# 15-780: Grad AI Lecture 20: Monte Carlo methods, Bayesian learning 

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## Admin

- Reminder: midterm March 29
- Tuomas's review session tomorrow, mine yesterday

- Reminder: project milestone reports due March 3I


## Review: factor graphs

- Undirected, bipartite graph
- one set of nodes represents variables
- other set represents factors in probability distribution-tables of nonnegative numbers
- need to compute normalizer in order to do anything useful
- Can convert back and forth to Bayes nets
- Hard v. soft constraints


## Review: factor graphs

- Graphical test for independence
- different results from Bayes net, even if we are representing the same distribution
- Inference by dynamic programming
- instantiate evidence, eliminate nuisance nodes, normalize, answer query
- elimination order matters
- treewidth
- Relation to logic


## Review: HMMs, DBNs

- Inference over time
- same graphical template repeated once for each time step-conceptually infinite
- Inference: forwardbackward algorithm (special case of belief propagation)



## Review: numerical integration

- Integrate a difficult function over a highdimensional volume
- narrow, tall peaks contribute most of the integral—difficult search problem
- Central problem for approximate inference
- e.g., computing normalizing constant in a factor graph


## Uniform sampling



## Importance sampling



## Variance

- How does this help us control variance?
- Suppose f big $==>$ Q big
- And Q small ==> f small
- Then $h=f / Q$ never gets too big
- Variance of each sample is lower ==> need fewer samples
- A good Q makes a good IS


## Importance sampling, part II

- Suppose

$$
\begin{aligned}
f(x) & =R(x) g(x) \\
\int f(x) d x & =\int R(x) g(x) d x \\
& =\mathbb{E}_{R}[g(x)]
\end{aligned}
$$

## Importance sampling, part II

- Use importance sampling w/ proposal $\mathrm{Q}(\mathrm{X})$ :
- Pick $N$ samples $x_{i}$ from $Q(X)$
- Average $w_{i} g\left(x_{i}\right)$, where $w_{i}=R\left(x_{i}\right) / Q\left(x_{i}\right)$ is importance weight

$$
\begin{aligned}
\mathbb{E}_{Q}(W g(X)) & =\int Q(x) \frac{R(x)}{Q(x)} g(x) \\
& =\int R(x) g(x) d x \\
& =\int f(x) d x
\end{aligned}
$$

## Parallel IS

- Now suppose $R(x)$ is unnormalized (e.g., represented by factor graph)—know only Z R(x)
- Pick $N$ samples $x_{i}$ from proposal $Q(X)$
- If we knew $w_{i}=R\left(x_{i}\right) / Q\left(x_{i}\right)$, could do IS
- Instead, set

$$
\hat{w}_{i}=Z R\left(x_{i}\right) / Q\left(x_{i}\right)
$$

## Parallel IS

$$
\begin{aligned}
\mathbb{E}(\hat{W}) & =\int Q(x) \frac{Z R(x)}{Q(x)} d x \\
& =\int_{Z} Z R(x) d x \\
& =Z
\end{aligned}
$$

- So, $\bar{w}=\frac{1}{N} \sum_{i} \hat{w}_{i}$ is an unbiased estimate of $\mathbf{Z}$


## Parallel IS

- So, $\hat{w}_{i} / \bar{w}$ is an estimate of $\mathrm{w}_{\mathrm{i}}$, computed without knowing Z
- Final estimate:

$$
\int f(x) d x \approx \frac{1}{n} \sum_{i} \frac{\hat{w}_{i}}{\bar{w}} g\left(x_{i}\right)
$$

## Parallel IS is biased


$E(\bar{W})=Z$, but $E(1 / \bar{W}) \neq 1 / Z$ in general
$Q:(X, Y) \sim N(1,1) \quad \theta \sim U(-\pi, \pi)$
$f(x, y, \theta)=Q(x, y, \theta) P(o=0.8 \mid x, y, \theta) / Z$


Posterior $E(X, Y, \theta)=(0.496,0.350,0.084)$


## Integration problem

- Recall: wanted

$$
\int f(x) d x=\int R(x) g(x) d x
$$

- And therefore, wanted good importance distribution $\mathrm{Q}(\mathrm{x})$ (close to R )


## Back to high dimensions

- Picking a good importance distribution is hard in high-D
- Major contributions to integral can be hidden in small areas
- recall, want (R big ==> Q big)
- Would like to search for areas of high $\mathrm{R}(\mathrm{x})$
- But searching could bias our estimates


## Markov-Chain Monte Carlo

- Design a randomized search procedure M over values of $x$, which tends to increase $R(x)$ if it is small
- Run M for a while, take resulting $x$ as a sample
- Importance distribution $\mathrm{Q}(\mathrm{x})$ ?


