

Fast Computation of the Barycenter Graph in the Spectral Domain

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Objective and Contributions

- ❖ Given a set of graph-valued data, how do we **estimate a mean graph** which inherits structural properties?
 - **Essential location parameter** of many ML algorithms working on networks
 - Challenge of capturing **topological information on various scales**
- ❖ Main contributions of our algorithm
 - Mean graph inherits large-scale topological information (community structure)
 - Independent of graph size → can compare **graphs of different sizes**
 - **Computationally efficient** (parametric approach)
 - Supported by **theoretical results**

Code implementation at https://github.com/ocourtney/frechet_mean.git

Algorithm Description

The Sample Fréchet Mean

Notation:

- n : graph size
 - A : adjacency matrix (unweighted, symmetric, $n \times n$)
 - λ : vector of ordered eigenvalues, $\lambda_1 \leq \lambda_2 \leq \dots \leq \lambda_n$
- ❖ We replace the algebraic notion of a mean with a minimization problem
- ❖ Finds the most central object, or ‘center of mass’

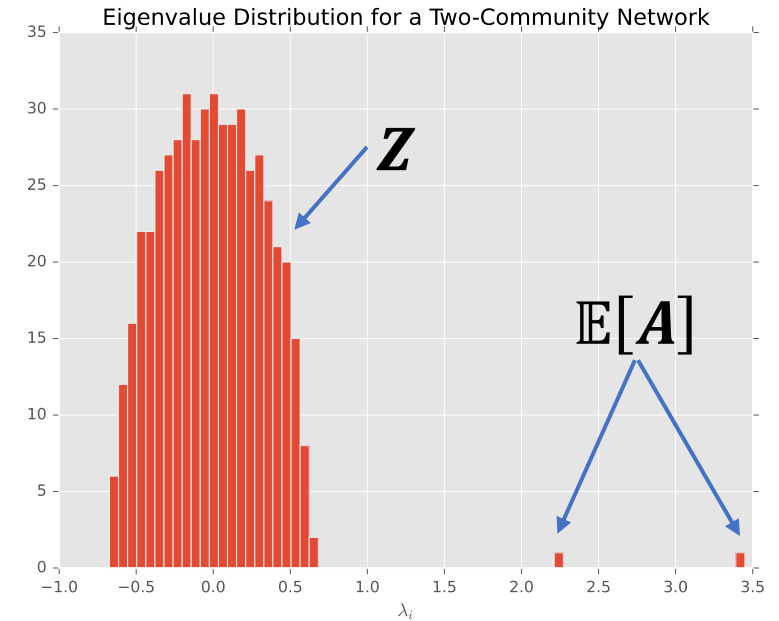
Given a sample of N graphs $\{G^{(1)}, G^{(2)}, \dots, G^{(N)}\}$ and distance d , the **sample Fréchet mean** is the solution to

$$\hat{\mathbb{E}}_N[G] = \arg \min_{G \in \mathcal{G}} \frac{1}{N} \sum_{k=1}^N d^2(G, G^{(k)}),$$

where \mathcal{G} is an unweighted, undirected graph.

Spectral Pseudo-Distance

- Common pairwise distances: Edit, Hamming, Frobenius Norm
 - Good for fine-scale features, but poorly recover large-scale features (i.e. community structure)
- The spectrum contains multi-scale information, so we use this distance



$$\mathbf{A} = \mathbb{E}[\mathbf{A}] + \mathbf{Z},$$

Given graphs G and G' with respective spectra $\lambda(\mathbf{A})$ and $\lambda'(\mathbf{A})$, the adjacency spectral distance between the two graphs is defined as

$$d(G, G') = \sqrt{\sum_{j=1}^n (\lambda_j - \lambda'_j)^2}$$

Problem Statement

- Use spectral distance in Fréchet sample mean definition
- Assume that the mean graph has a Stochastic Block Model (SBM(n, \mathbf{P})) structure
 - Random generative model to construct graphs with community structure
 - Has universal approximating properties
 - k communities C_1, \dots, C_k , $a_{ij} \sim \text{Bernoulli}(p_{ij})$
 - Edge probabilities stored in \mathbf{P}
- We prove that
 - $\lambda_i(\mathbb{E}[\mathbf{A}])$ depend linearly on the density probabilities
 - The error in $\lambda_i(\mathbf{A}) - \lambda_i(\mathbb{E}[\mathbf{A}])$ is bounded
- So we can effectively optimize this difference as a function of \mathbf{P}

