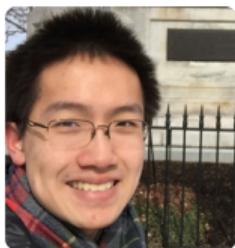


# Limited Correlations, Fractional Log-Concavity, and Fast Mixing

Nima Anari



based on joint works with



Kuikui  
Liu



Shayan  
OveisGharan

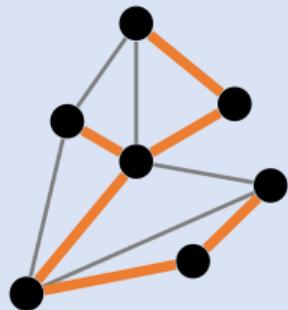


Cynthia  
Vinzant

Efficient sampling/counting/inference when pairwise correlations are ...

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Negative



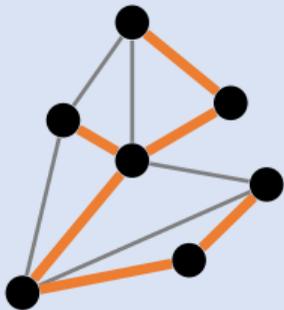
random spanning tree  
 $\mathbb{P}[\text{tree } T] \propto 1$

[Feder-Mihail'92, ...]

$$\mathbb{P}[\text{edge } e \in T \mid \text{edge } f \in T] \leq \mathbb{P}[\text{edge } e \in T] \text{ for } e \neq f$$

Efficient sampling/counting/inference when pairwise correlations are ...

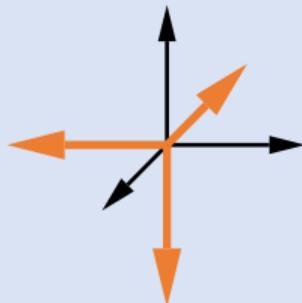
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Spectrally Negative



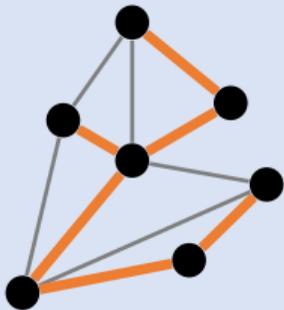
matroid  
 $\mathbb{P}[\text{basis } B] \propto 1$

[previous talk ...]

spectral bound on matrix of correlations  $\equiv$  log-concave polynomial

Efficient sampling/counting/inference when pairwise correlations are ...

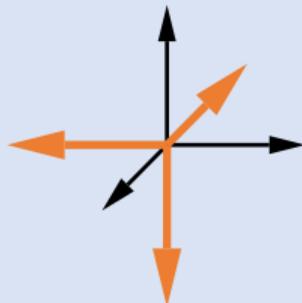
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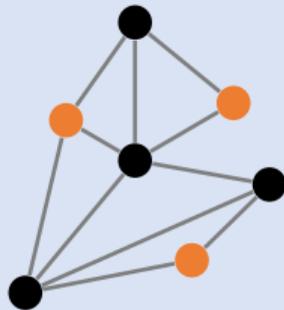
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[previous talk ...]

Decaying



hardcore model  
 $\mathbb{P}[\text{stable } S] \propto \lambda^{|S|}$

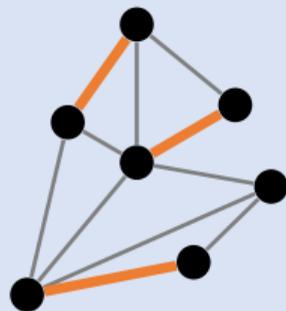
[Weitz'06, ...]

$\mathbb{P}[\text{vertex } u \in S \mid \text{vertex } v \in S] \simeq \mathbb{P}[\text{vertex } u \in S]$  for distant  $u, v$  when  $\lambda < \lambda_c$

What kind of correlations are useful for  
sampling/counting/inference?

# Example with strange correlations

## Monomer-Dimer System

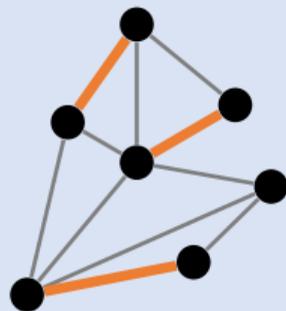


$$\mathbb{P}[\text{matching } M] \propto \prod_{e \in M} \text{weight}(e) \cdot \prod_{v \notin M} \text{weight}(v)$$

# Example with strange correlations

- ▶ Special case is mystery:  
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## Monomer-Dimer System



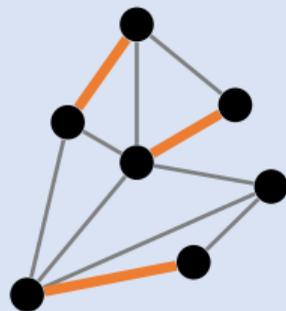
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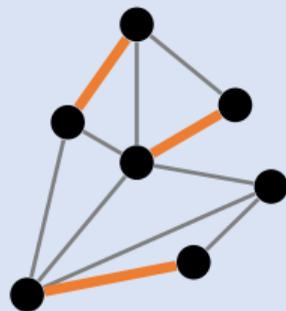
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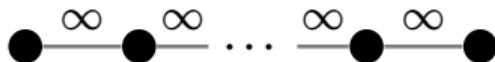
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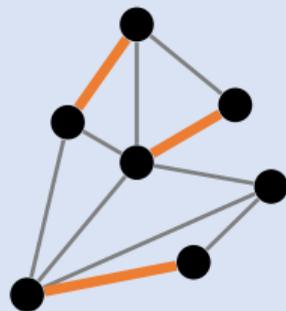
- ▶ Positive correlations:



- ▶ Long-range correlations:



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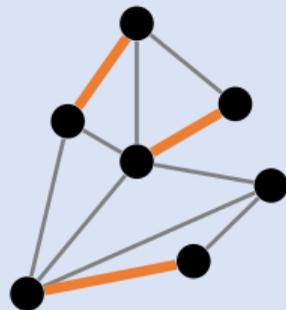


- ▶ Long-range correlations:



- ▶ Correlations are “limited” ...

## Monomer-Dimer System



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- ▶ Efficient sampling/counting/inference:

