COMPSCI 614: Randomized Algorithms with Applications to Data Science

Prof. Cameron Musco University of Massachusetts Amherst. Spring 2024. Lecture 21 I released Problem Set 5 yesterday, due 5/13 at 11:59pm.
This problem set is optional – it can be used to replace your lowest grade on the first four problem sets.

Last Time: Markov Chain Mixing Times

- Total variation distance and it<u>s dual characterizations</u>.
- Basic results on mixing time.
- Coupling as a technique for bounding mixing time.

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Today: Mixing Time Analysis

- Finish up coupling and example applications.
- Start on algorithmic applications Markov Chain Monte Carlo (MCMC).

Total Variation Distance

Definition (Total Variation (TV) Distance)

For two distributions $p, q \in [0, 1]^m$ over state space [m], the total variation distance is given by:

$$\|p-q\|_{TV} = \frac{1}{2} \sum_{i \in [m]} |p(i) - q(i)| = \max_{A \subseteq [m]} |p(A) - q(A)|.$$

Kontorovich-Rubinstein duality: Let P, Q be possibly correlated random variables with marginal distributions p, q. Then

$$\|p - q\|_{TV} \leq \Pr[P \neq Q].$$

$$\|p - \gamma\|_{TV} \leq \Pr[P \neq Q].$$

This dual notion is the key idea behind mixing time analysis via coupling.

Definition (Mixing Time)

Consider a Markov chain X_0, X_1, \ldots with unique stationary distribution π . Let $q_{i,t}$ be the distribution over states at time t assuming $X_0 = i$. The mixing time is defined as:

$$au(\epsilon) = \min\left\{t: \max_{i\in[m]} \|q_{i,t} - \pi\|_{TV} \le \epsilon\right\}.$$

Definition (Coupling)

1.
$$\mathbf{X}_0 = i$$
 and $\mathbf{Y}_0 = j$ for some $i, j \in [m]$.

2.
$$\Pr[\mathbf{X}_t = j | \mathbf{X}_{t-1} = i] = \Pr[\mathbf{Y}_t = j | \mathbf{Y}_{t-1} = i] = P_{i,j}$$

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Definition (Coupling)

For a finite Markov chain $X_0, X_1, ...$ with transition matrix $P \in \mathbb{R}^{m \times m}$, a coupling is a joint process $(X_0, Y_0), (X_1, Y_1), ...$ such that:

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$$\mathbf{X}_0 = i$$
 and $\mathbf{Y}_0 = j$ for some $i, j \in [m]$.

2.
$$\Pr[\mathbf{X}_t = j | \mathbf{X}_{t-1} = i] = \Pr[\mathbf{Y}_t = j | \mathbf{Y}_{t-1} = i] = P_{i,j}$$

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Theorem (Mixing Time Bound via Coupling)

For a finite, irreducible, and aperiodic Markov chain X_0, X_1, \ldots and any valid coupling $(X_0, Y_0), (X_1, Y_1), \ldots$ letting $T_{i,j} = \min\{t : X_t = Y_t | X_0 = i, Y_0 = j\},$

$$\max_{i \in [m]} \|q_{i,t} - \pi\|_{TV} \leq \max_{i,j \in [m]} \|q_{i,t} - q_{j,t}\|_{TV} \leq \max_{i,j \in [m]} \Pr[\mathsf{T}_{i,j} > t]. \quad \leq \varepsilon$$

How many times do we need to swap a random card to the top of the deck so that the distribution of orderings on our cards is ϵ -close in TV distance to the uniform distribution over all permutations?

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- Can check that this is a valid coupling since X_t, Y_t have the correct marginal distributions, and since
 X_t = Y_t ⇒ X_{t+1} = Y_{t+1}
- Observe that X_t = Y_t as soon as all c unique cards have been swapped at least once. How many swaps does this take?

$$\begin{split} \max_{i \in [m]} \|q_{i,t} - \pi\|_{TV} &\leq \max_{i,j \in [m]} \Pr[\mathsf{T}_{i,j} > t] \\ &\leq \Pr[< c \text{ unique cards are swapped in } t \text{ swaps}] \end{split}$$

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$$\leq \Pr[< c \text{ unique cards are swapped in } t \text{ swaps}]$$

By coupon collector analysis for $t \ge c \ln(c/\epsilon)$, this probability is bounded by ϵ . In particular, by the fact that $\left(1 - \frac{1}{c}\right)^{c \ln c/\epsilon} \le \frac{\epsilon}{c}$ plus a union bound over c cards.

