Motivation

- Gaussian Boson Sampling effectively samples submatrices of a given matrix according to their Hafnians
- Calculating the Hafnian is #P-hard, but there are probabilistic estimators for non-negative matrices
- Depending on statistical properties, these estimators could "dequantize" some proposed applications of GBS

Gaussian Boson Sampling

A perfect Gaussian boson sampler with M modes is specified by m squeeze parameters $r_i \in \mathbb{R}$ and the interferometer unitary $U \in \mathcal{U}(M)$. The **kernel matrix** \mathcal{A} is then defined as $A \oplus A^*$, where

 $A = U \operatorname{diag}(\tanh r_1 \ldots \tanh r_M) U^T. \quad (1)$ A measurement outcome $\mathbf{n} = (n_1, ..., n_M)$, where n_i is the number of photons measured in mode j, corresponds to a submatrix $A_{\mathbf{n}}$ of \mathcal{A} where *j*-th and

(j+m)-th row and column are taken n_j times. The probability of observing such an outcome is proportional to

$$P(\mathbf{n}) \propto \frac{1}{n_1! \dots n_M!} \operatorname{Haf}(\mathcal{A}_{\mathbf{n}}).$$
 (2)

Here the **Hafnian** function treats a matrix as an adjacency matrix of a graph and counts its **perfect** matchings:

Haf
$$(\square) = \square + \square + \square$$

For edge-weighted graphs, a perfect matching contributes a product of its edge weights.

Denser graphs tend to have more perfect matchings, so GBS can be used as a heuristic to find cliques in a graph.

Randomized estimators of the Hafnian of a non-negative matrix

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Randomized estimators

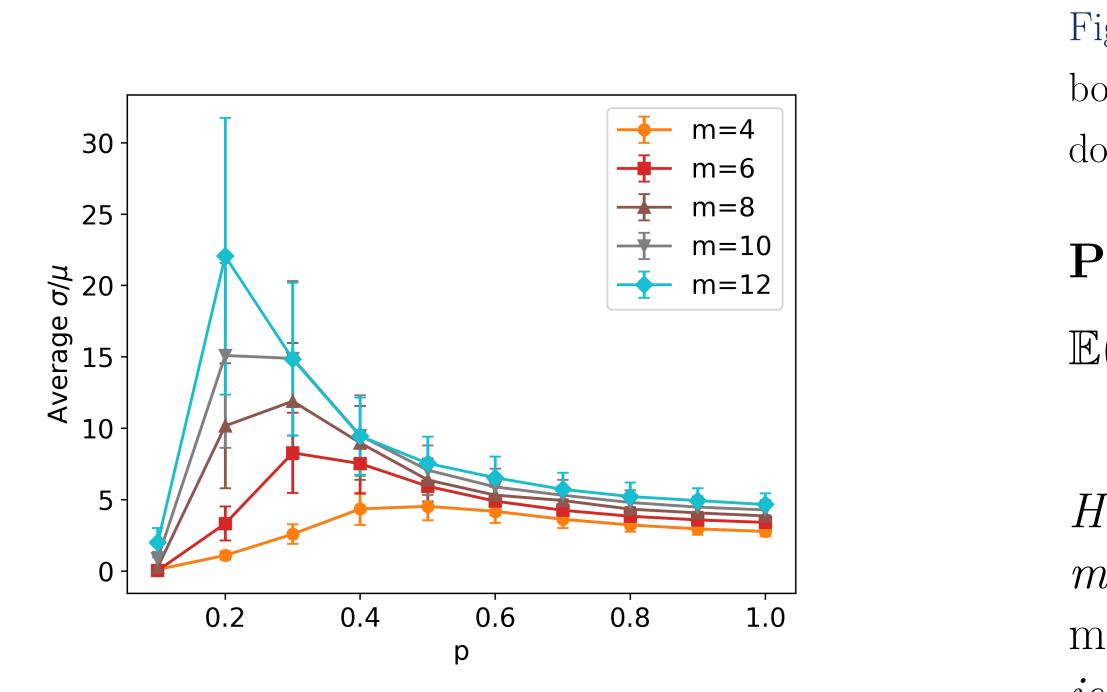
Assuming that $A \in \mathbb{R}^{2m \times 2m}$ is a non-negative matrix (i.e. all $a_{ij} \ge 0$), we can construct an estimator of the Hafnian. Let $W \in \mathbb{R}^{2m \times 2m}$ be a random skew-symmetric matrix such that $\mathbb{E}w_{ij} = 0$, $\mathbb{E}w_{ij}^2 = 1$, and all above-diagonal entries are i.i.d. Define G to be a matrix with $g_{ij} = w_{ij} \sqrt{a_{ij}}$. Then $\mathbb{E} \det G = \operatorname{Haf} A.$