spectrally limited correlations



fast mixing of simplicial  
complex walks

# Simplicial complex walks

## Canonical form

Distribution defined by

$$\mu : \binom{[n]}{k} \rightarrow \mathbb{R}_{\geq 0}$$

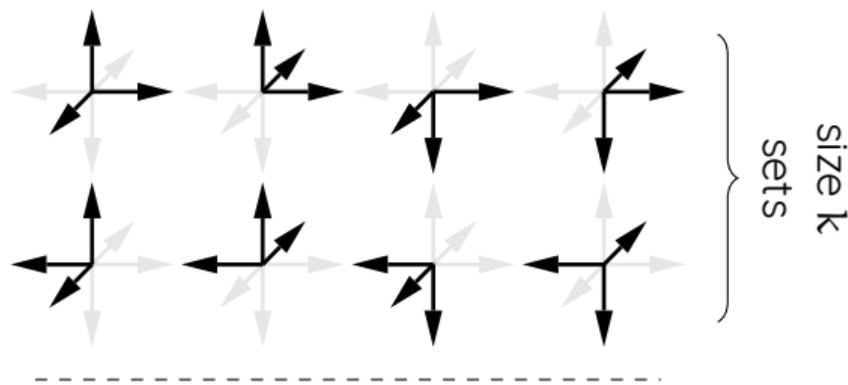
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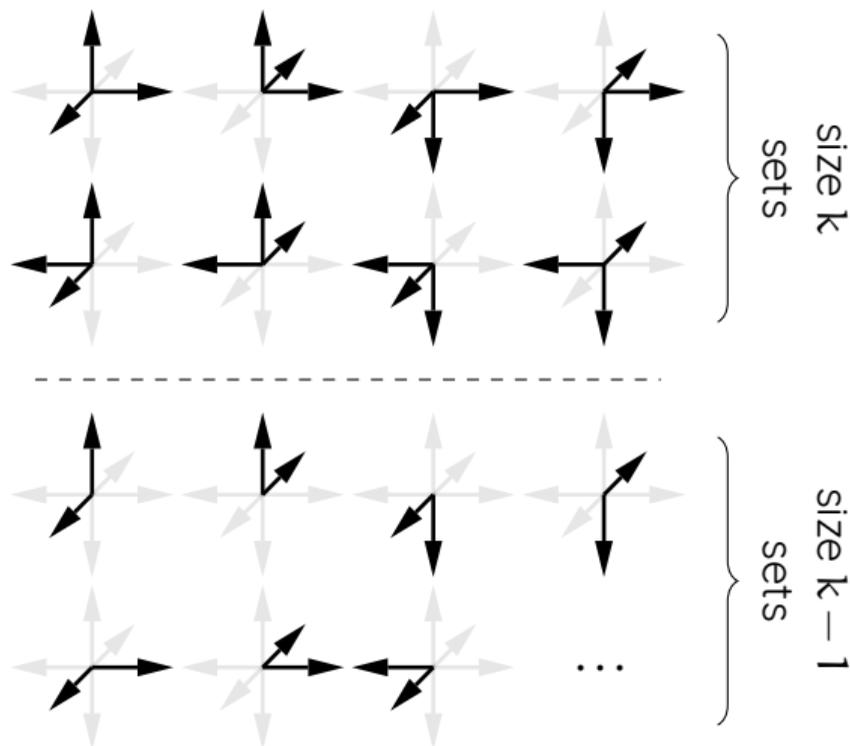
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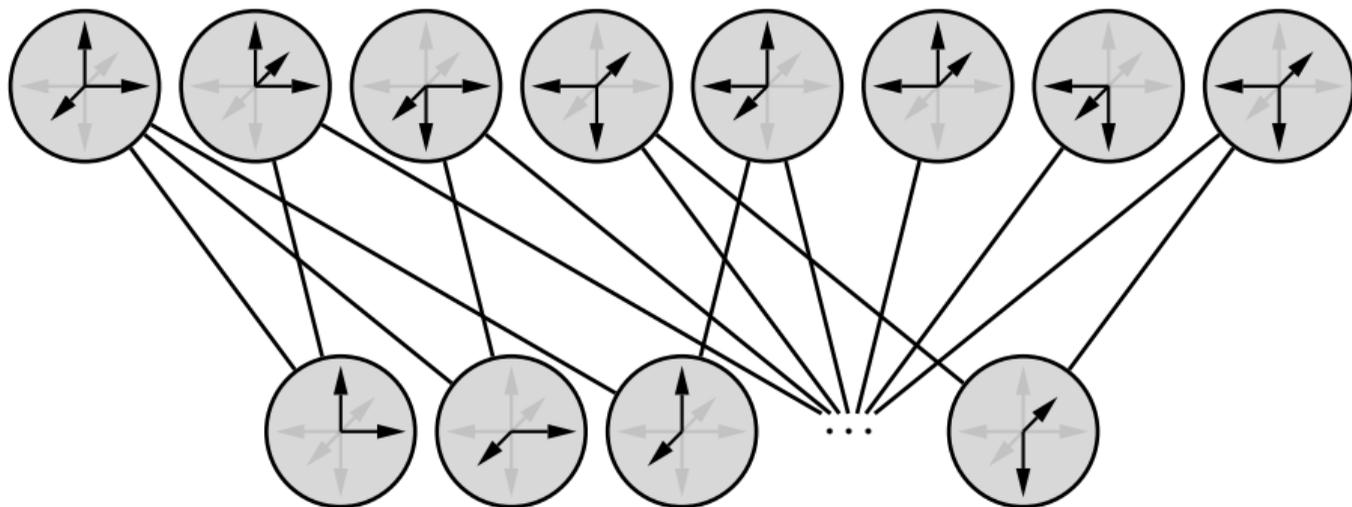
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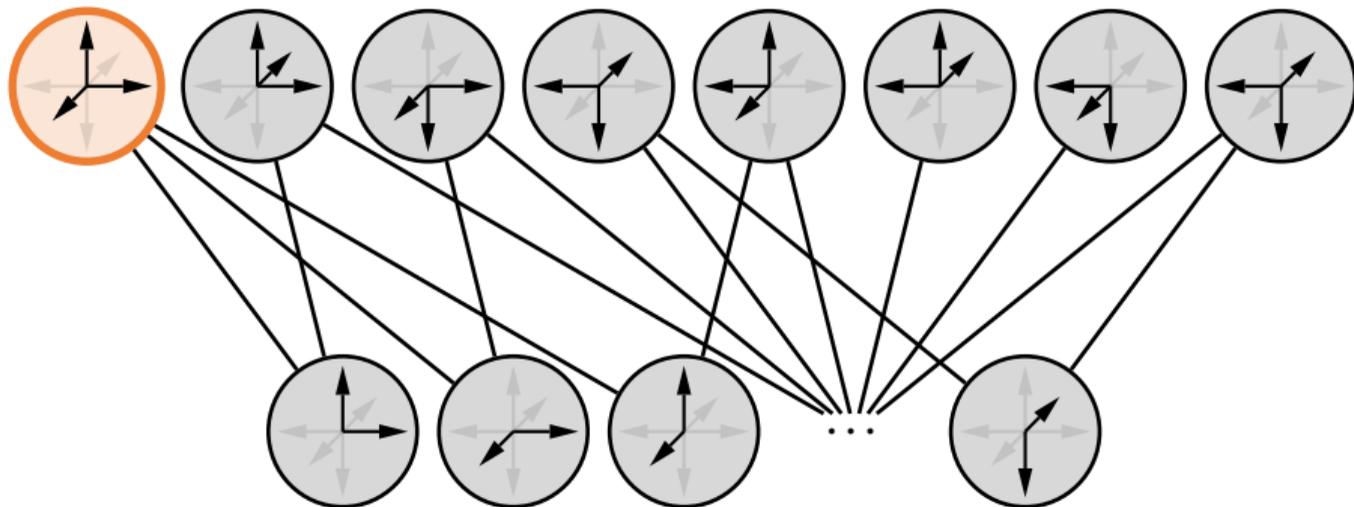
- ▶ Matroids conveniently already of this form.
- ▶ Natural random walk between  $\binom{[n]}{k}$  and  $\binom{[n]}{k-1}$ .



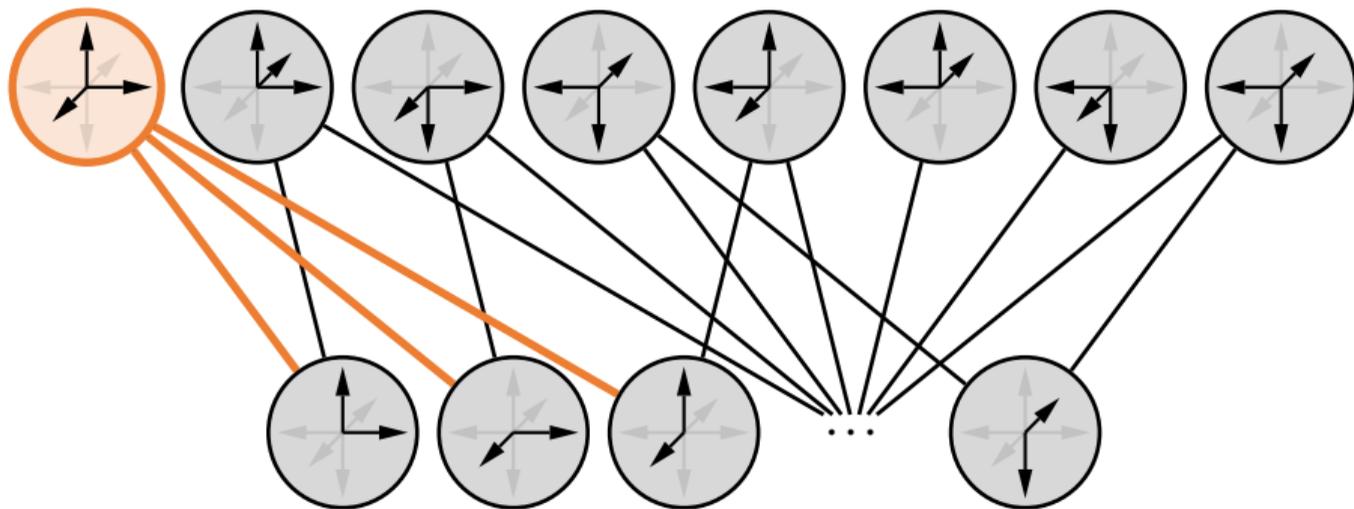
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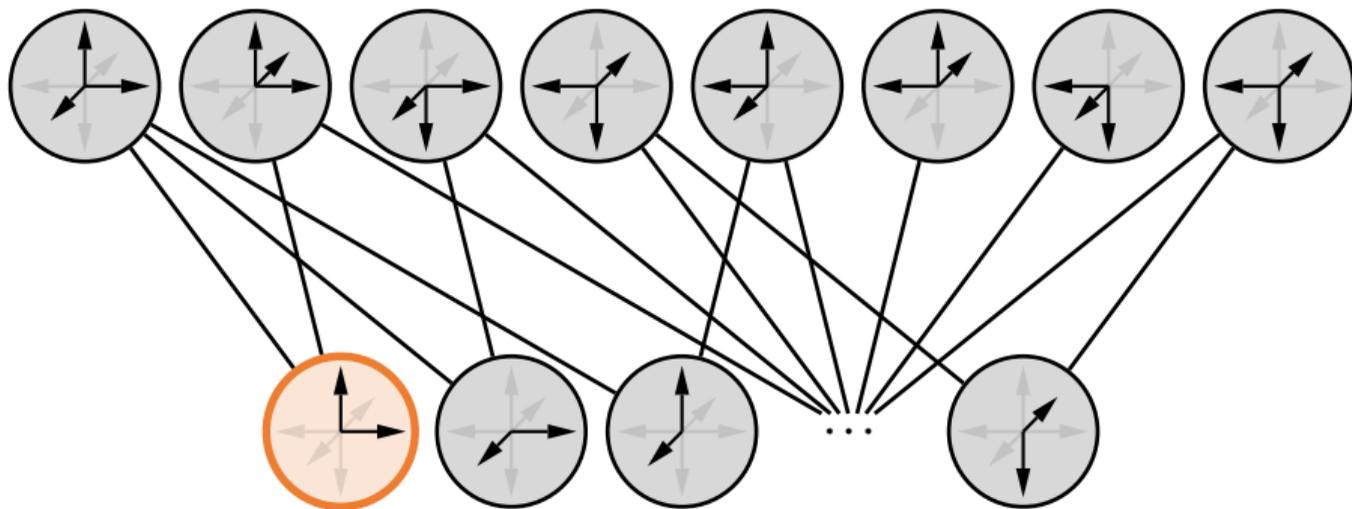


# Random Walk



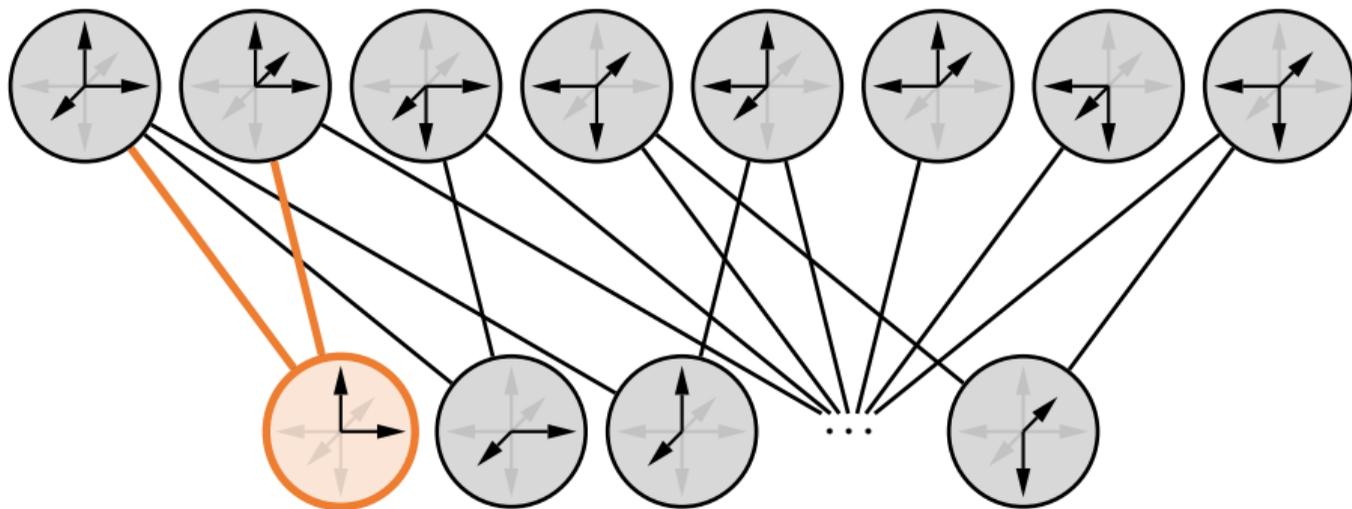
- 1 Drop an element uniformly at random.