## Markov-Chain Monte Carlo

- Design a randomized search procedure M over values of $x$, which tends to increase $R(x)$ if it is small
- Run M for a while, take resulting $x$ as a sample
- Importance distribution $\mathrm{Q}(\mathrm{x})$ ?
- $\mathrm{Q}=$ stationary distribution of $\mathrm{M} . .$.


## Stationary distribution

- Run HMM or DBN for a long time; stop at a random point
- Do this again and again
- Resulting samples are from stationary distribution



## Designing a search chain

$$
\int f(x) d x=\int R(x) g(x) d x
$$

- Would like $\mathrm{Q}(\mathrm{x})=\mathrm{R}(\mathrm{x})$
- makes importance weight = I
- Turns out we can get this exactly, using Metropolis-Hastings


## Metropolis-Hastings

- Way of designing chain $w / Q(x)=R(x)$
- Basic strategy: start from arbitrary $x$
- Repeatedly tweak $x$ to get $x^{\prime}$
- If $R\left(x^{\prime}\right) \geq R(x)$, move to $x^{\prime}$
- If $R\left(x^{\prime}\right) \ll R(x)$, stay at $x$
- In intermediate cases, randomize


## Proposal distribution

- Left open: what does "tweak" mean?
- Parameter of MH: Q(x' $\mid x)$
- one-step proposal distribution
- Good proposals explore quickly, but remain in regions of high $R(x)$
- Optimal proposal?


## MH algorithm

- Sample $x^{\prime} \sim \mathrm{Q}\left(\mathrm{x}^{\prime} \mid \mathrm{x}\right)$
- Compute $\mathrm{p}=\frac{R\left(x^{\prime}\right)}{R(x)} \frac{Q\left(x^{\prime} \mid x\right)}{Q\left(x \mid x^{\prime}\right)}$
- With probability $\min (1, \mathrm{p})$, set $\mathrm{x}:=\mathrm{x}$ '
- Repeat for T steps; sample is $\mathrm{X}_{1}, \ldots, \mathrm{X}_{\boldsymbol{T}}$ (will usually contain duplicates)


## MH algorithm

- Sample $x^{\prime} \sim \mathrm{Q}\left(\mathrm{x}^{\prime} \mid \mathrm{x}\right)$
note: we don't need
- Compute $\mathrm{p}=\frac{R\left(x^{\prime}\right)}{R(x)} \frac{Q\left(x^{\prime} \mid x\right)}{Q\left(x \mid x^{\prime}\right)}$
- With probability $\min (1, \mathrm{p})$, set $\mathrm{x}:=\mathrm{x}$ '
- Repeat for $T$ steps; sample is $\mathrm{X}_{1}, \ldots, \mathrm{X}_{\mathrm{T}}$ (will usually contain duplicates)


## MH example



## Acceptance rate

- Moving to new $x$ ' is accepting
- Want acceptance rate (avg $p$ ) to be large, so we don't get big runs of the same $x$
- Want $\mathrm{Q}\left(\mathrm{x}^{\prime} \mid \mathrm{x}\right)$ to move long distances (to explore quickly)
- Tension between Q and P (accept):

$$
\mathrm{p}=\frac{R\left(x^{\prime}\right)}{R(x)} \frac{Q\left(x^{\prime} \mid x\right)}{Q\left(x \mid x^{\prime}\right)}
$$

## Mixing rate, mixing time

- If we pick a good proposal, we will move rapidly around domain of $R(x)$
- After a short time, won't be able to tell where we started-we have reached stationary dist'n
- This is short mixing time = \# steps until we can't tell which starting point we used
- Mixing rate $=1 /($ mixing time)


