

Introduction & Motivation

Graphical models represent dependencies (e.g., genetics, neuroscience). Inference often uses correlation (\mathbf{C}) or precision matrices.

Challenge: Generate *theoretical* \mathbf{C} with specific sparsity (graph \mathcal{G}) for benchmarking. Existing methods often yield mean-zero correlations.

Our Goal: Flexible method for $\mathbf{C} \in \mathcal{C}(\mathcal{G})$ with controlled mean, mimicking real data (e.g., brain connectivity [1]).

Key Definitions:

- \mathbf{C} : symmetric, PSD, $c_{ii} = 1$, $|c_{ij}| \leq 1$.
- $\mathcal{G} = (\mathcal{V}, \mathcal{E})$: sparsity means $c_{ij} = 0$ if $(i, j) \notin \mathcal{E}$.
- Graph density (d): $2|\mathcal{E}|/(p(p-1))$.

Related Work

Cholesky/Polar Parametrizations [2, 6]:

- $\mathbf{C} = \mathbf{U}\mathbf{U}^\top$.
- **Pro:** Uniform sampling in $\mathcal{C}(\mathcal{G})$.
- **Con:** Only for chordal graphs.

Diagonal Dominance [3]:

- Diagonally dominant $\tilde{\mathbf{C}}$, then normalize.
- **Pro:** Works for any \mathcal{G} .
- **Con:** Very small correlations.

Partial Orthogonalization [2]:

- Orthogonalize rows based on \mathcal{E} .
- **Pro:** Non-chordal graphs.
- **Con:** Sensitive to initialization, mean-zero matrices.

Proposed Approach: Convex Optimization

Generate $\mathbf{C} \in \mathcal{C}(\mathcal{G})$ by solving:

$$\begin{aligned} & \underset{\mathbf{C}}{\text{minimize}} && \frac{1}{2} \|\mathbf{C} - \bar{\mathbf{C}}\|_F^2 \\ & \text{subject to} && \mathbf{C} \succeq 0 \quad (\text{PSD}) \\ & && \text{diag}(\mathbf{C}) = \mathbf{1} \\ & && |c_{ij}| \leq 1, \quad \forall (i, j) \\ & && c_{ij} = 0, \quad \forall (i, j) \notin \mathcal{E} \\ & && \frac{1}{2|\mathcal{E}|} \sum_{(i,j) \in \mathcal{E}} c_{ij} \geq b \quad (*) \end{aligned}$$

Novelty: Mean constraint

$\bar{\mathbf{C}}$ = target matrix, b = mean of edge correlations.

Key Aspects:

- Solved via convex optimization (CVXPY [4]).
- Objective: feasible \mathbf{C} close to $\bar{\mathbf{C}}$.
- Constraint (*) controls mean via b .
- Unique solution if feasible ($b \leq 0$ guarantees feasibility).
- **Post-processing:** If needed, $\tilde{\mathbf{C}}_\epsilon = \bar{\mathbf{C}} + \epsilon \mathbf{I}$, $\mathbf{C} = \tilde{\mathbf{C}}_\epsilon / (1 + \epsilon)$.

Results: Influence of Graph Structure

Tested: ER, BA, WS, SBM, Chordal.