# Random Walk



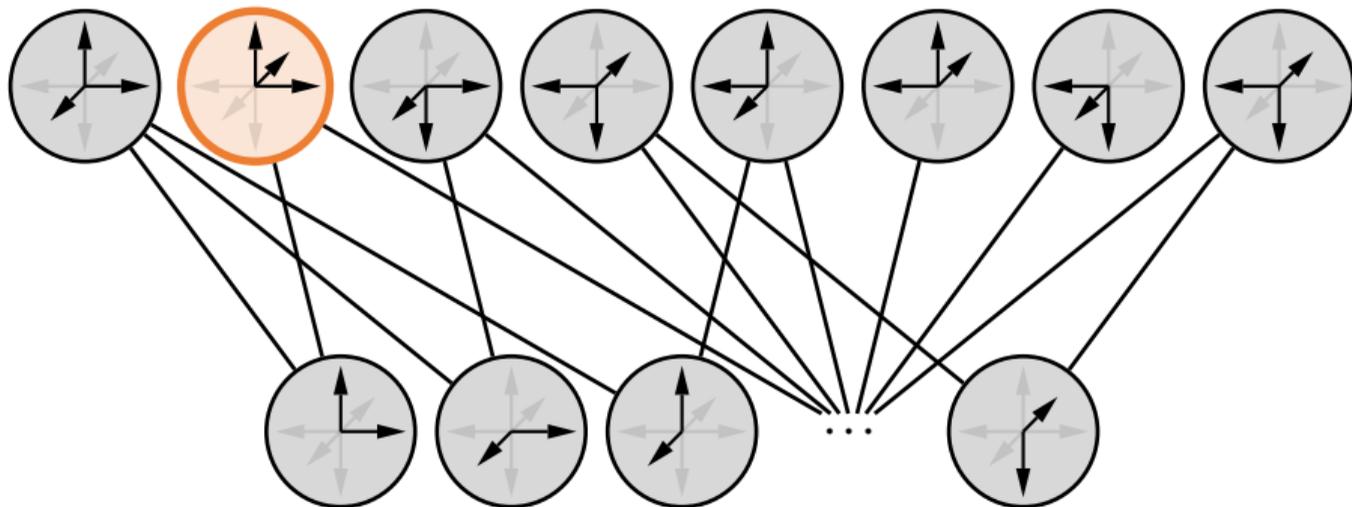
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- 1 Drop an element uniformly at random.
- 2 Add an element with probability  $\propto \mu(\text{resulting set})$ .

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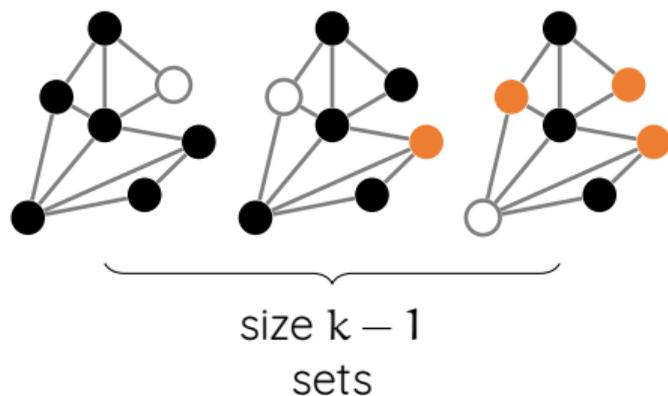
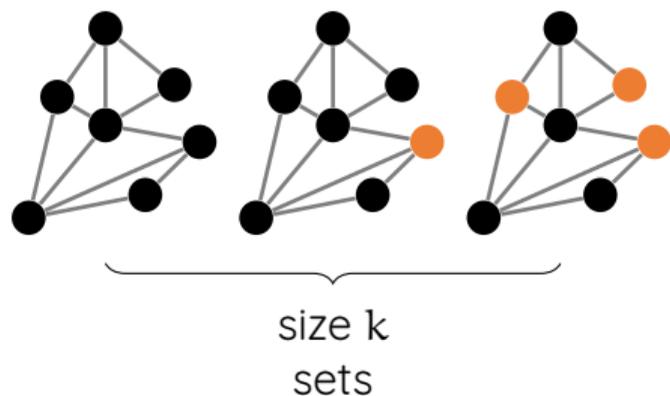
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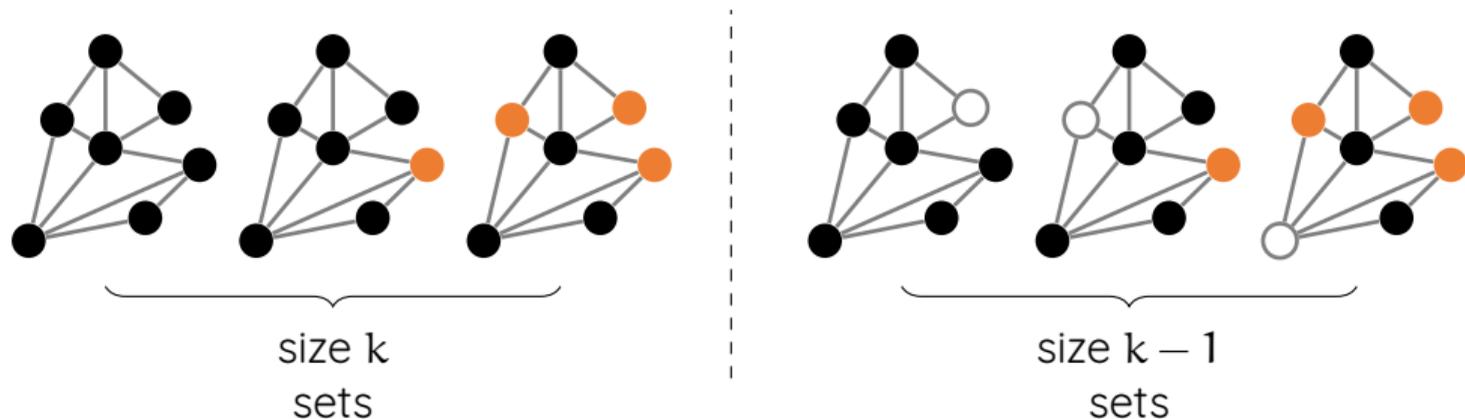
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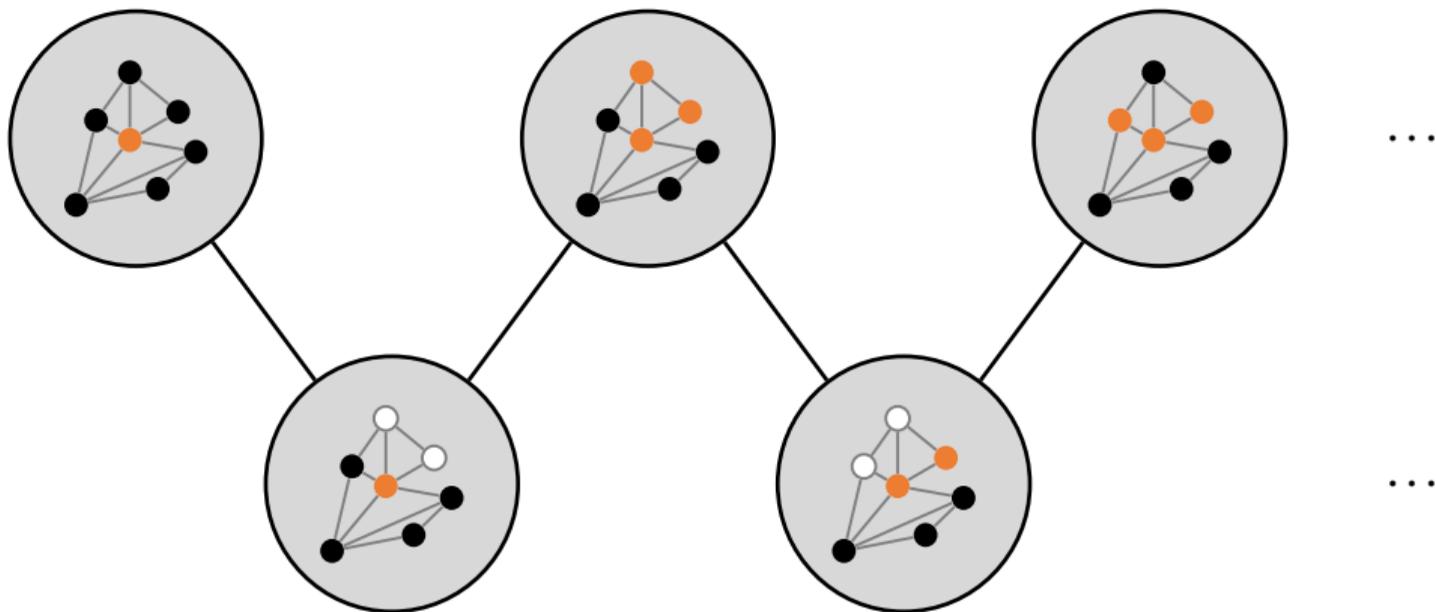
- ▶ Hardcore model and monomers are not naturally in canonical form.
- ▶ There is a canonical form with  $n = 2 \times \# \text{vertices}$  and  $k = \# \text{vertices}$ .



- ▶ Random walk is Glauber dynamics:
  - 1 Unmark vertex uniformly at random.
  - 2 Make a choice for it with probability  $\propto \mu(\text{resulting configuration})$ .

