

# Sublinear-Time Algorithms for Monomer-Dimer Systems on Bounded Degree Graphs

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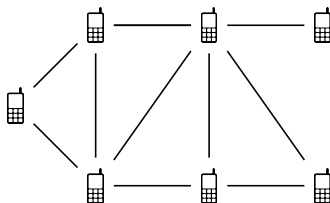
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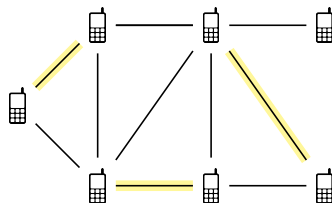
# Background

- Sublinear-time algorithms for graph problems
- Optimization vs. **Counting and Statistics**



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# Monomer-Dimer systems

- $G = (V, E)$  undirected graph with  $|V| = n$
- Maximum degree  $\Delta$
- Monomer  $\leftrightarrow$  one vertex, Dimer  $\leftrightarrow$  two adjacent vertices
- Monomer-Dimer arrangement  $\leftrightarrow$  matching

# Definition

- $\mathbb{M}$ : set of all matchings of  $G$
- **Partition function:**  $Z(G, \lambda) = \sum_{M \in \mathbb{M}} \lambda^{|M|}$
- **Gibbs distribution:**

$$\pi_{G, \lambda}(M) = \frac{\lambda^{|M|}}{Z(G, \lambda)}, \quad \forall M \in \mathbb{M}$$

- **Marginal probability:**

$$p_{G, \lambda}(v) := \sum_{M \ni v} \pi_{G, \lambda}(M), \quad \forall v \in V$$

- Partition function:
  - $\#P$ -complete (Valiant 1979, Vadhan 2002)
  - Randomized polynomial-time approximation scheme (Sinclair 1993, Jerrum Sinclair 1997)
  - Deterministic polynomial-time approximation scheme (Bayati *et al.* 2007)
- Matching statistics
  - $\#P$ -hard (Sinclair Srivastava 2013)
- Permanent of expander graphs
  - Randomized polynomial-time approximation scheme (Jerrum Sinclair Vigoda 2004)
  - Deterministic polynomial-time approximation algorithm (Gamarnik Katz 2010)

# Our Main Results

Local computations for **marginal probability**

- Approximation algorithm
- Complexity lower bound

Randomized **sublinear-time** approximation algorithms for

- **Partition function**
- **Matching statistics**
- **Permanent of expander graphs**

# Notations

- $\epsilon$ -approximation solution : within additive error  $\epsilon$
- $\epsilon$ -approximation algorithm : outputs an  $\epsilon$ -approximation solution with probability at least  $2/3$
- Oracles:  $\mathcal{D}(v)$  and  $\mathcal{N}(v, i)$
- Our focus: Query complexity



# Marginal probability

$$p_{G,\lambda}(v) = \frac{1}{1 + \lambda \sum_{u \in N(G,v)} p_{G \setminus \{v\}, \lambda}(u)}$$

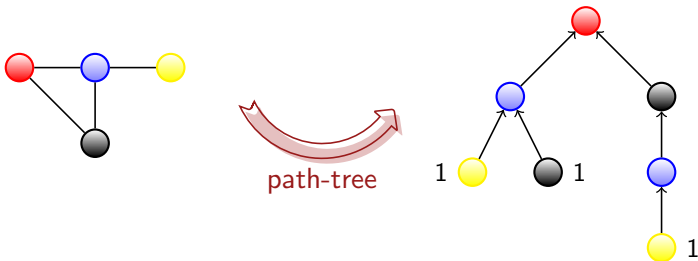
What is the probability that  is matched? ( $\lambda = 1$ )