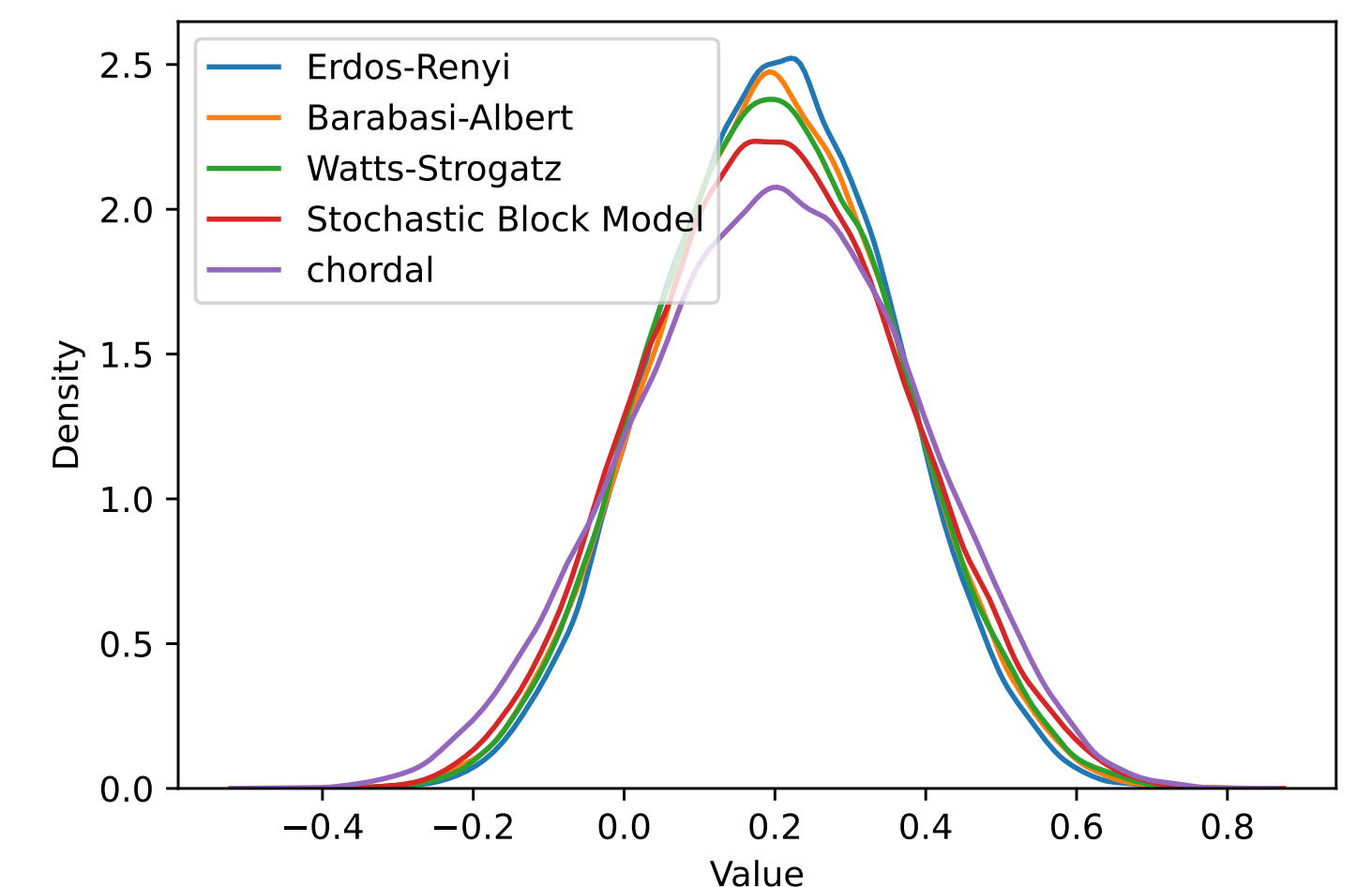


Figure 3: Density plots ($p = 51$, $d = 0.5$, $b = 0.2$). Minimal impact of graph structure at this scale.

Observations:

- For $p = 51$, graph type has limited impact on correlation distribution given fixed d and b .
- Influence may increase for larger p .
- More computationally intensive but better control over statistical properties.

Results: Comparison & Mean Control

Simulations: $p = 51$, 50 runs. Target $\bar{\mathbf{C}} \sim U(-1, 1)$ unless stated.

Comparison (ER, $d = 0.5$, $b = -1$):