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- ▶ Random walk between  $k$ -sets and  $(k - 2)$ -sets:

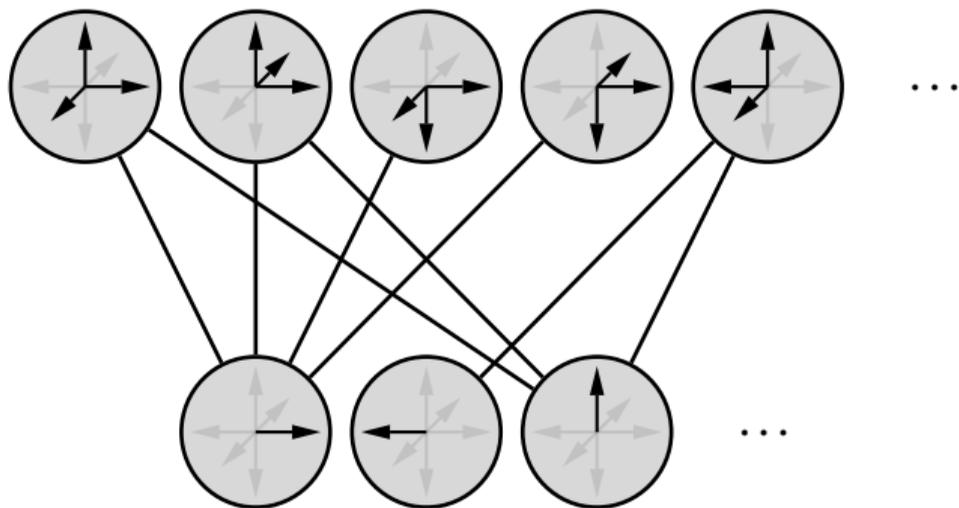


# Extreme random walks

- ▶ Random walk between  $k$ -sets and  $0$ -sets trivially mixes in 1 step.

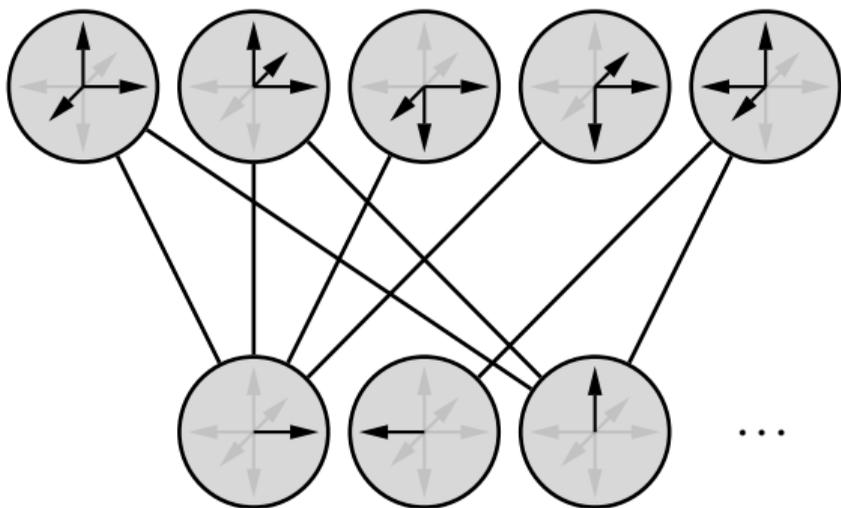
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... Transition probability matrix from 1-sets to 1-sets:

$$\frac{1}{k} \begin{bmatrix} \vdots & & \dots & & \vdots \\ \vdots & \mathbb{P}_{S \sim \mu}[j \in S \mid i \in S] & & & \vdots \\ \vdots & & \dots & & \vdots \end{bmatrix}$$

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- ▶ Obtain the following for all of our examples:

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- ▶ Show that this implies  $\text{poly}(k)$  relaxation time for the random walk between  $k$ -sets and  $(k - O(1))$ -sets.

# Fractional log-concavity

To our distribution  $\mu : \binom{[n]}{k} \rightarrow \mathbb{R}_{\geq 0}$  we can associate polynomial

