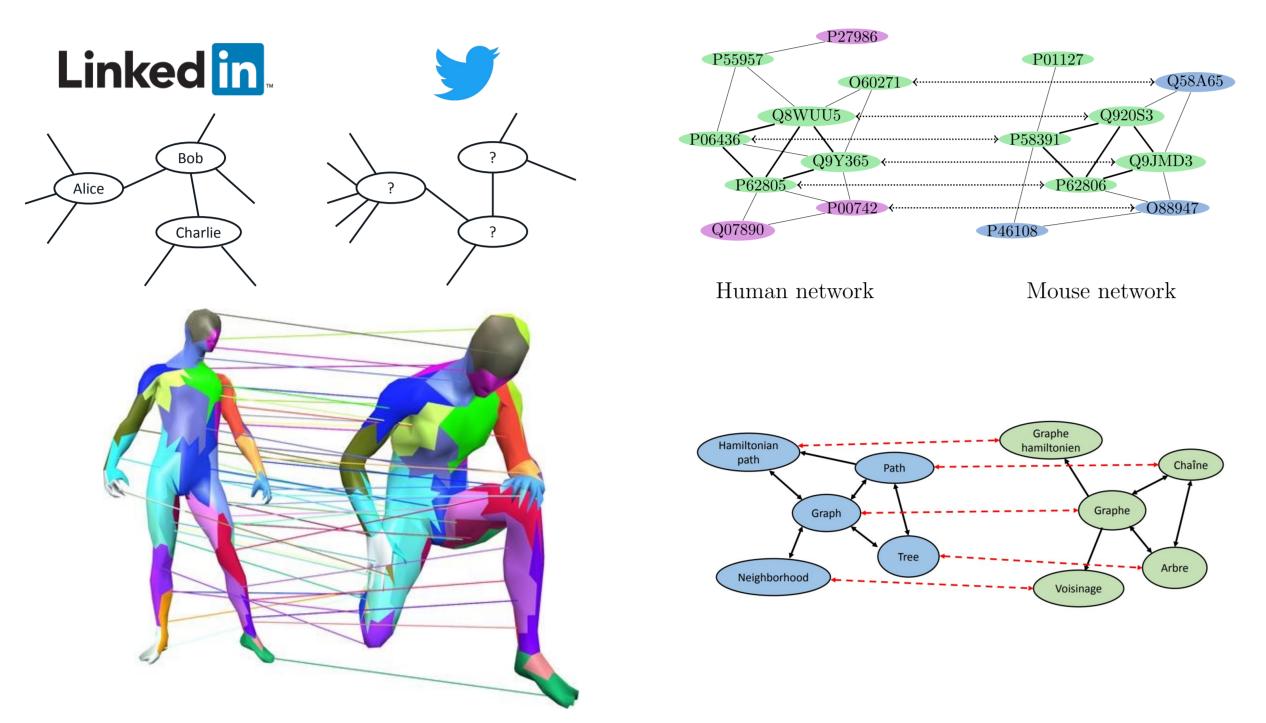
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1. Motivation

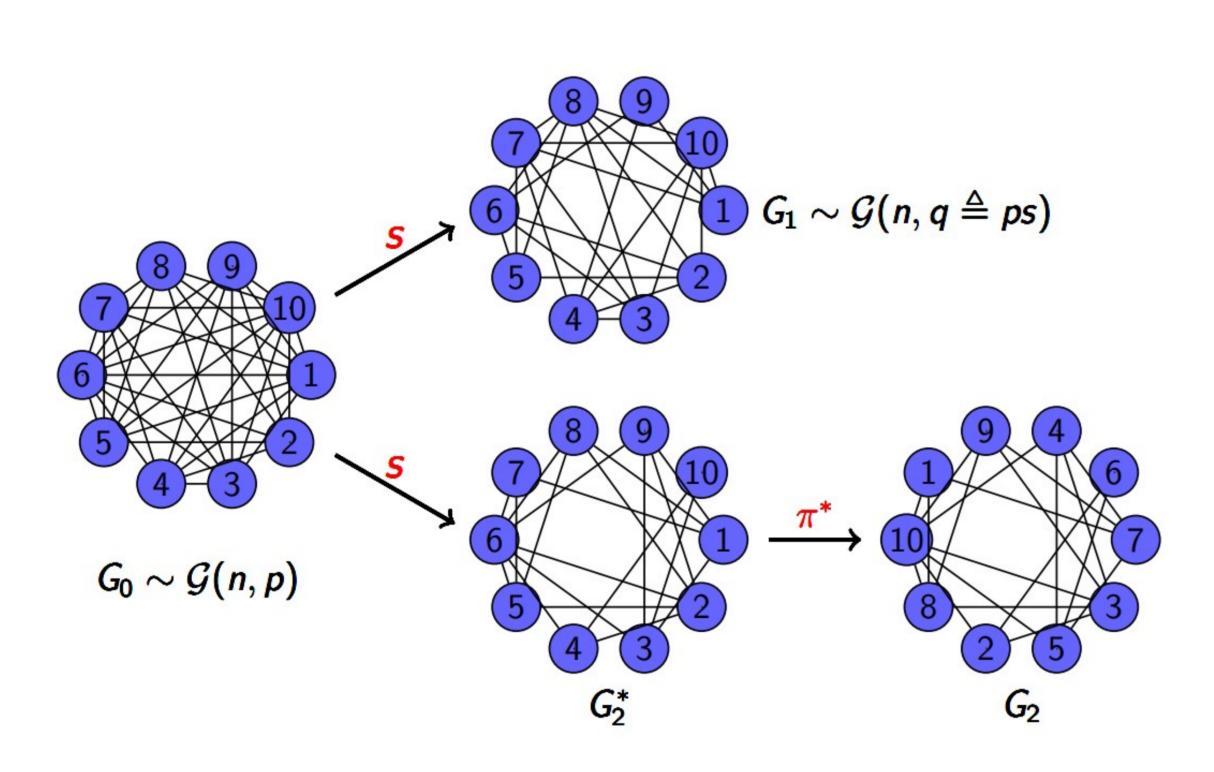
- Consider a pair of graphs with (positive) correlated structure through certain vertex correspondence.
- In most scenarios, the labels of vertices are lost and the only observable is the **topology structure** of graphs.
- People are concerned about recovering the hidden correspondence based on such sheer information, as well as the closely related correlation detection problem.
- Questions of this type are important combinatorial optimization problems which have applications in various fields:
 - social network de-anonymization; protein-protein interaction; computer vision;
 - machine translation.