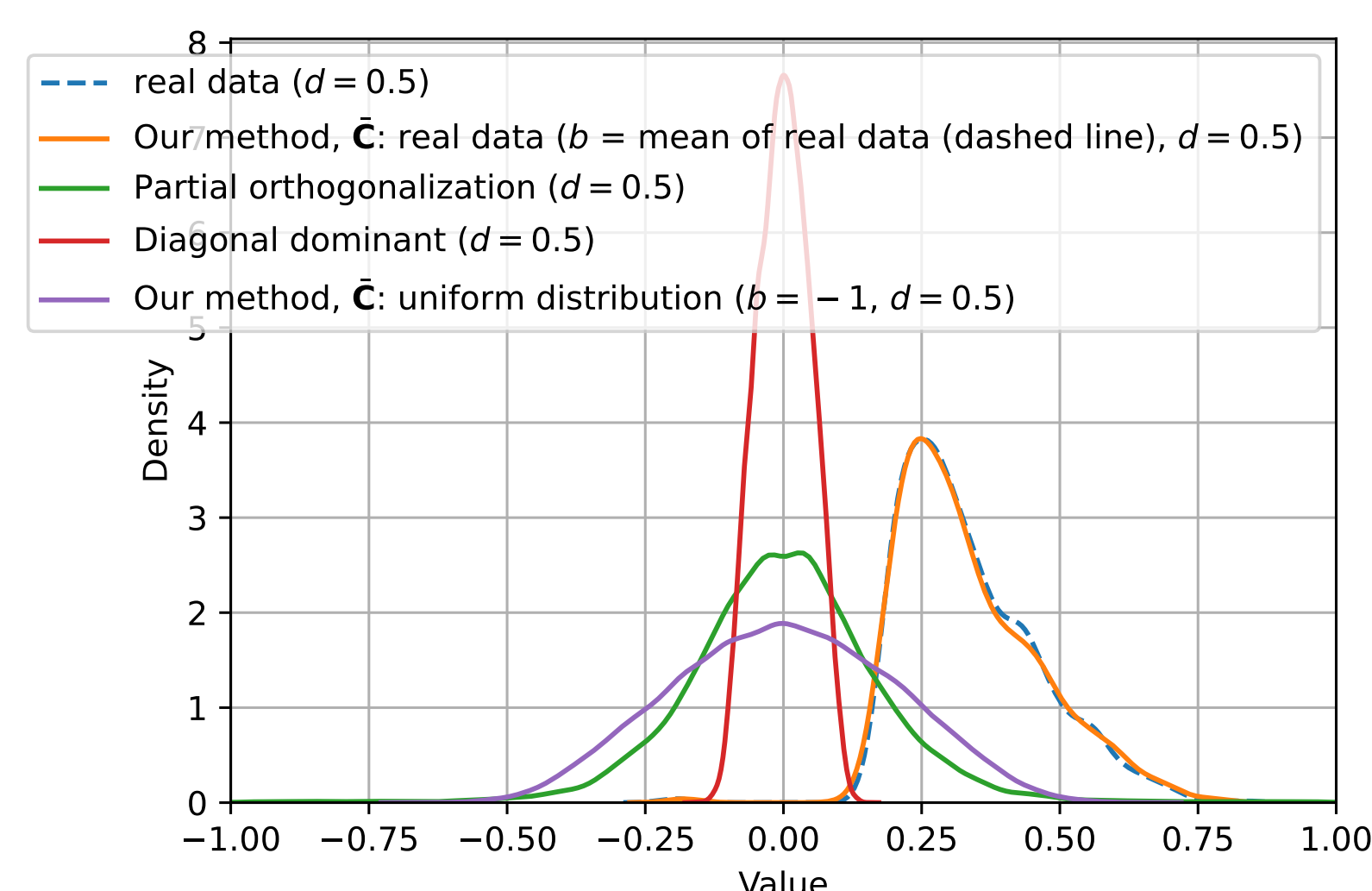


Figure 1: Density plots. Our method (purple) vs. others (red, green). Approximates real fMRI (blue) when tuned (orange).

Effect of Mean Constraint (b):

- Fig. 1 (orange) fits fMRI [5] using empirical b and $\bar{\mathbf{C}}$.
- Feasibility depends on b and graph density d .