$$g(z_1, \dots, z_n) = \sum_{S \in \binom{[n]}{k}} \mu(S) \prod_{i \in S} z_i.$$

## Fractional Log-Concavity

We have  $\lambda_{\max}(\text{correlation matrix}) = O(1)$  if and only if

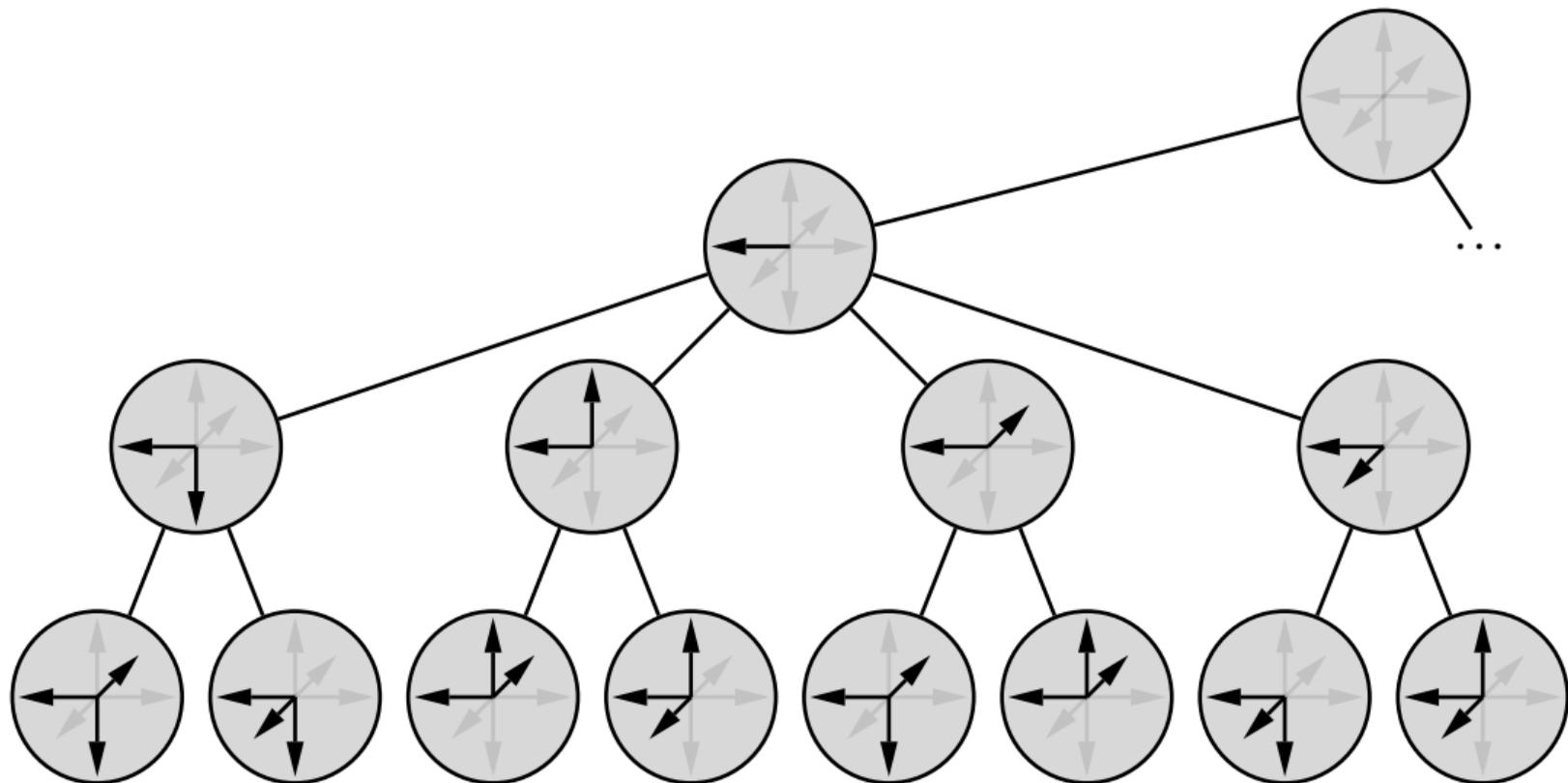
$$g(z_1^\alpha, \dots, z_n^\alpha)$$

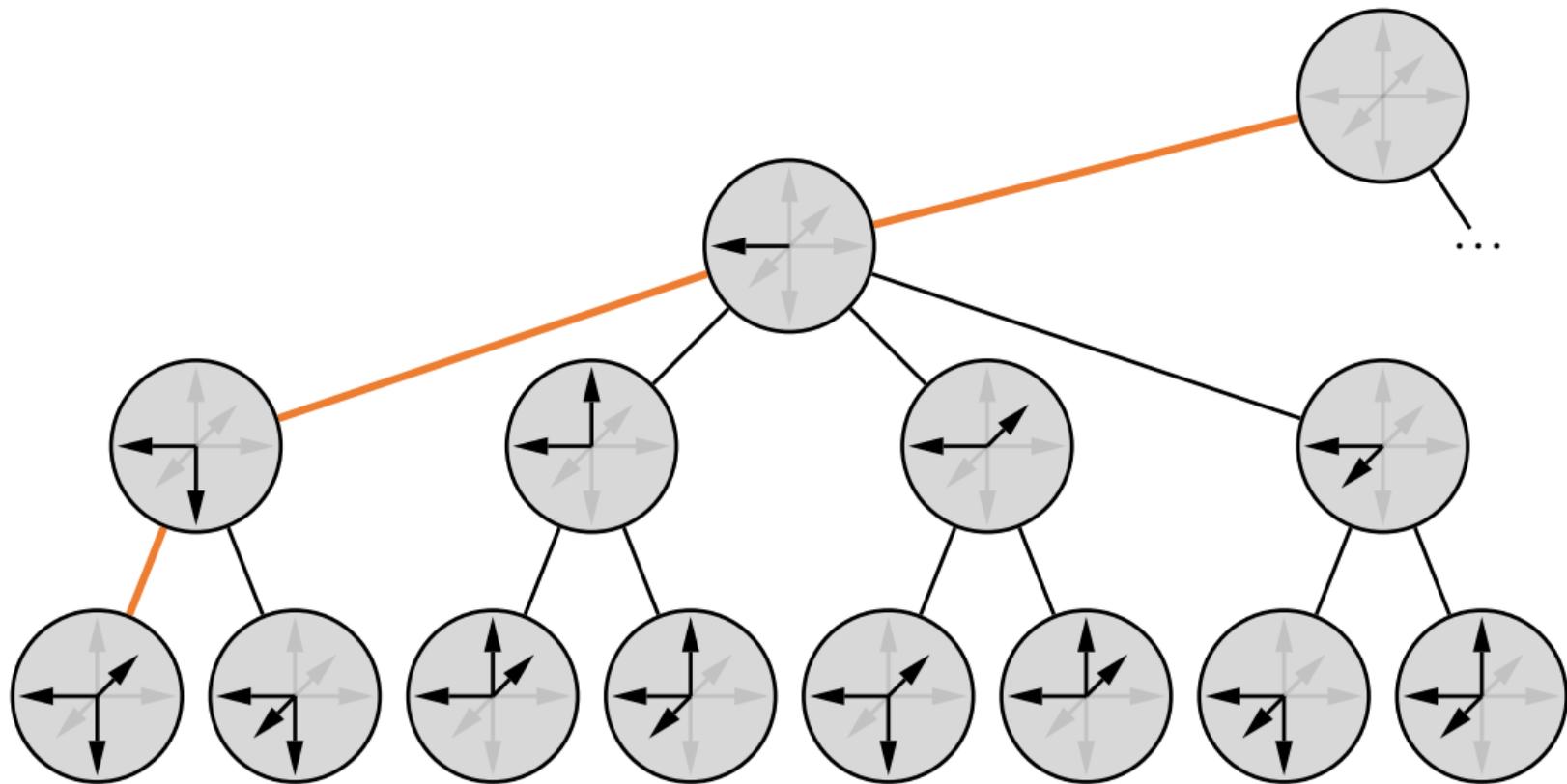
is log-concave around  $z_1 = \dots = z_n = 1$  for some  $\alpha = \Omega(1)$ .

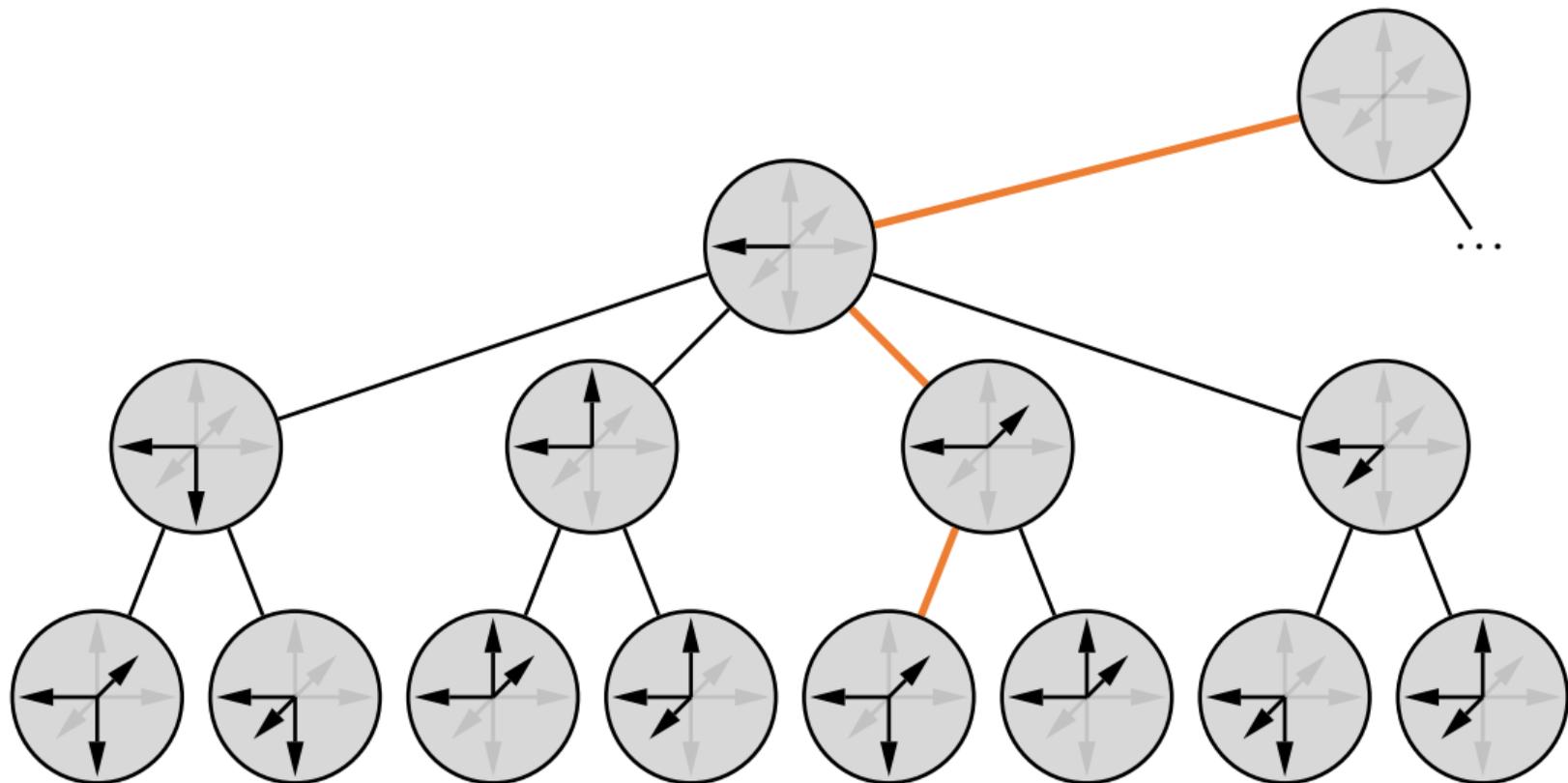
# Why does fractional log-concavity imply rapid mixing?