## MH estimate

- Once we have our samples $x_{1}, x_{2}, \ldots$
- Optional: discard initial "burn-in" range
- allows time to reach stationary dist'n
- Estimated integral:

$$
\frac{1}{N} \sum_{i=1}^{N} g\left(x_{i}\right)
$$

## In example

- $g(x)=x^{2}$
- True $\mathrm{E}(\mathrm{g}(\mathrm{X}))=0.28 \ldots$
- Proposal: $Q\left(x^{\prime} \mid x\right)=N\left(x^{\prime} \mid x, 0.25^{2} I\right)$
- Acceptance rate 55-60\%
- After 1000 samples, minus burn-in of I00:

```
final estimate 0.282361
final estimate 0.271167
final estimate 0.322270
final estimate 0.306541
final estimate 0.308716
```


## Gibbs sampler

- Special case of MH
- Divide $\mathbf{X}$ into blocks of r.v.s $B(I), B(2), \ldots$
- Proposal Q:
- pick a block i uniformly (or round robin, or any other schedule)
- sample $\mathbf{X}_{B(i)} \sim \mathrm{P}\left(\mathbf{X}_{B(i)} \mid \mathbf{X}_{\neg B(i)}\right)$


## Gibbs example



## Gibbs example



## Why is Gibbs useful?

- For Gibbs, $\mathrm{p}=\frac{P\left(x_{i}^{\prime}, x_{\neg i}^{\prime}\right)}{P\left(x_{i}, x_{\neg i}\right)} \frac{P\left(x_{i} \mid x_{\neg i}^{\prime}\right)}{P\left(x_{i}^{\prime} \mid x_{\neg i}\right)}$


## Gibbs derivation

$$
\begin{aligned}
& \frac{P\left(x_{i}^{\prime}, x_{\neg i}^{\prime}\right)}{P\left(x_{i}, x_{\neg i}\right)} \frac{P\left(x_{i} \mid x_{\neg i}^{\prime}\right)}{P\left(x_{i}^{\prime} \mid x_{\neg i}\right)} \\
= & \frac{P\left(x_{i}^{\prime}, x_{\neg i}\right)}{P\left(x_{i}, x_{\neg i}\right)} \frac{P\left(x_{i} \mid x_{\neg i}\right)}{P\left(x_{i}^{\prime} \mid x_{\neg i}\right)} \\
= & \frac{P\left(x_{i}^{\prime}, x_{\neg i}\right)}{P\left(x_{i}, x_{\neg i}\right)} \frac{P\left(x_{i}, x_{\neg i}\right) / P\left(x_{\neg i}\right)}{P\left(x_{i}^{\prime}, x_{\neg i}\right) / P\left(x_{\neg i}\right)} \\
= & 1
\end{aligned}
$$

## Gibbs in practice

- Proof of $p=1$ means Gibbs is often easy to implement
- Often works well
- if we choose good blocks (but there may be no good blocking!)
- Fancier version: adaptive blocks, based on current $\mathbf{X}$


## Gibbs failure example



## Sequential sampling



- In an HMM or DBN, to sample $P\left(\mathbf{X}_{T}\right)$, start from $\mathbf{X}_{1}$ and sample forward step by step - $\mathbf{X}_{\mathrm{t}+1} \sim P\left(\mathbf{X}_{\mathrm{t}+1} \mid \mathbf{X}_{\mathrm{t}}\right)$
- $\mathrm{P}\left(\mathbf{X}_{1: T}\right)=\mathrm{P}\left(\mathbf{X}_{1}\right) \mathrm{P}\left(\mathbf{X}_{2} \mid \mathbf{X}_{1}\right) \mathrm{P}\left(\mathbf{X}_{3} \mid \mathbf{X}_{2}\right) \ldots$


## Particle filter

- Can sample $\mathbf{X}_{t+1} \sim P\left(\mathbf{X}_{t+1} \mid \mathbf{X}_{\mathrm{t}}\right)$ using any algorithm from above
- If we use parallel importance sampling to get N samples at once from each $\mathrm{P}\left(\mathbf{X}_{\mathrm{t}}\right)$, we get a particle filter
- also need one more trick: resampling
- Write $\mathbf{x}_{\mathrm{t}, \mathrm{i}}(\mathrm{i}=\mathrm{I} \ldots \mathrm{N})$ for sample at time t