$$\mathbb{E}[\mathbf{A}] = \begin{bmatrix} p_1 & \cdots & p_1 & q & \cdots & q \\ \vdots & & \vdots & \vdots & & \vdots \\ p_1 & \cdots & p_1 & q & \cdots & q \\ q & \cdots & q & p_2 & \cdots & p_2 \\ \vdots & & \vdots & \vdots & & \vdots \\ q & \cdots & q & p_2 & \cdots & p_2 \end{bmatrix}$$

$$\hat{G} = \arg \min_{\mathbf{P} \in [0,1]^k} \frac{1}{N} \sum_{i=1}^N \sum_{j=1}^k (\lambda_j(\mathbb{E}[\mathbf{A}]_{\hat{\mathbf{P}}}) - \lambda_j(\mathbf{A}^i))^2$$

The Sample Fréchet Mean Algorithm

- Input: Sample networks $\mathbf{G}_1, \mathbf{G}_2, \dots, \mathbf{G}_N$ (or simply their dominant eigenvalues)
- Output: Probability matrix from which \hat{G} can be generated
- Objective function: $F(\mathbf{P}) = \frac{1}{N} \sum_{i=1}^N \sum_{j=1}^k \|\lambda_j(\mathbb{E}[\hat{\mathbf{A}}]_{\mathbf{P}}) - \lambda_j(\mathbf{A}^{(i)})\|^2$

Algorithm 1 Approximation of the Sample Fréchet Mean

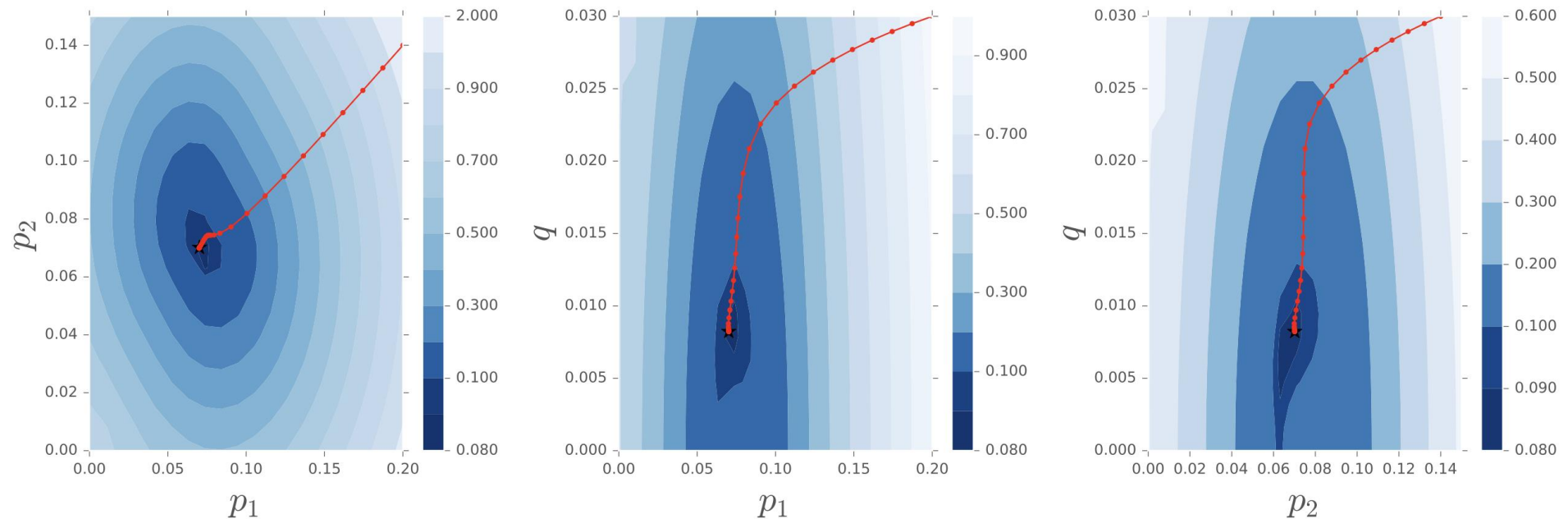
Require: A set of graphs $G = \{G^{(i)}\}_{i=1}^N$.