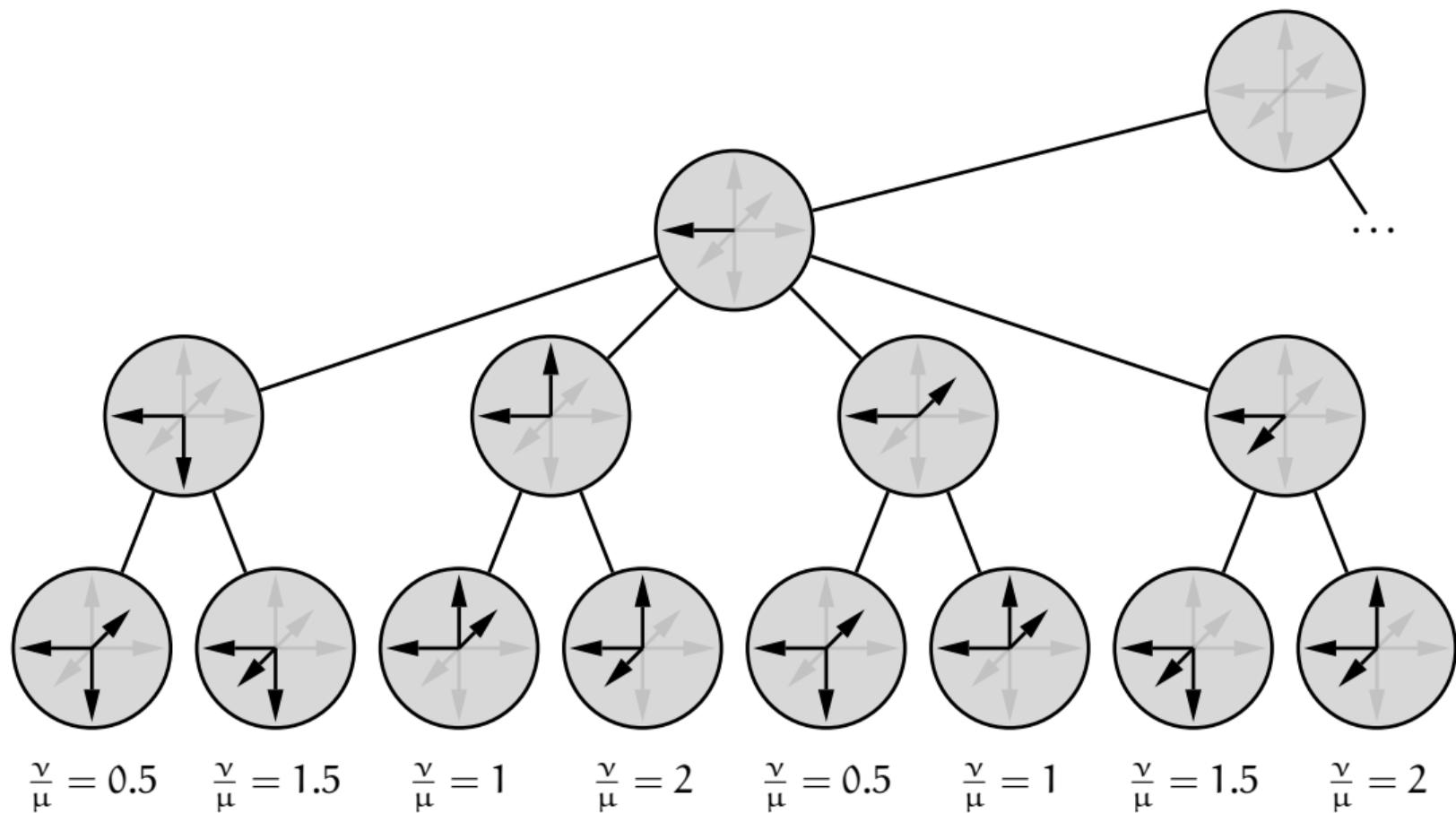
Local-to-global expansion phenomenon

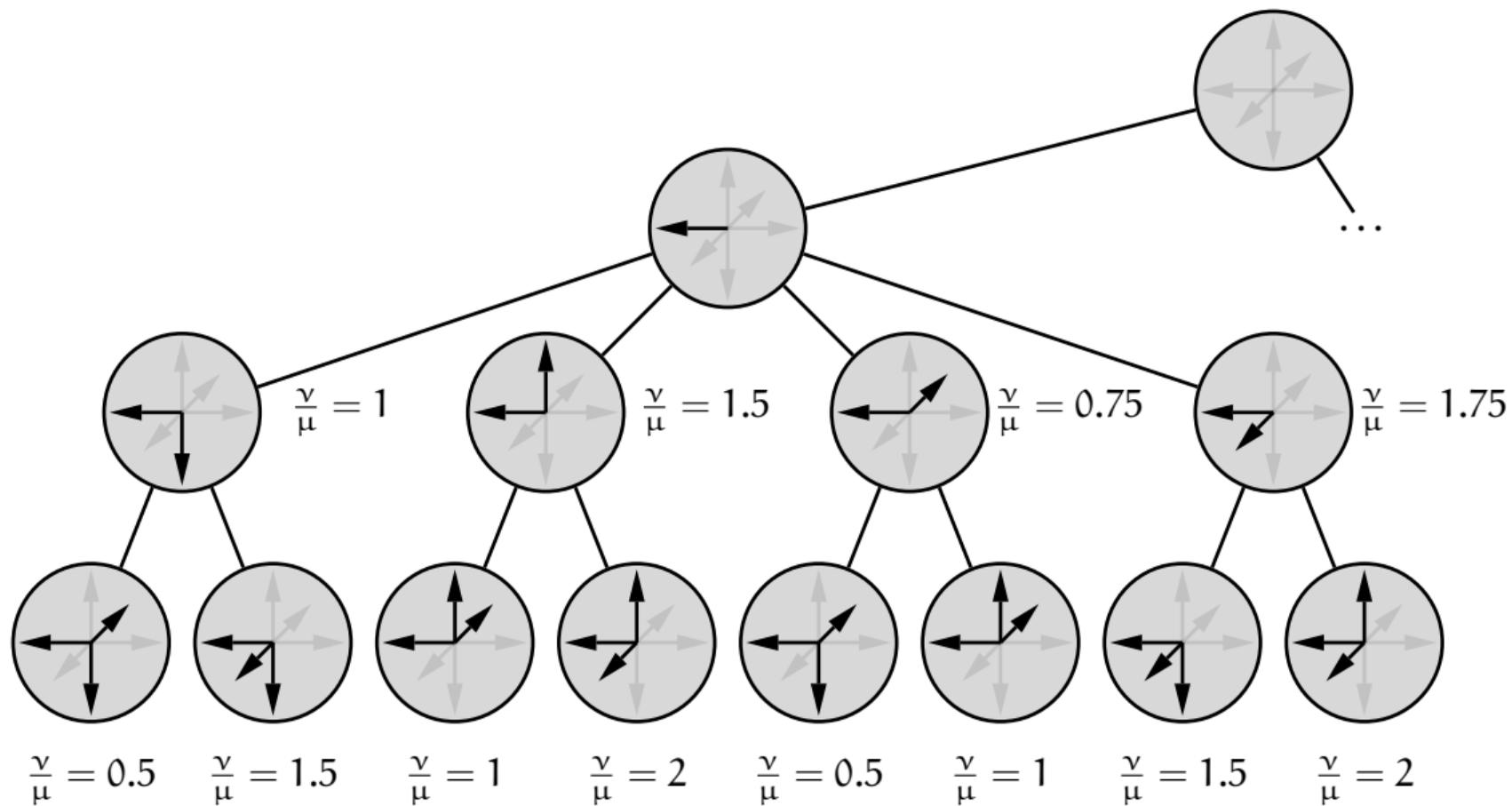
[Kaufman-Oppenheim'17, Cryan-Guo-Mousa'19, Alev-Lau'19,  
A-Liu-OveisGharan-Vinzant'19].

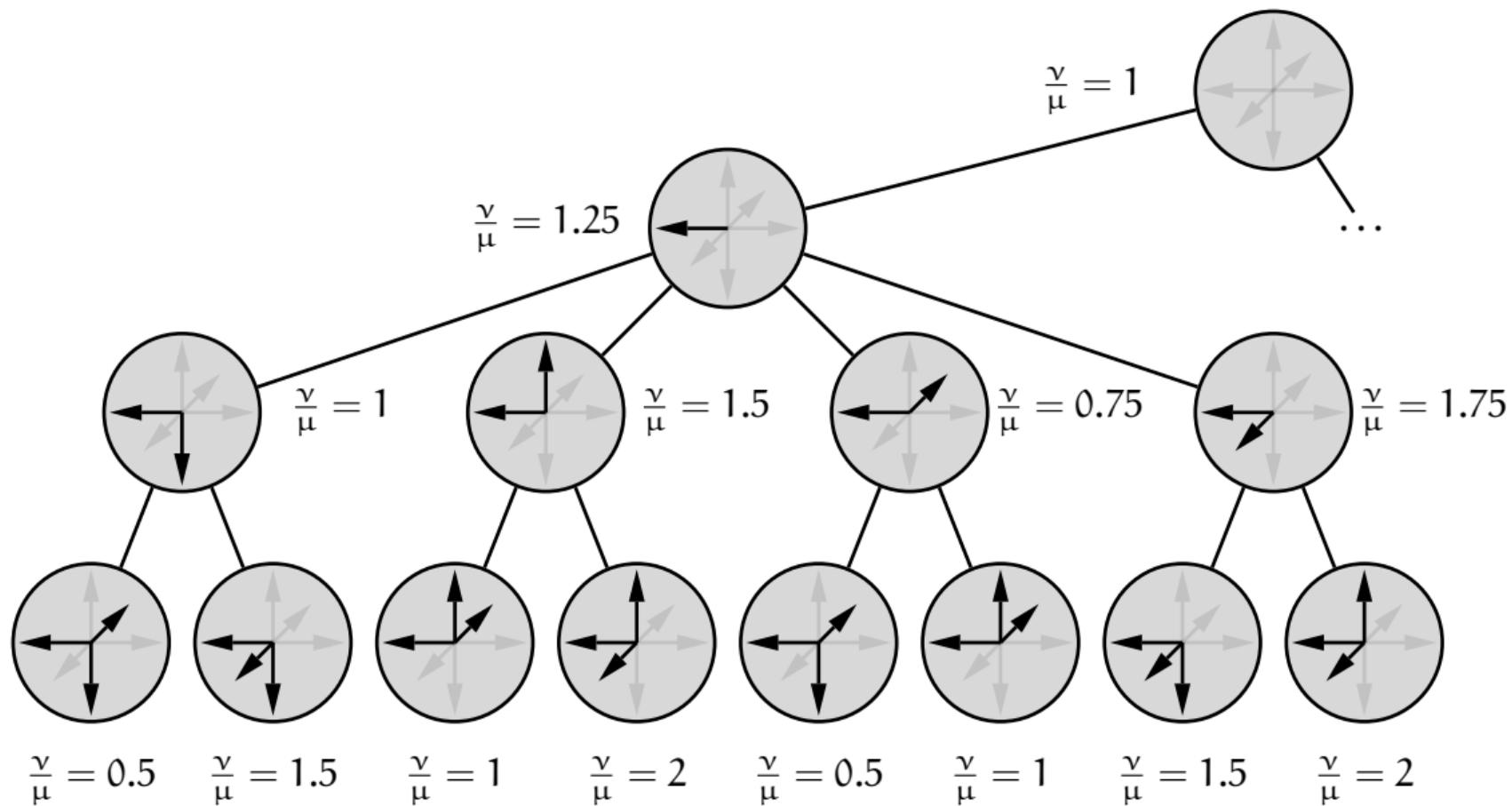






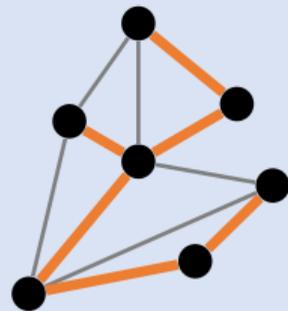






# Negative correlations

## Negative Correlations



random spanning tree

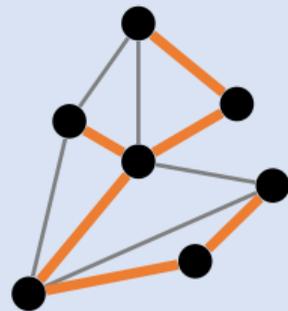
$$\mathbb{P}[\text{tree } T] \propto 1$$

# Negative correlations

- The  $\ell_1$ -norm of rows of correlation matrix:

$$\sum_f |\mathbb{P}[f \in T | e \in T] - \mathbb{P}[f \in T]|.$$

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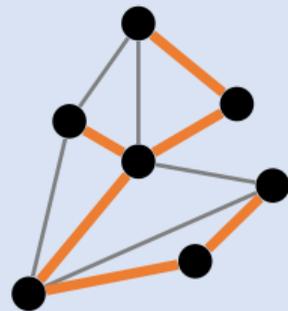
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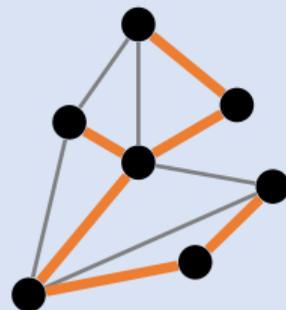
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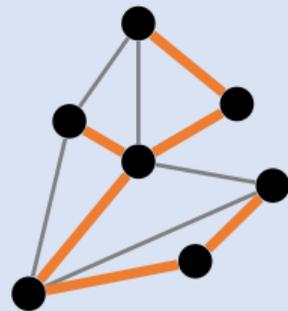
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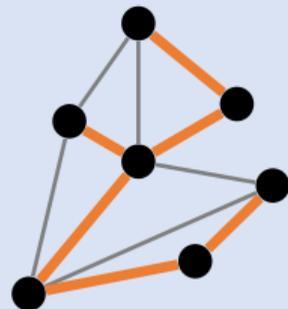
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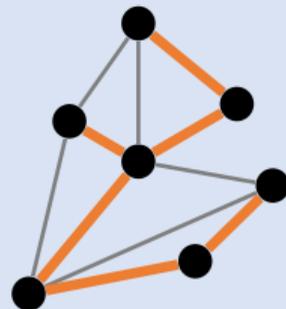
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- ▶ This implies  $\lambda_{\max} = O(1)$ .