## Particle filter

- Want one sample from each of $P\left(\mathbf{X}_{t+1} \mid \mathbf{X}_{t, i}\right)$
- Have only Z P( $\left.\mathbf{X}_{\mathrm{t}+1} \mid \mathbf{x}_{\mathrm{t}, \mathrm{i}}\right)$
- For each i, pick $\mathbf{x}_{\mathrm{t}+1, \mathrm{i}}$ from proposal $\mathrm{Q}(\mathrm{x})$
- Compute unnormalized importance weight

$$
\hat{w}_{i}=Z P\left(\mathbf{x}_{t+1, i} \mid \mathbf{x}_{t, i}\right) / Q\left(\mathbf{x}_{t+1, i}\right)
$$

## Particle filter

- Normalize weights:

$$
\bar{w}=\frac{1}{N} \sum_{i} \hat{w}_{i} \quad w_{i}=\hat{w}_{i} / \bar{w}
$$

- Now, $\left(w_{i}, \mathbf{X}_{t+1, i}\right)$ is an approximate weighted sample from $\mathrm{P}\left(\mathbf{X}_{\mathrm{t}+1}\right)$
- What will happen if we do this for $T=I, 2, \ldots$ ?


## Resampling

- To get an unweighted sample, resample
- Sample $N$ times (with replacement) from $\boldsymbol{X}_{t+1, i}$ with probabilities $w_{i} / N$
- alternately: deterministically take floor( $w_{i}$ ) copies of $\boldsymbol{x}_{t+1, i}$ and sample only from fractional part [ $w_{i}$ - floor $\left(w_{i}\right)$ ]
- Each $\boldsymbol{X}_{t+1, i}$ appears $w_{i}$ times on average, so we're still a sample from $P\left(\boldsymbol{X}_{t+1}\right)$


## Particle filter example



## Learning

- Basic learning problem: given some experience, find a new or improved model
- Experience: a sample $x_{1}, \ldots, x_{N}$
- Model: want to predict $x_{N+1}, \ldots$


## Example

- Experience = range sensor readings \& odometry from robot
- Model = map of the world



## Example

- The "botanist learning problem"
- Experience = physical measurements of surveyed specimens \& expert judgements of their true species
- Model = factor graph relating species to measurements


## Sample data

| sepal <br> length | sepal <br> width | petal <br> length | petal <br> width | species |
| :---: | :---: | :---: | :---: | :---: |
| 5.1 | 3.5 | 1.4 | 0.2 | Iris setosa |
| 5.6 | 3.0 | 4.5 | 1.5 | Iris <br> versicolor |
| 4.9 | 3.0 | 1.4 | 0.2 | Iris setosa |
| 6.4 | 2.8 | 5.6 | 2.1 | Iris <br> virginica |
| 5.8 | 2.7 | 4.1 | 1.0 | Iris <br> versicolor |

Factor graph


- One of many possible factor graphs
- Values of $\Phi s$ not shown, but part of model

Factor graph



## Factor graph



## In general

- For our purposes, a model $M$ is exactly a distribution $P(\boldsymbol{X} \mid M)$ over possible samples
- When is $M$ better than $M$ ? When $P(X \mid M)$ is more accurate than $P\left(\boldsymbol{X} \mid M^{\prime}\right)$.
- Bayes rule encodes this: from prior $P(M)$ and evidence $\boldsymbol{X}$, compute posterior $P(M \mid \boldsymbol{X})$
- $P(M \mid \boldsymbol{X})=P(\boldsymbol{X} \mid M) P(M) / P(\boldsymbol{X})$
- better predictions (higher $P(\boldsymbol{X} \mid M)$ ) yield higher posterior