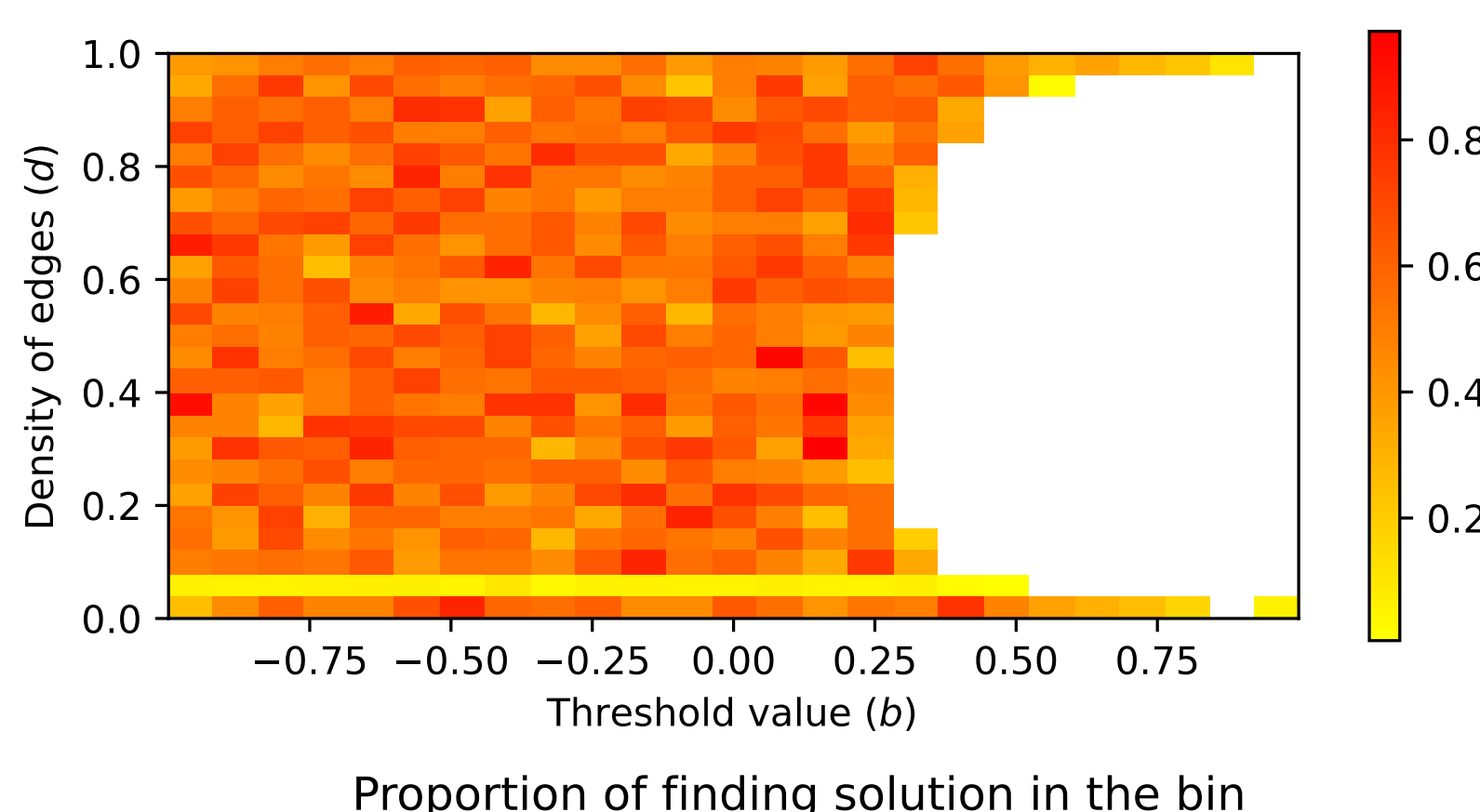


Figure 2: Feasibility map for ER graphs ($p = 51$). White = infeasible (d, b).

Conclusion

- Convex optimization framework to generate \mathbf{C} with graph structure \mathcal{G} .
- **Key advantage:** Explicit control over mean correlation (b) for realistic data generation.
- **Improvement:** Supports any graph, avoids near-zero correlations, introduces mean control.
- Application: benchmarking inference methods on controlled graph structures.
- Future: scaling to large p , deeper study of graph structure role.

Code: [7]

Acknowledgements & References

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Selected References:

- [1] S. Achard et al. (2022). NeuroImage.
- [2] I. Córdoba et al. (2020). J Comput Graph Stat.
- [3] I. Córdoba et al. (2020). TEST.
- [4] S. Diamond & S. Boyd (2016). JMLR.
- [5] G. Becq et al. (2020). NeuroImage, 219, 116945.
- [6] M. Pourahmadi (2015). Stat & Prob Let.
- [7] A. Fakhar et al. Code: https://github.com/alifakhar7/corr_mat_gen