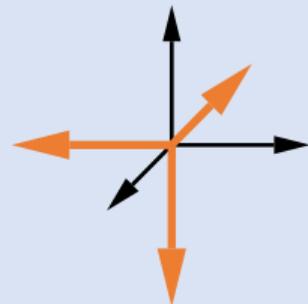
## Negative Correlations



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## Log-Concave Polynomial

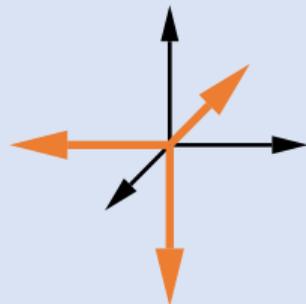


matroid  
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▶ Previous talk implies

$$\lambda_{\max}(\text{correlation matrix}) = O(1).$$

## Log-Concave Polynomial



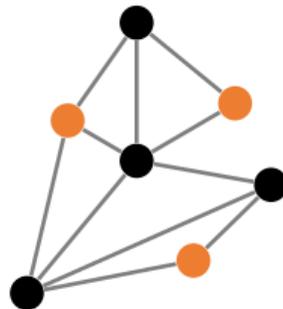
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## Hardcore Model

Given graph  $G = (V, E)$  and  $\lambda > 0$ ,  
sample a stable set  $S$  with

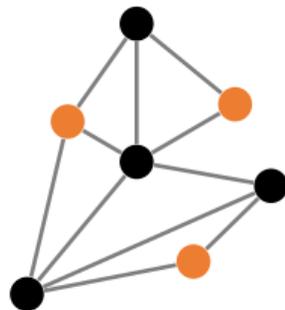
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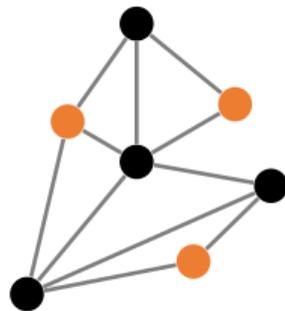
► When degrees are bounded by  $\Delta$ , there is a computational threshold

$$\lambda_c(\Delta) := (\Delta - 1)^{\Delta-1} / (\Delta - 2)^\Delta \simeq e / (\Delta - 2).$$

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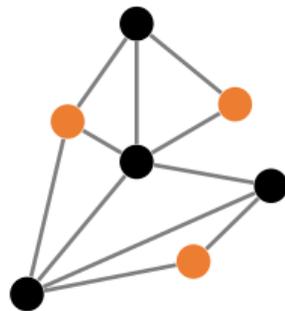
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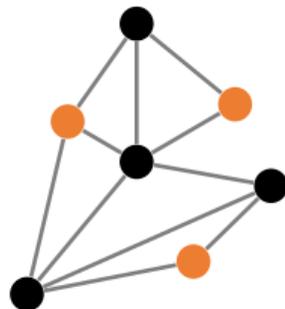
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▶ When  $\lambda < (1 - \epsilon)\lambda_c$ , sampling is possible in time  $n^{O(f(\epsilon) \log \Delta)}$  [Weitz'06].

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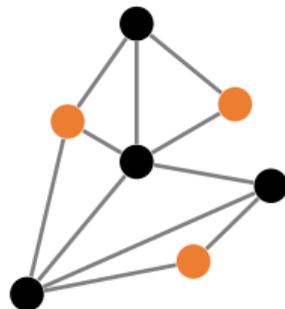
$$\lambda_c(\Delta) := (\Delta - 1)^{\Delta-1} / (\Delta - 2)^\Delta \simeq e / (\Delta - 2).$$

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## New Result [A-Liu-OveisGharan'19]

Normal Glauber dynamics mixes in time  $n^{f(\epsilon)}$ .

- ▶ Correlation between  $\mathbf{u}$  and  $\mathbf{v}$  decays exponentially

$$|\mathbb{P}[\mathbf{u} | \mathbf{v}] - \mathbb{P}[\mathbf{u}]| \leq e^{-\Omega(\text{dist}(\mathbf{u}, \mathbf{v}))}.$$

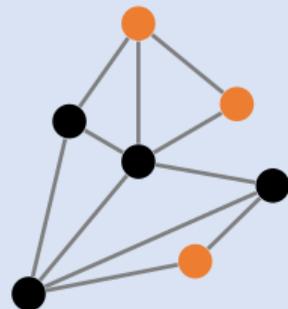
- ▶ Use this to bound  $\ell_1$  norm of columns of correlation matrix

$$\sum_{\mathbf{v}} |\mathbb{P}[\mathbf{u} | \mathbf{v}] - \mathbb{P}[\mathbf{u}]|$$

- ▶ Unfortunately there are  $(\Delta - 1)^d$  nodes at distance  $d$ , so not trivial to use decay of correlation.
- ▶ Nevertheless, Weitz's self-avoiding walk tree recursion can still be used to show the  $\ell_1$  norm is  $O_\epsilon(1)$ .

# Monomers

## Monomer Distribution



monomer distribution

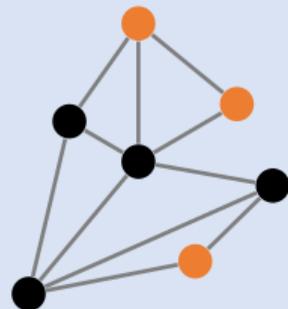
$$\mathbb{P}[S] \propto \text{\#perfect matchings in } S^c$$

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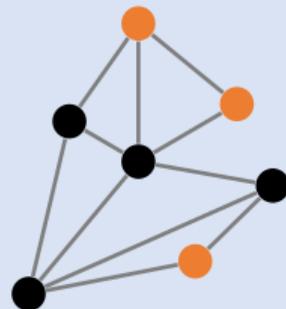
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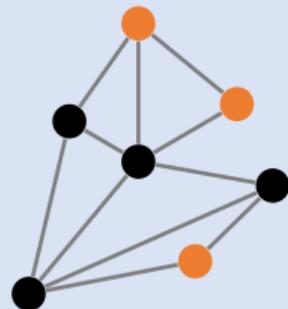
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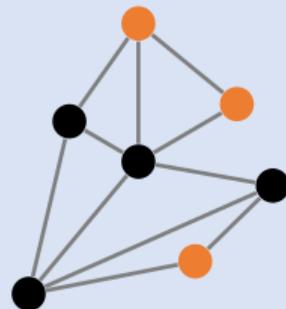
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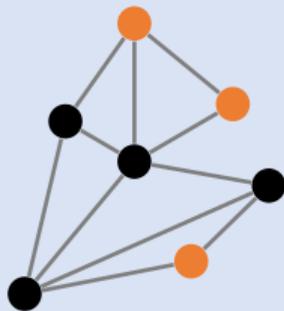
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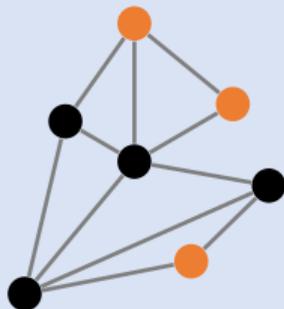
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- ▶ This implies efficient sampling for the monomer-dimer systems on planar graphs (hardness shown by [Jerrum'97]).

## Monomer Distribution



monomer distribution

$$\mathbb{P}[S] \propto \# \text{perfect matchings in } S^c$$

► Let  $T$  be the set of positive terms in the  $\ell_1$  norm:

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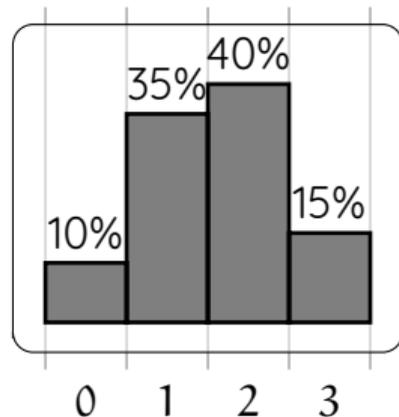
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▶ We will show that  $X/2$  is distributed as

$$\text{constant} + \text{Bernoulli}(p_1) + \dots + \text{Bernoulli}(p_m).$$

# Distributions $\leftrightarrow$ Polynomials



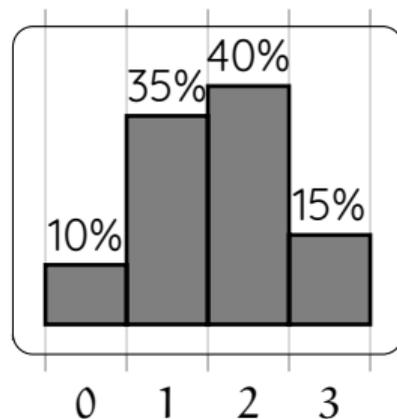
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## Generalized Binomials

Compute the roots of  $g(z)$  and verify that all are real (none are complex).