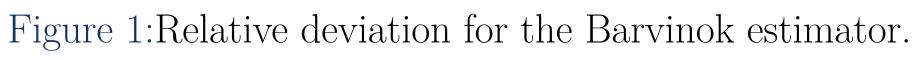
Here we look at two estimators:

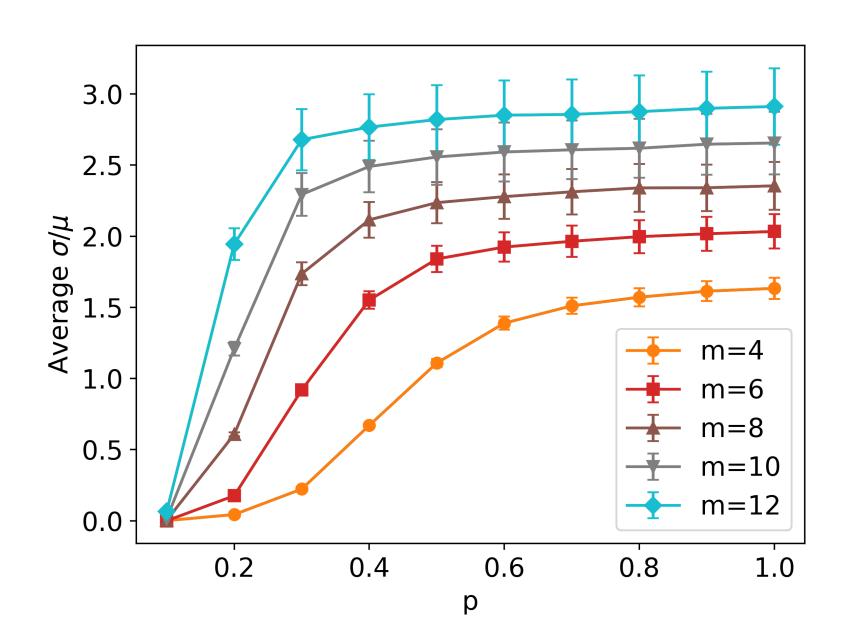
- Barvinok estimator: $w_{ij} \sim \mathcal{N}(0, 1)$.
- Godsil-Gutman estimator: $w_{ij} \in \{-1, 1\},\$

sampled with equal probability.

