

Lecture 14

①

- Recall J-S MC for sampling matchings in unweighted graphs $G=(V, E)$, according to the Gibbs distn': $\pi_\lambda(m) = \frac{1}{Z(\lambda)} \lambda^{|m|}$.
- Running time: $\text{poly}(n, \lambda)$, $n=|V|$
- $Z(\lambda) = \sum_k m_k \lambda^k$ partition function (# of k-matchings) (matching polynomial)
- Last time: Estimation of $Z(\lambda)$.
- Today: Estimation of coefficients m_k .

Estimating coefficients m_k

exact estimation: when $k = \frac{|V|}{2}$, m_k is # perfect matchings if G is bipartite with n vertices on each side, computing $m_n \equiv$ computing $\text{per}(A)$

def: If A is $n \times n$, $\text{per}(A) = \sum_{\sigma} \prod_{i=1}^n A(i, \sigma(i))$, where the summation is over all permutations σ of $\{1, \dots, n\}$

(permanent of a matrix) adjacency matrix

contrast this, with the determinant:

def: If A is $n \times n$, the determinant of A is $\det(A) = \sum_{\sigma} (-1)^{\text{sign}(\sigma)} \prod_{i=1}^n A(i, \sigma(i))$

where $\text{sign}(\sigma)$ is # of pairs $\{i, j\}$ s.t. $\sigma(j) > \sigma(i)$

Claim 2: If $\lambda = \frac{m_{k-1}}{m_k}$, then $m_{k'} \lambda^{k'}$ is maximized at $k'=k$ & $k'=k-1$. (3)

Proof: First verify that: $m_k \lambda^k = m_k \lambda \cdot \lambda^{k-1} = m_{k-1} \lambda^{k-1}$

Moreover,

for all $k' \leq k-1$: $\frac{m_{k'-1} \lambda^{k'-1}}{m_{k'} \lambda^{k'}} = \frac{1}{\lambda} \cdot \frac{m_{k'-1}}{m_{k'}} = \frac{m_k}{m_{k-1}} \cdot \frac{m_{k'-1}}{m_{k'}} \leq \frac{m_{k'} \cdot m_{k'-1}}{m_{k'-1} \cdot m_{k'}} \leq 1$ using claim 1

for all $k' \geq k$: $\frac{m_{k'+1} \lambda^{k'+1}}{m_{k'} \lambda^{k'}} = \lambda \frac{m_{k'+1}}{m_{k'}} = \frac{m_{k-1}}{m_k} \cdot \frac{m_{k'+1}}{m_{k'}} \leq \frac{m_{k'} \cdot m_{k'+1}}{m_{k'+1} \cdot m_{k'}} \leq 1$ using claim 1

□

Claims 1, 2 suggest the following algorithm; for estimating $\frac{m_{k'}}{m_{k'-1}}$
 for $k'=2, \dots, k$: (starting at $\frac{m_0}{m_1} = \frac{1}{|E|}$)
 i) gradually raise λ and sample from Gibbs distn π_λ repeatedly until we see "lots of" $(k'-1)$ - and k' -matchings, where "lots of" \equiv they appear w/ prob. at least $\frac{\epsilon}{n}$, for some constant c (Claim 2, confirms that we'll not need to raise λ beyond $\frac{m_{k'+1}}{m_{k'}}$)

ii) For the λ chosen in step i, sample MC repeatedly to estimate $\frac{m_{k'}}{m_{k'-1}}$; take as estimator the ratio of the number of k' -matchings to the number of $(k'-1)$ -matchings multiplied by $\frac{1}{\lambda}$.

ex 1pt: Can estimate $\frac{m_{k'}}{m_{k'-1}}$ to within a factor of $(1 \pm \frac{\epsilon}{n})$.

w/ probability at least $1 - \frac{1}{4n}$ in time $\text{poly}(n, \frac{1}{\epsilon}, \frac{m_{k'-1}}{m_{k'}})$.

- computing $\det(A)$ can be done efficiently w/ Gaussian Elimination
- computing $\text{per}(A)$ is #P-complete (Valiant '79).

estimation of m_k :

- if we could compute $Z(\lambda)$ exactly at any desired λ , we would obtain all the m_k 's exactly; indeed $Z(\lambda)$ is a polynomial of degree n w/ coefficients m_0, \dots, m_n , so we could do that by computing $Z(\lambda)$ at $n+1$ points and interpolating.
- however, approach is not robust to error.
- We use a different approach, aiming at estimating ratios of the form $\frac{m_k}{m_{k-1}}$ and taking a telescoping product:

$$m_k = \left(\prod_{k'=1}^k \frac{m_{k'}}{m_{k'-1}} \right) \cdot m_0 \quad (*)$$

$\nwarrow m_0 = 1$ (trivially)
 so we only need estimates of $\frac{m_{k'}}{m_{k'-1}}$ for all $k'=1, \dots, k$.

- Claim 1: The sequence $\{m_k\}_k$ is log-concave, i.e.

$$m_{k-1} \cdot m_{k+1} \leq m_k^2, \text{ for all } k.$$

exercise (1 pt): Show Claim 1, by defining an injective mapping from pairs of $(k+1)$ and $(k-1)$ -matchings to pairs of k -matchings.

- plugging the above estimator into (*) and noticing that the worst possible ratio $\frac{m_{k-1}}{m_k}$ is $\frac{m_{k-1}}{m_k}$ (by claim 1), we obtain that we can estimate m_k to within a factor of $(1 \pm \epsilon)$ w/ probability $\geq \frac{3}{4}$ in time $\text{poly}(n, \frac{1}{\epsilon}, \frac{m_{k-1}}{m_k})$.

NOTE: there is nothing special about $\frac{3}{4}$; we can boost this by repeating many times & outputting the median of the results.

Ex 1 pt: If k^* is the (unknown) size of a maximum matching in a given graph, show how to use the above algorithm to compute a matching of size $\geq (1-\epsilon) \cdot k^*$ in time $n^{O(1/\epsilon)}$.

• How large can $\frac{m_{k-1}}{m_k}$ be?

- In many interesting classes of graphs, such as
 - dense graphs (every degree is $\geq \frac{n}{2}$)
 - random graphs (e.g. in the $(n, 1/2)$ model, whp)
 - regular lattices (finite square regions of \mathbb{Z}^2)

the ratio $\frac{m_{k-1}}{m_k}$ is polynomially bounded in n .

- However, in general it can be very bad, e.g.



- $|V| = 4l + 2 = 2^{2l}$, where $l = \# \text{ squares}$
easy to see $m_n = 1$ (unique perfect matching), but $m_{n-1} \geq 2^l$.