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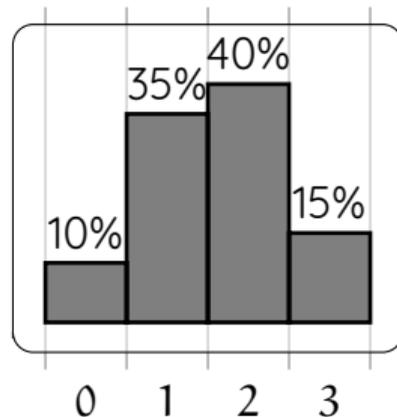
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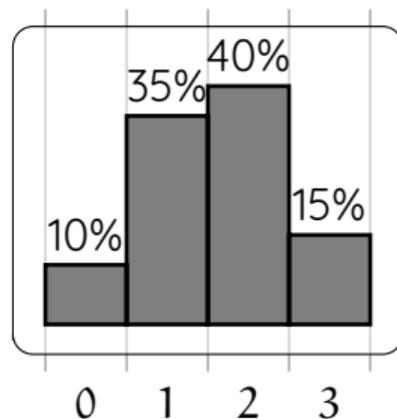
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► Roots correspond to biases:

$$\text{coin bias } p \leftrightarrow 1 - 1/p \text{ root of } g$$



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[Heilmann-Lieb'72]

If  $\mu$  is a monomer distribution, the polynomial

$$\sum_S \mu(S) \prod_{u \in S} z_u$$

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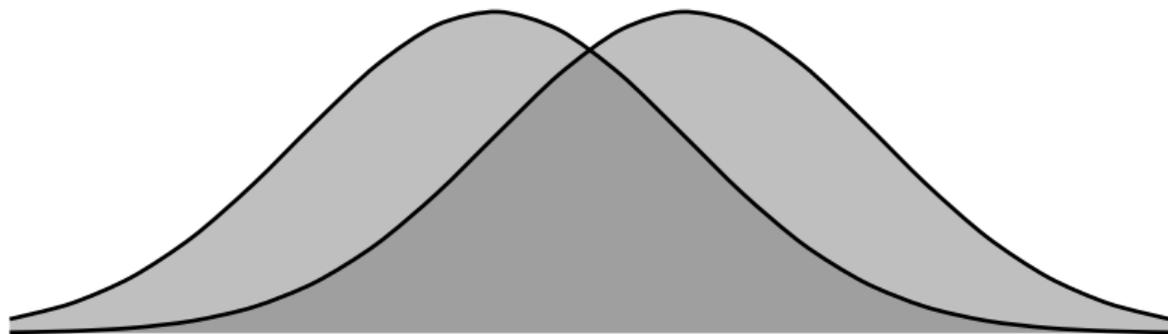
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Massaging this, we get that  $X$  is distributed as  
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# Test your intuition

▶ Two Gaussians. Are mixtures unimodal? Log-concave?

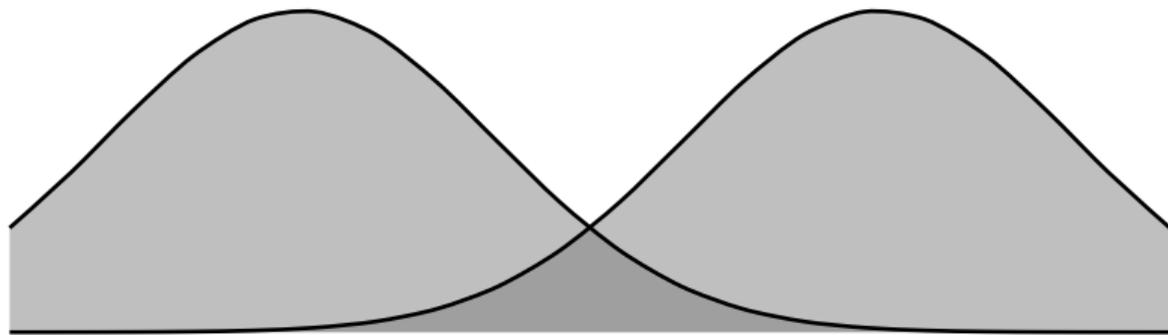


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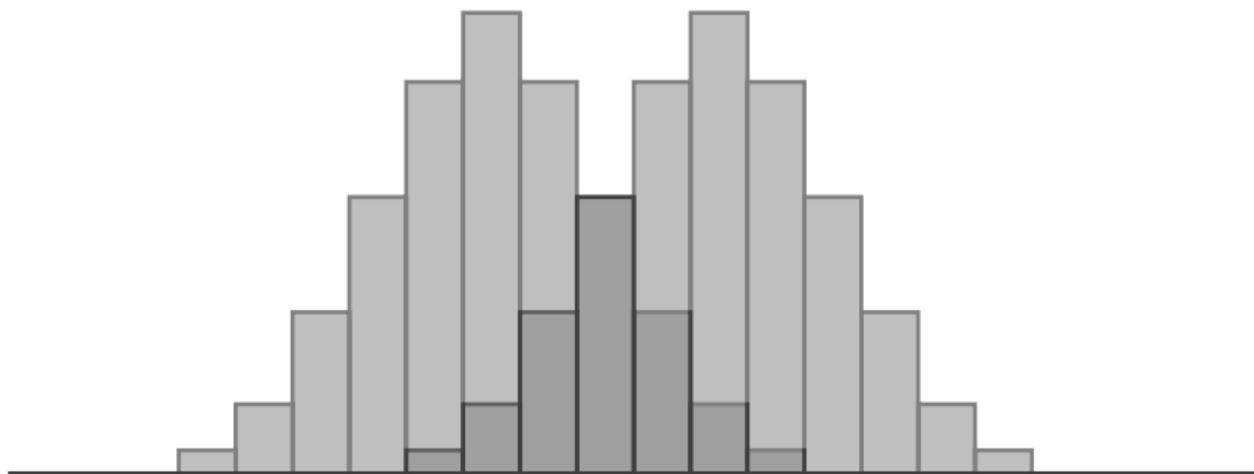


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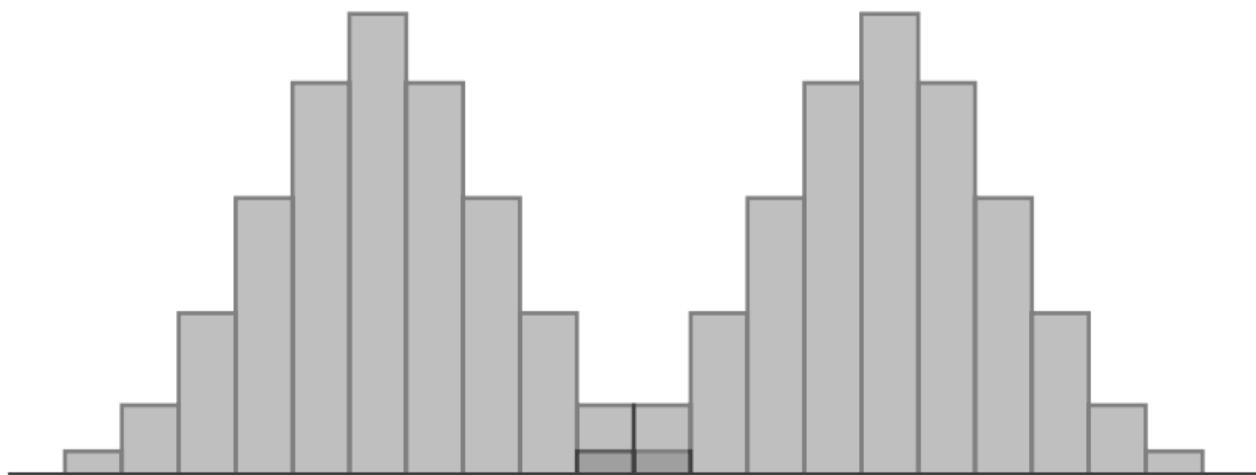


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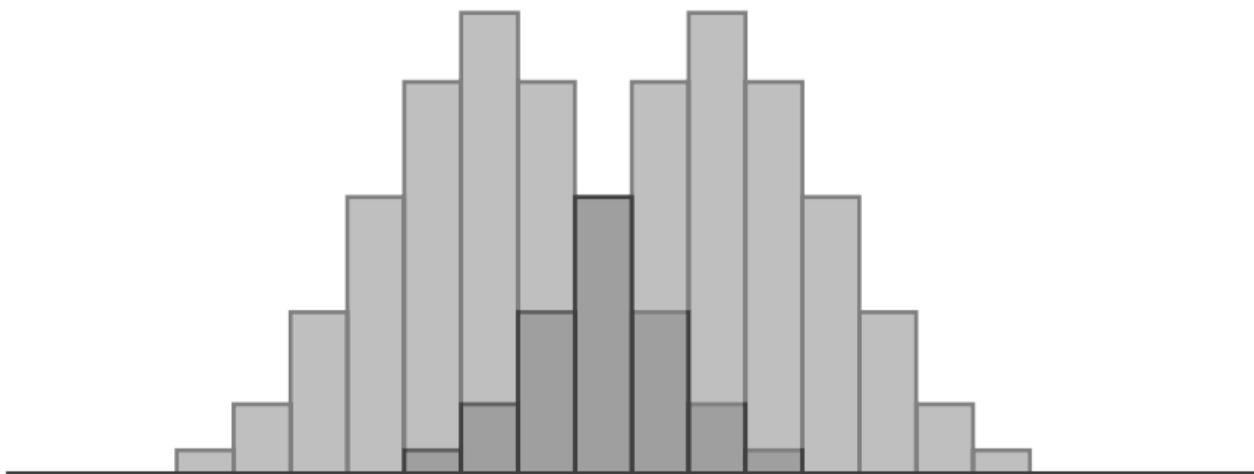


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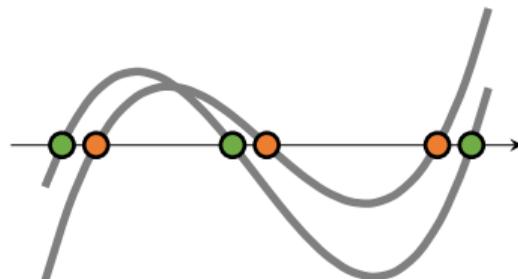
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[Folklore, used by e.g., MSS'13]

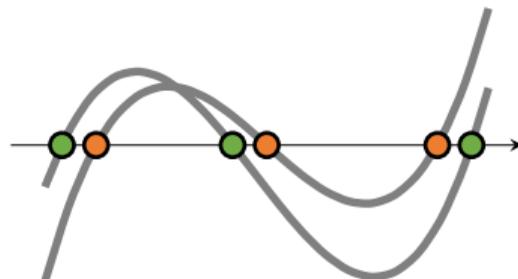
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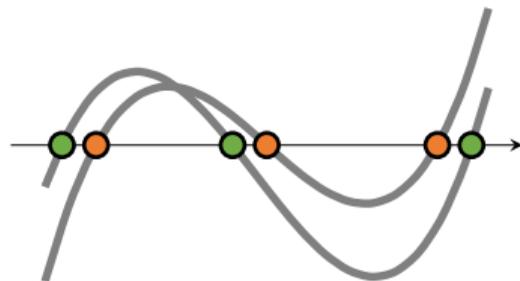
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► Corollary: The means of  $\mu$  and  $\nu$  can be off by  $\leq 1$ .

$$|(p_1 + \cdots + p_n) - (q_1 + \cdots + q_n)| \leq 1$$

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Thank you!