- 1: Set k and initialize $\mathbf{P} = \mathbf{P}_0$ via the desired method **.
 - 2: For $i = 1, \dots, N$, compute $\boldsymbol{\lambda}^i = [\lambda_1(\mathbf{A}^i), \dots, \lambda_k(\mathbf{A}^i)]$.
 - 3: **while** The relative change in \mathbf{P} is greater than the tolerance **do**
 - 4: Estimate the gradient of $F(\mathbf{P})$ (given by (4)) via finite differences.
 - 5: Update \mathbf{P} via a projected gradient step.
 - 6: **end while**
 - 7: **return** \mathbf{P} .
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Numerical Results

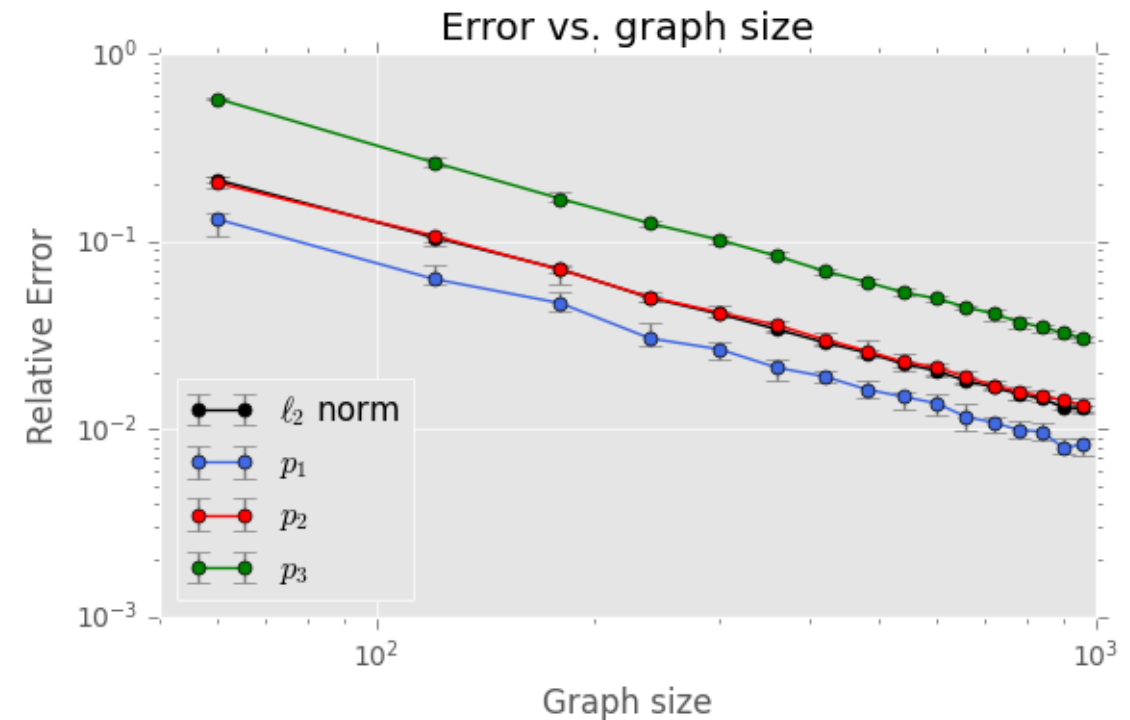
Convergence of gradient descent

- Sample: real network data¹
 - Temporal social contact networks of a primary school
 - $n = 242$
 - We see convergence over the objective function