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Thus, for
$$t \ge c \ln(c/\epsilon)$$
,
 $\max_{i\in[m]} \|q_{i,t} - \pi\|_{TV} \le \max_{i,j\in[m]} \|q_{i,t} - q_{j,t}\|_{TV} \le \epsilon$
I.e., $\tau(\epsilon) \le c \ln(c/\epsilon)$.

Let X_0, X_1 be a Markov chain over state space $\{0, 1\}^n$. In each step, pick a random position $i \in [n]$ and set $X_t(i) = 0$ with probability 1/2 and $X_t(i) = 1$ with probability 1/2.



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What is a coupling $(X_0, Y_0), (X_1, Y_1), \ldots$ on this chain that we can use to bound the mixing time of this walk?

In each step, pick a single random position $i \in [n]$ and let $X_t(i) = Y_t(i) = 0$ with probability 1/2 and $X_t(i) = Y_t(i) = 1$ with probability 1/2.



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How large must we set t so that $\Pr[X_t \neq Y_t] \leq \epsilon$?

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Upshot: The mixing time of the *n*-dimensional hypercube is $\tau(\epsilon) = O(n \log(n/\epsilon)).$

Claim: If X_0, X_1, \ldots is finite, irreducible, and aperiodic, then for any c < 1/2 and any $\epsilon > 0$, $\tau(\epsilon) \le \tau(c) \cdot O(\log(1/\epsilon))$.

I.e., it suffices to bound the mixing time for any small constant c and then can boost this result to any $\epsilon > 0$.

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Proof:

• After $t = \tau(c)$ steps, for any i we have $||q_{i,t} - \pi||_{TV} \le c$. So, for any i, j we have $||q_{i,t} - q_{j,t}||_{TV} \le 2c < 1$.

Claim: If X_0, X_1, \ldots is finite, irreducible, and aperiodic, then for any $(\begin{array}{c} u & v \end{array})$ c < 1/2 and any $\epsilon > 0$, $\tau(\epsilon) \le \tau(c) \cdot O(\log(1/\epsilon))$.

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Proof:

- After $t = \tau(c)$ steps, for any i we have $||q_{i,t} \pi||_{TV} \le c$. So, for any i, j we have $||q_{i,t} q_{j,t}||_{TV} \le 2c < 1$.
- This implies a coupling between two chains X_0, X_1, \ldots and Y_0, Y_1, \ldots starting in any initial states such that $Pr[X_t \neq Y_t] \le 2c < 1.$

$$m = P_{\tau}(X_{t} \neq V_{t})^{2} = ||q_{ii+} = q_{j+}||_{TV}^{2} \geq 2c^{2}$$



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Claim: If X_0, X_1, \ldots is finite, irreducible, and aperiodic, then for any c < 1/2 and any $\epsilon > 0$, $\tau(\epsilon) \leq \underline{\tau(c)} \cdot O(\log(1/\epsilon))$.

I.e., it suffices to bound the mixing time for any small constant c and then can boost this result to any $\epsilon > 0$.

Proof:

- After $t = \tau(c)$ steps, for any i we have $||q_{i,t} \pi||_{TV} \le c$. So, for any i, j we have $||q_{i,t} q_{j,t}||_{TV} \le 2c < 1$.
- This implies a coupling between two chains X_0, X_1, \ldots and $Y_0, Y_{\overline{1}, \ldots}$ starting in any initial states such that $Pr[X_t \neq Y_t] \leq 2c < 1.$
- So after $\tau(c) \cdot O(\log(1/\epsilon))$ steps, $\Pr[X_t \neq Y_t] \leq (2c)^{O(\log 1/\epsilon)} \leq \epsilon$
- This establishes that $\tau(\epsilon) \leq \tau(c) \cdot O(\log(1/\epsilon))$.

Markov Chain Monte Carlo

Many applications in computational biology, machine learning, theoretical computer science, etc. require sampling from complex dis<u>tributions</u>, which are difficult to write down in closed form, and difficult to directly sample from.

A very common approach is to design a Markov chain whose stationary distribution π is equal to the distribution of interest.

By running this Markov chain for at least $\tau(\epsilon)$ steps (burn-in time), one can draw a sample which is nearly from the distribution of interest.

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By running this Markov chain for at least $\tau(\epsilon)$ steps (burn-in time), one can draw a sample which is nearly from the distribution of interest.

Note: A major focus is on designing and analyzing Markov chains where $\tau(\epsilon)$ is small. For today, we'll just focus on getting the stationary distribution right, and mostly ignore runtime.

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Very non-obvious how to sample from this distribution. Exactly counting the number of independent sets, which is closely related to sampling, is #P-hard.