Numerical results for Erdős-Rényi graphs







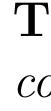
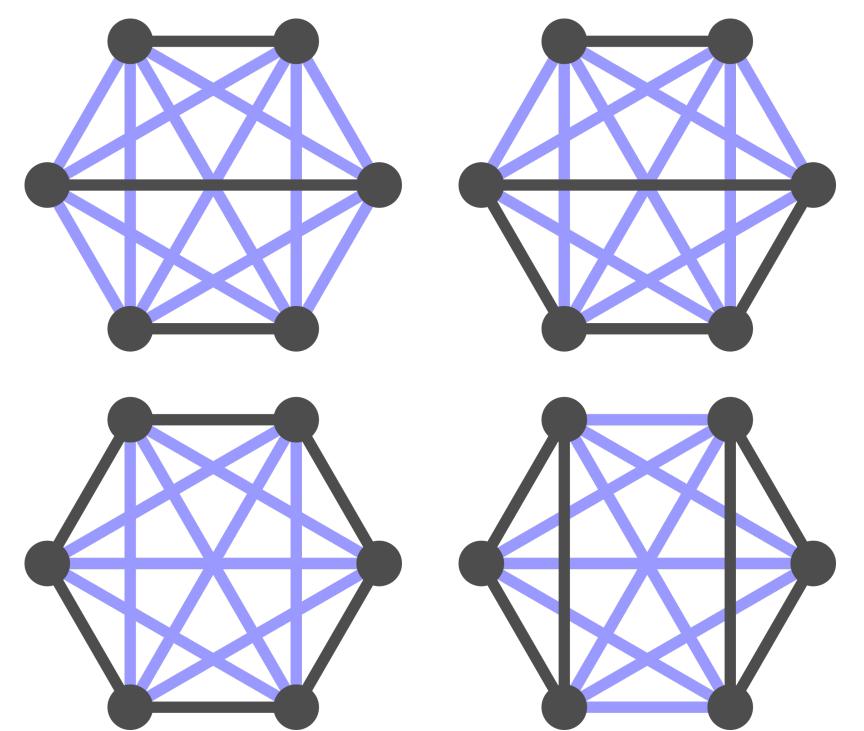


Figure 2:Relative deviation for the Godsil-Gutman estimator.

Analytical results

The variance can be expressed in terms of **perfect 2-matchings**, i.e. spanning subgraphs such that all connected components are either cycles or isolated edges:



Examples like this can be cracked with the FKT algorithm which calculates the Hafnian for planar graphs in polynomial time. However, adding a few more edges to make the graph non-planar will make FKT useless as well.

Figure 3:Examples of perfect 2-matchings. The one on the bottom right contains cycles of odd length; such 2-matchings do not contribute to the variance.

Proposition 1. Let $\mathbb{E}w_{ij}^3 = 0$, $\mathbb{E}w_{ij}^4 = \eta$. Then $\mathbb{E}(\det G)^2 = \sum \eta^{|\mathrm{match}(d)|} 6^{|\mathrm{cycle}(d)|}$ $a_{ij}^2 a_{kl}.$ $\{i,j\} \in \mathrm{match}(d)$ $\{k,l\} \in d \setminus \mathrm{match}(d)$

Here the sum is taken over all perfect 2machings d that contain cycles of even length; match(d) is the set of isolated edges in d; cycle(d)is the set of even-length cycles in d.

For Barvinok estimator $\eta = 3$, while for Godsil-Gutman estimator $\eta = 1$.

$$(\operatorname{Haf} A)^{2} = \sum_{d} 2^{|\operatorname{cycle}(d)|} \prod_{\substack{(i,j) \in \operatorname{match}(d) \\ (k,l) \in d \setminus \operatorname{match}(d)}} a_{ij}^{2} a_{kl}. \quad (3)$$

Theorem 1. Let A be the adjacency matrix of a complete graph with 2m vertices. Then

 $\frac{\mathbb{E}\det G^2}{(\operatorname{Haf} A)^2} = \sqrt{\pi}m e^{\frac{\eta-3}{2}} + O(1), \ m \to \infty.$ (4)

racy:

- graphs.





Special cases

A graph like this will require an exponential number of samples to get the Hafnian with a constant accu-

Conclusions

• We investigated the statistical properties of the Hafnian estimators and found that the Hafnian of a random graph is typically easy to estimate. • We derived the expression for the estimator variance in terms of perfect 2-matchings. The

Godsil-Gutman estimator always has a smaller variance than the Barvinok estimator.

• We prove that both esimators demonstrate a linear scaling of relative variance for for complete

Open question: is there a fully polynomial randomized approximation scheme (FPRAS) for the Hafnian? If yes, GBS experiments based on random

graphs would be classically simulable.

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