• In our works [DL22+] and [DL23+], we consider the algorithmic perspective for this problem in *correlated Erdős-Rényi graphs model*.

2. Mathematical settings

- Denote $\mathcal{G}(n,p)$ for the law of Erdős-Rényi graph on n vertices with edge density p (i.e. each edge is kept independently with probability p).
- Consider the following idealistic model:
- fix $n \in \mathbb{N}$, $p, s \in (0, 1)$;
- Sample a "mother graph" $G_0 \sim \mathcal{G}(n, p)$;
- Independently subsample $G_1, G_2^* \subset G_0$ with subsampling probability s;
- Relabel G_2^* by a uniform permutation π^* to obtain G_2 .



- \mathbb{P} : the law of (G_1, G_2) generated as above; \mathbb{Q} : the law of two independent $\mathcal{G}(n, ps)$ graphs.
- The fundamental problems for this model:
- **Detection**: given (G_1, G_2) , determine whether it is sampled from \mathbb{P} or \mathbb{Q} . (e.g. applications in distinguishing objects by computer.)
- **Recovery**: given $(G_1, G_2) \sim \mathbb{P}$, recover π^* as good as possible. (e.g., applications in de-anonymizing social networks.)
- Believed to exhibit **information-computation gap**, a major challenge in many random combinatorial optimization problems.
- By the collective effort of the community (for example, see [WXY21], [DD22a] and [DD22b]), now we have a fairly complete understanding of the information thresholds for the problem of correlation detection and matching recovery. In contrast, the understanding of the computational aspect is far from being complete.
- Intuitively, the matching π^* that maximizes the common edge between two graphs (i.e. the MLE) should be the most effective estimator for recovering the latent matching π .
- This intuition helps us to reduce the problem into the **network alignment** problem of correlated random graphs.
- Unfortunately, the classical graph alignment problem is a **NP-hard** optimization problem, so we must seek help from randomness.
- Previously, arguably the best result is the recent work [MWXY23] that obtained a polynomial time algorithm that succeeds as long as the correlation is above the square root of the Otter's constant.

3. Previous results

- The existing algorithms are essentially of two types:
- the optimization-based method that relies on "convex relaxation and rounding": original optimization problem is hard to solve, but feasible if we enlarge the space of potential solutions ([FMWX22a, FMWX22b]);
- the signature-based method that relies on "computing and comparing signatures": for each vertex, compute a "signature" and match pairs of vertices with similar signatures. ([BCL+19, DMWX21]).
- In a later breakthrough [MRT23], the authors found the first polynomial time algorithm (based on some sophisticated partition tree) that succeeds for exact matching with correlation a constant close to 1;
- In a recent breakthrough [MWXY23], the authors substantially improved [MRT23] and obtained a polynomial time algorithm which succeeds as long as the correlation is above $\sqrt{\alpha}$, where $\alpha \approx 0.338$ is the Otter's constant.

3.Our results

Theorem 1 (D.-L.'2022). There is a polynomial time iterative algorithm for matching Gaussian matrices as long as the correlation is non-vanishing.

- Strongly suggest the barrier at Otter's constant in [MWXY23] is not real in **dense region**;
- New feature: signal is stored in a vector where each coordinate is a pair of sets, and signal per coordinate decreases with iteration but compensated by increase in dimension;
- Might shed light on many other matching problems too.

Sketch of our iterative algorithm:

- (1) Obtain a seed $u_i, \pi(u_i) : 1 \le i \le K$ via brutal-force searching, where K is a large constant;
- (2) At each time t, construct pairs of sets $(\Gamma_i^{(t)}, \Pi_i^{(t)}), 1 \le i \le K_t$ according to the edge weights of each vertex v to the seeds or $(\Gamma_i^{(t-1)}, \Pi_i^{(t-1)}), 1 \le i \le K_{t-1}$.
- (3) Main observation: although the signal in each pair $(\Gamma_k^{(t)}, \Pi_k^{(t)})$ will decrease, we may take many linear combinations of those edge weights to increase the number of paired sets, so the total signal is increasing in t.

Theorem 2 (D.-L.'2023). There is a polynomial time iterative algorithm for matching dense Erdős-Rényi graphs (i.e., when the edge-density q of Erdős-Rényi graphs satisfies $q \ge n^{-1+\alpha+o(1)}$ for some constant $\alpha > 0$) as long as the correlation is non-vanishing.

- Demonstrates the robustness of the iterative matching algorithm we proposed in [DL22+], this type of "algorithmic universality" is closely related to the universality phenomenon in random matrix theory.
- Expected to be sharp in the following sense: as the correlation tends to 0, no polynomial time algorithm with a fixed power would be able to match two random graphs.

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