## Conditional model

- Split variables into (X, Y)
- Suppose we always observe $\boldsymbol{X}$
- Two ways $P(\boldsymbol{X}, \boldsymbol{Y})$ and $P^{\prime}(\boldsymbol{X}, \boldsymbol{Y})$ can differ:
- $P(\boldsymbol{X}) \neq P^{\prime}(\boldsymbol{X})$, and/or
- $P(\boldsymbol{Y} \mid \boldsymbol{X}) \neq P^{\prime}(\boldsymbol{Y} \mid \boldsymbol{X})$
- First way doesn't matter for decisions
- Conditional model: only specifies $P(\boldsymbol{Y} \mid \boldsymbol{X}, M)$


## Conditional model example

- Experience $=$ samples of $(\mathbf{X}, \mathbf{Y})$
- $\boldsymbol{X}=$ features of object
- $\boldsymbol{Y}=$ whether object is a "framling"
- Model = rule for deciding whether a new object is a framling


## Sample data \& possible model

| tall | pointy | blue | framling |
| :---: | :---: | :---: | :---: |
| T | T | F | T |
| T | F | F | T |
| F | T | F | F |
| T | T | T | F |
| T | F | F | T |

$$
\mathrm{H}=\text { tall } \wedge \neg \text { blue }
$$

## Hypothesis space

- Hypothesis space $\mathscr{K}=$ set of models we are willing to consider
- for philosophical or computational reasons
- E.g., all factor graphs of a given structure
- Or, all conjunctions of up to two literals
- Prior is a distribution over $\mathscr{K}$


## A simple learning algorithm

## = Bayes Rule

- Conditional learning: samples ( $\mathbf{x}_{i}, y_{i}$ )
- Let $\mathscr{K}$ be a set of propositional formulae
- $\mathscr{K}=\left\{H_{1}, H_{2}, \ldots\right\}$
- $H$ is consistent if $H\left(\mathbf{x}_{i}\right)=y_{i}$ for all $i$
- Version space $V=\{$ all consistent $H\} \subseteq \mathscr{K}$
- Version space algorithm: predict $y=$ majority vote of $H(\mathbf{x})$ over all $H \in V$


## Framlings

| tall | pointy | blue | framling |
| :---: | :---: | :---: | :---: |
| T | T | F | T |
| T | F | F | T |
| F | T | F | F |
| T | T | T | F |
| T | F | F | T |

- $\mathscr{K}=\{$ conjunctions of up to 2 literals $\}=\{\mathrm{T}, \mathrm{F}$, tall, pointy, blue, $\neg$ tall, $\neg$ pointy, $\neg$ blue, tall $\wedge$ pointy, tall $\wedge$ blue, pointy $\wedge$ blue, $\neg$ tall $\wedge$ pointy,.. \}

Framlings

| tall | pointy | blue | framling |
| :---: | :---: | :---: | :---: |
| T | T | F | T |
| T | F | F | T |
| F | T | F | F |
| T | T | T | F |
| T | F | F | T |

$$
\begin{aligned}
& x_{1}: T_{\text {, tall, pointy, }} \text { ablue, tall ipointy, tall } 1 \text { a blve } \\
& \text { ponity }{ }_{x} \text { a blue }
\end{aligned}
$$

$$
\begin{aligned}
& x_{j} \text { itall, talln } \operatorname{slin}^{x}
\end{aligned}
$$

## Analysis

- Mistake = make wrong prediction
- If some $\mathrm{H} \in \mathscr{K}$ is always right, eventually we'll eliminate all competitors, and make no more mistakes
- If no $\mathrm{H} \in \mathscr{K}$ is always right, eventually V will become empty
- e.g., if label noise or feature noise


## Analysis

## - Suppose $|\mathscr{K}|=\mathrm{N}$

- How many mistakes could we make?

$$
\Gamma \log _{2} N 7
$$

## Analysis

- Suppose $|\mathscr{T}|=\mathrm{N}$
- How many mistakes could we make?
- Since we predict w/ majority ofV, after any mistake, we eliminate half (or more) of $V$
- Can't do that more than $\log _{2}(\mathrm{~N})$ times


## Discussion

- In example, $N=20, \log _{2}(N)=4.32$
- Made only 2 mistakes
- Mistake bound: limits wrong decisions, as desired
- But, required strong assumptions (no noise, true H contained in $\mathscr{K}$ )
- Could be very slow!