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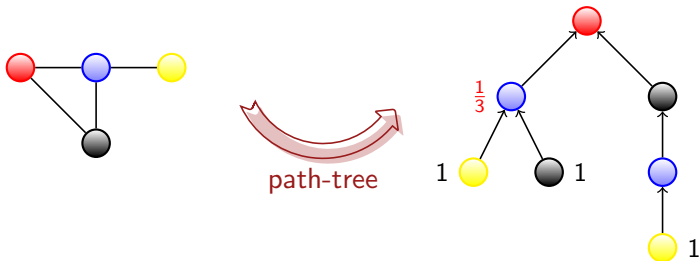
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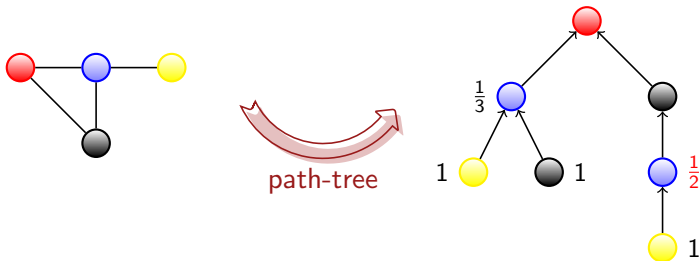
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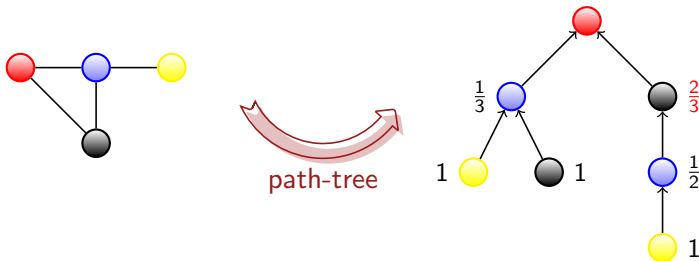
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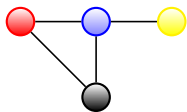
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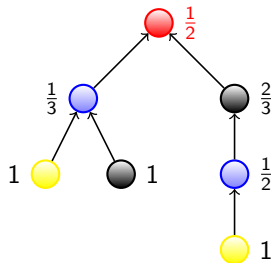
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path-tree



- $T(v)$  : the path-tree rooted at  $v$
- $x_u(v) := \frac{1}{1 + \lambda \sum_{w \succ u} x_w(v)}$ , for every  $u \in T(v)$
- $x_v(v) = p_{G,\lambda}(v)$

# Truncated path-tree

- $T^h(v)$  :  $T(v)$  truncated at depth  $h$
- $x_v^h(v)$  : the solution of the recursions in  $T^h(v)$

Correlation decay property (Bayati *et al.* 2007)

$$|\log x_v^h(v) - \log p_{G,\lambda}(v)| \leq \epsilon, \text{ for any } h \geq h(\epsilon, \Delta) = \tilde{O}\left(\sqrt{\lambda\Delta} \log(1/\epsilon)\right).$$



## Proposition

*We can compute an  $\epsilon$ -approximation of  $p_{G,\lambda}(v)$  using  $O(\Delta^{h(\epsilon,\Delta)})$  queries.*

## Proposition (lower bound)

*Any approximation algorithm requires at least the above query complexity.*

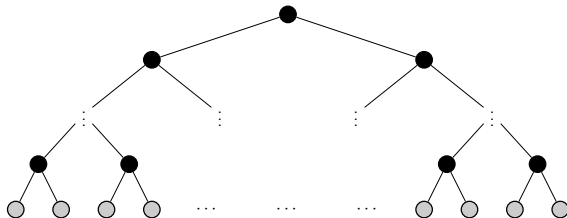
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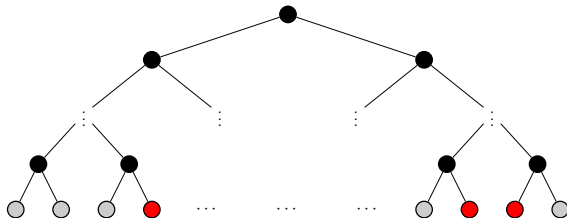
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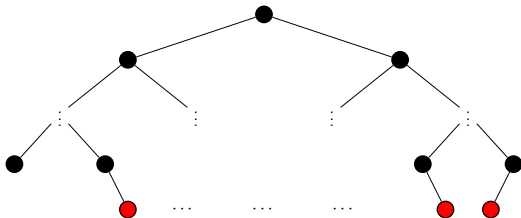
# Idea of the lower bound