Design a Markov chain X_0, X_1, \ldots whose states are exactly the independent sets. E.g., let X_{t+1} be chosen uniformly at random from

 $\mathcal{N}(X_t) = \{Y : \text{independent, set formed by adding/removing a node from } X_t\}.$



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Unfortunately, the stationary distribution of this chain may not be uniform. It places higher probability on independent sets with lots of neighboring independent sets.

Define a Markov chain X_0, X_1, \ldots over independent sets with transition function:

- Pick a random vertex v.
- $\cdot \text{ If } v \in X_t \text{, set } X_{t+1} = X_t \setminus \{v\}.$
- If $v \notin X_t$ and $X_t \cup \{v\}$ is independent, set $X_{t+1} = X_t \cup \{v\}$.
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For any two independent sets i, j, what is $P_{i,i}$?

$$j \notin N(i)$$
 $P_{ij} = 0$ c
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Thus, the Markov chain is symmetric, so by our claim from two classes ago, the stationary distribution is uniform.



Known as the 'hard-core model' in statistical physics.



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• Assume access to some symmetric transition function with transition probability matrix $Q \in [0, 1]^{m \times m}$.

A very generic way of designing a Markov chain over state space [m] with stationary distribution $\pi \in [0, 1]^m$.

- Assume the ability to efficiently compute a density $p(X) \propto \pi(X)$.
- Assume access to some symmetric transition function with transition probability matrix $Q \in [0, 1]^{m \times m}$.
- At step *t*, generate a 'candidate' state X_{t+1} from X_t according to *Q*.
- With probability min $\left(1, \frac{p(X_{t+1})}{p(X_t)}\right)$, 'accept' the candidate. Else 'reject' the candidate, setting $X_{t+1} = X_t$.

Metropolis-Hastings Intuition



$$[pP](i) = \underbrace{\sum_{j} p(j) \cdot Q_{j,i} \cdot \min\left(1, \frac{p(i)}{p(j)}\right)}_{aceptances} + \underbrace{p(i) \cdot \sum_{j} Q_{i,j}\left(1 - \min\left(1, \frac{p(j)}{p(i)}\right)\right)}_{rejections}$$

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$$= \underbrace{\sum_{j} Q_{i,j} \cdot \min\left(p(j), p(i)\right) + p(i) \cdot \sum_{j} Q_{i,j} - \sum_{j} Q_{i,j} \cdot \min\left(p(i), p(j)\right)}_{j}$$

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$$= p(i) \cdot \underbrace{\sum_{j} Q_{i,j}}_{j} = p(i).$$

Want to sample an independent set X with probability $\pi(X) = \frac{\lambda^{|X|}}{\sum_{Y \text{ independent }} \lambda^{|Y|}}.$

- Let $p(X) = \lambda^{|X|}$ and let the transition function Q be given by:
 - Pick a random vertex v.
 - If $v \in X_t$, set $X_{t+1} = X_t \setminus \{v\}$
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- Need to accept the transition with probability min $\left(1, \frac{\rho(X_{t+1})}{\rho(X_t)}\right)$.

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 - Pick a random vertex v.
 - If $v \in X_t$, set $X_{t+1} = X_t \setminus \{v\}$ with probability min(1, 1/ λ).
 - If $v \notin X_t$ and $X_t \cup \{v\}$ is independent, set $X_{t+1} = X_t \cup \{v\}$.
 - Else set $X_{t+1} = X_t$ with probability min(1, λ).
- Need to accept the transition with probability min $\left(1, \frac{p(X_{t+1})}{p(X_t)}\right)$.

Want to sample an independent set X with probability $\pi(X) = \frac{\lambda^{|X|}}{\sum_{Y \text{ independent }} \lambda^{|Y|}}.$

- Let $p(X) = \lambda^{|X|}$ and let the transition function Q be given by:
 - Pick a random vertex v.
 - If $v \in X_t$, set $X_{t+1} = X_t \setminus \{v\}$ with probability min(1, 1/ λ).
 - If $v \notin X_t$ and $X_t \cup \{v\}$ is independent, set $X_{t+1} = X_t \cup \{v\}$.
 - Else set $X_{t+1} = X_t$ with probability min(1, λ).
- Need to accept the transition with probability min $\left(1, \frac{p(X_{t+1})}{p(X_t)}\right)$.

The key challenge then becomes to analyze the mixing time.

For the related Glauber dynamics, Luby and Vigoda showed that for graphs with maximum degree Δ , when $\lambda < \frac{2}{\Delta-2}$, the mixing time is $O(n \log n)$. But when $\lambda > \frac{c}{\Delta}$ for large enough constant *c*, it is NP-hard to approximately sample from the hard-core model.