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

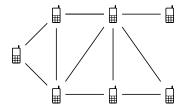
## Marc Lelarge<sup>1,2</sup> and Hang Zhou<sup>1</sup>

<sup>1</sup>École Normale Supérieure de Paris, France

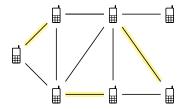
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- Sublinear-time algorithms for graph problems
- Optimization vs. Counting and Statistics



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

## Definition

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

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

• Marginal probability:

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

#### • 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)

Local computations for marginal probability

- Approximation algorithm
- Complexity lower bound

Randomized sublinear-time approximation algorithms for

- Partition function
- Matching statistics
- Permanent of expander graphs

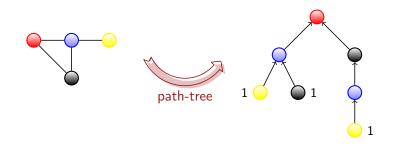
- $\epsilon$ -approximation solution : within additive error  $\epsilon$
- $\epsilon$ -approximation algorithm : outputs an  $\epsilon$ -approximation solution with probability at least 2/3

- Oracles:  $\mathcal{D}(v)$  and  $\mathcal{N}(v, i)$
- Our focus: Query complexity

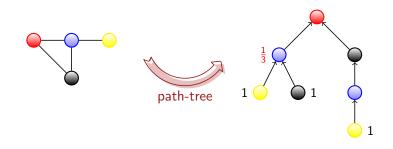
$$p_{G,\lambda}(v) = \frac{1}{1 + \lambda \sum_{u \in N(G,v)} p_{G \setminus \{v\},\lambda}(u)}$$



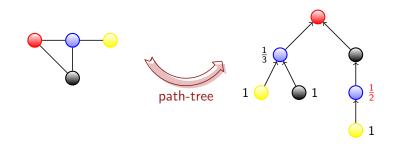
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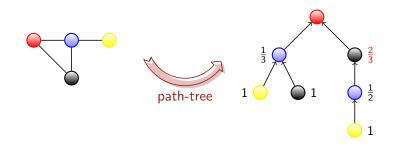
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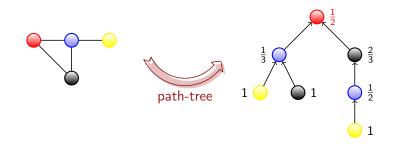
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• 
$$T(v)$$
 : the path-tree rooted at  $v$ 

• 
$$x_u(v) := rac{1}{1 + \lambda \sum_{w \succ u} x_w(v)}, ext{ for every } u \in T(v)$$

•  $x_v(v) = p_{G,\lambda}(v)$ 

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

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

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

#### Proposition

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

#### Proposition (lower bound)

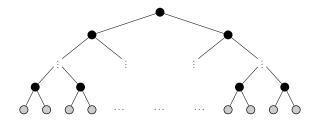
Any approximation algorithm requires at least the above query complexity.

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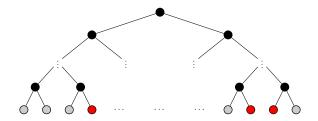
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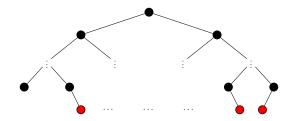
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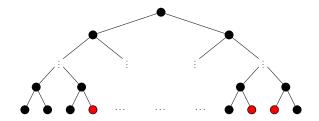
Sublinear-Time Algorithms for Monomer-Dimer Systems



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

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

- Take  $\Theta(1/\epsilon^2)$  samples uniformly at random from  $[1, \ldots, n]$ ;
- 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

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

#### 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.

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

#### 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:

 $|N(S)| \ge (1 + \alpha)|S|$ , for  $S \subset X$  or  $S \subset Y$  with  $|S| \le n/2$ 

• PERM : Permanent of the adjacency matrix of G

#### Lemma (Gamarnik Katz 2010)

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

#### Corollary

We have an  $\epsilon$ n-approximation algorithm for  $\log PERM$  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!