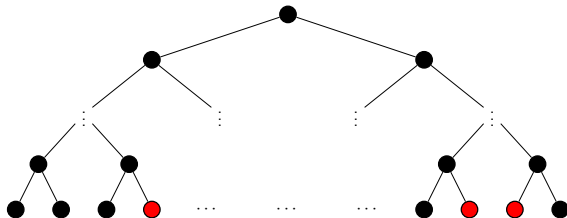
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$$Z(G, \lambda) = \prod_{1 \leq k \leq n} p_{G_k, \lambda}^{-1}(v_k)$$

## Algorithm for $\log Z(G, \lambda)$

- Take  $\Theta(1/\epsilon^2)$  samples uniformly at random from  $[1, \dots, n]$ ;
- Compute an  $\epsilon$ -approximation of  $\log p_{G_k, \lambda}^{-1}(v_k)$  for every sample  $k$ ;
- Return  $n \cdot$  (the average of the estimates).

## Main Theorem

*We have an  $\epsilon n$ -approximation algorithm for the logarithm of the partition function with  $\tilde{O}\left((1/\epsilon)^{\tilde{O}(\sqrt{\Delta})}\right)$  queries. In addition, any  $\epsilon n$ -approximation algorithm needs  $\Omega(1/\epsilon^2)$  queries.*



# Average matching size

$$\begin{aligned} E(G, \lambda) &:= \sum_{M \in \mathbb{M}} |M| \cdot \pi_{G, \lambda}(M) \\ &= n - \frac{1}{2} \sum_{1 \leq k \leq n} p_{G, \lambda}(v_k) \end{aligned}$$

## Theorem

*We have an  $\epsilon n$ -approximation algorithm for the average matching size with  $\tilde{O}\left((1/\epsilon)^{\tilde{O}(\sqrt{\Delta})}\right)$  queries. In addition, any  $\epsilon n$ -approximation algorithm needs  $\Omega(1/\epsilon^2)$  queries.*

# Entropy of a matching

$$\begin{aligned} S(G, \lambda) &:= - \sum_{M \in \mathcal{M}} \pi_{G, \lambda}(M) \log \pi_{G, \lambda}(M) \\ &= \log Z(G, \lambda) + \log \lambda \cdot E(G, \lambda) \end{aligned}$$

## Corollary

*We have an  $\epsilon n$ -approximation algorithm for the entropy of a matching with  $\tilde{O}\left((1/\epsilon)^{\tilde{O}(\sqrt{\Delta})}\right)$  queries. In addition, any  $\epsilon n$ -approximation algorithm needs  $\Omega(1/\epsilon^2)$  queries.*

# Application: Permanent of expander graphs

- **Bi-partite** graph  $G$  with vertices  $X \cup Y$ ,  $|X| = |Y| = n$
- **$\alpha$ -expander** graph:  
 $|N(S)| \geq (1 + \alpha)|S|$ , for  $S \subset X$  or  $S \subset Y$  with  $|S| \leq n/2$
- **PERM** : Permanent of the adjacency matrix of  $G$

Lemma (Gamarnik Katz 2010)

$$1 \leq \frac{Z(G, \lambda)}{\lambda^n \cdot \text{PERM}} \leq e^{O(n\lambda^{-1} \log^{-1}(1+\alpha) \log \Delta)}.$$

Corollary

*We have an  $\epsilon n$ -approximation algorithm for  $\log \text{PERM}$  with query complexity  $\tilde{O}\left((1/\epsilon)^{\tilde{O}(\sqrt{\Delta}/(\epsilon\alpha))}\right)$ .*

- Marginal probability (*correlation decay property*)
- Sublinear-time approximation for monomer-dimer systems
- Extension to the partition function of independent sets
- Experiments on large real-world graphs

Thank you!