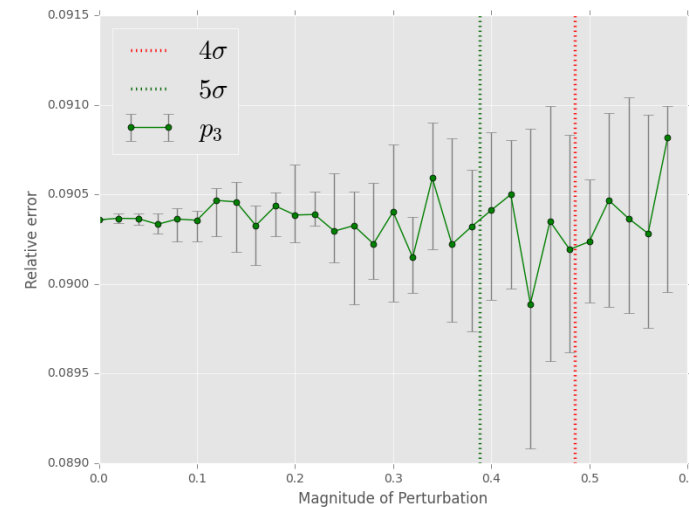
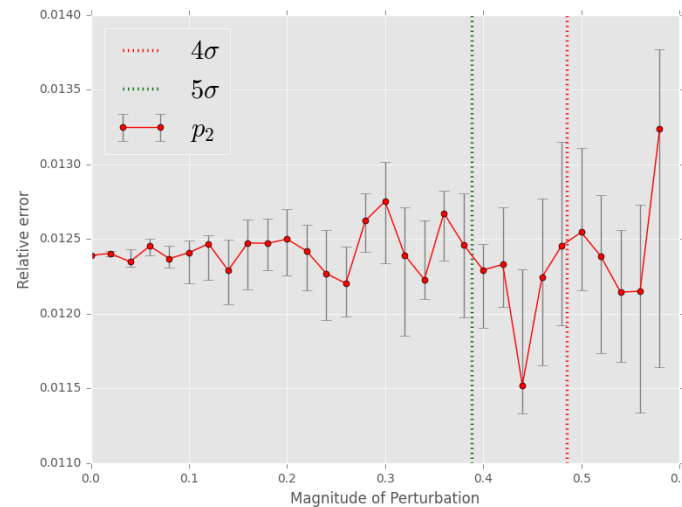
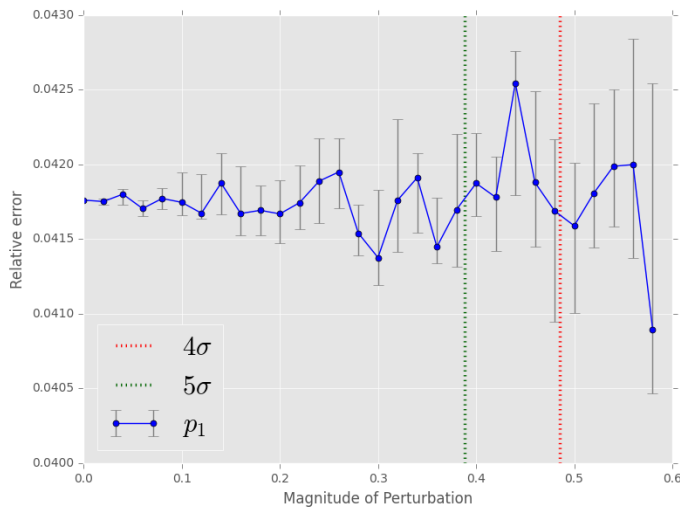
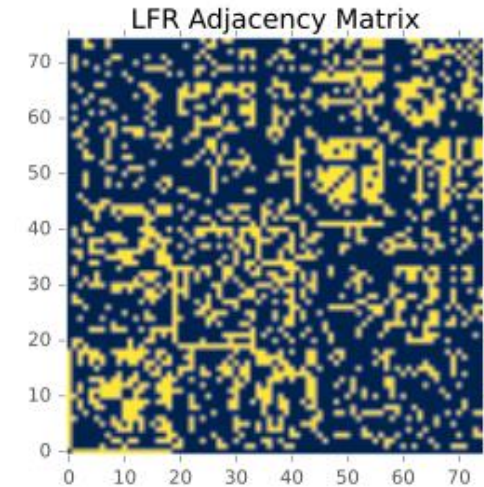
Convergence in the spectral domain

- Sample: SBM networks
 - $n \in [50, 1000]$, $N = 30$
 - $k = 3$, with distinct sizes that are relative to n
 - \mathbf{P} = constant, but distinct p densities
 - Median of 20 trials is plotted
- Spectral error: $\|\hat{\lambda} - \lambda\|_2$
- ❖ We see the algorithm converges to the true mean network as the graph size (n) grows.



Real Network

- True mean: LFR-Benchmark² network
 - $n = 75$, unweighted, undirected
 - $k = 3$, with distinct sizes, non-overlapping
 - Median of 20 trials is plotted
- Recovery of the true mean network given a noisy sample
- Density error: $\|\hat{\mathbf{P}} - \mathbf{P}\|_2$
- Algorithm will on expectation return true mean graph, acts as an unbiased estimator



Conclusion

- ❖ Given a sample of networks, we can **approximate the mean** using a parameterized approach in the spectral domain
- ❖ Using the eigenvalues to measure proximity between graphs allows for the **recovery of community structure**
- ❖ **Limitation:** algorithm relies on our ability to approximate the mean graph with a SBM
- ❖ **Future work:**
 - ❖ Modular analysis of errors from SBM approximation and from algorithm design
 - ❖ Experiments on graphs with no community structure

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