Sampling

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February 23, 2015

Content

Motivation

Sampling

Monte Carlo methods Markov chain Monte Carlo methods

MCMC parsing

Conclusions

A PCFG $G = \langle \Sigma, N, S, R, p \rangle$

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from here let's assume CNF

Let us define the parameters $\boldsymbol{\theta} \in [0,1]^{|R|}$ where $p(r) = \theta_r$ where $r = A \to \alpha$ is a rule in R

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Expectations: Inside-Outside dynamic program $O(|V|^3|\mathbf{w}|^3)$

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If we could solve (1)

then
$$\hat{\Phi} \equiv \frac{1}{N} \sum_{i=1}^N \phi(x^{(i)})$$

Robert and Casella [2004]

Monte Carlo estimates

Accuracy of an MC estimate is independent of dimensionality

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However, it is **very hard** to sample from high dimensional spaces!

Sampling from chart

Given a string \mathbf{w} , assume we can build the chart $\mathcal{T}(\mathbf{w})$

• $\langle i, A, j \rangle$ where $A \in N$ and $0 \le i < j \le |\mathbf{w}|$ represents a chart cell

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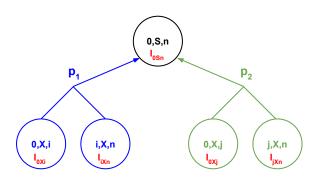
then,

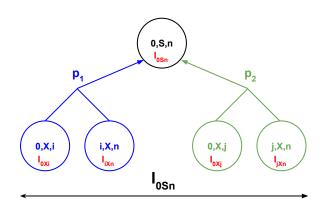
expectations trivial Inside-Outside run sampling trivial random tree traversal from start symbol

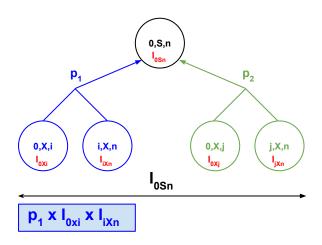
Top-down sampling illustration

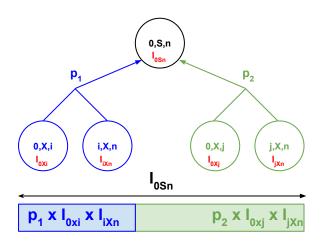


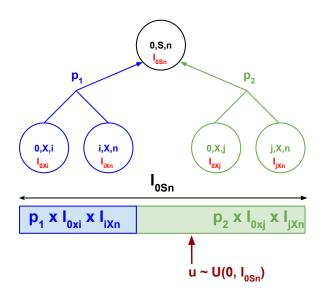
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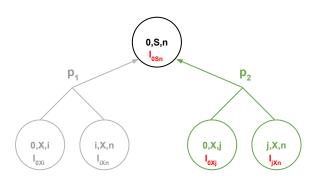


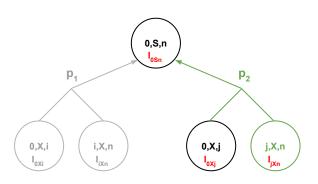


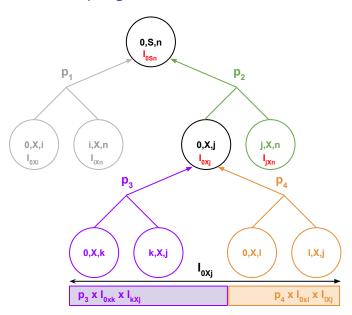


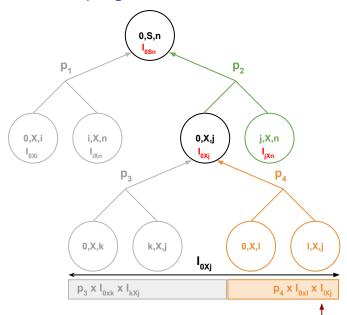












Sampling from chart

Given a string \mathbf{w} , assume we can build the chart $\mathcal{T}(\mathbf{w})$

We care about the cases in which we cannot instantiate the chart!

Why is it hard to sample from high dimensional spaces?

Let's rewrite the density

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In parsing

it is the inside at the root of the chart but we cannot afford building the chart!

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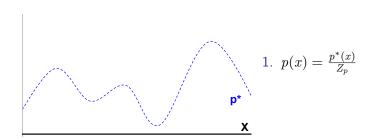
We could sample uniformly directly from the support ${\mathcal X}$

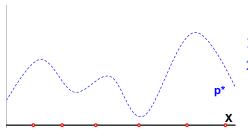
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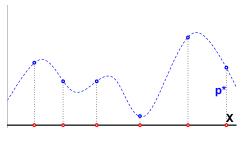
We could sample uniformly directly from the support \mathcal{X} approximating Z_p by how much of it we have seen

$$Z_N = \sum_{i=1}^{N} p^*(x^{(i)})$$



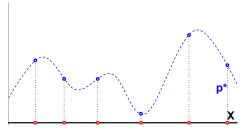


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• the typical set T $|T| \approx 2^{H(x)}$

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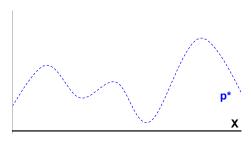
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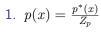
Suppose, 10^3 bits think of it as rules in a chart for $|\mathbf{w}| = 10$

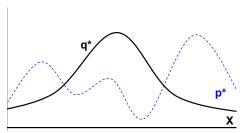
 $ho 2^{500} pprox 10^{150}$ trials square of the number of particles in the universe [MacKay, 1998]

Lessons

- 1. assessing \mathbb{Z}_p in high dimensional spaces is hard
- 2. sampling is hard even when $p^*(x)$ is easy to evaluate (and direct access to Z_p is not required)



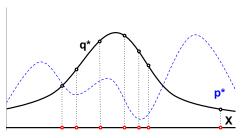




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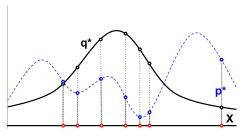


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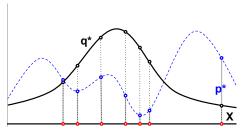


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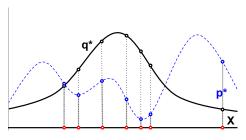
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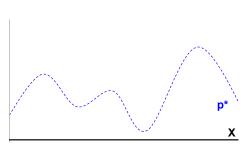
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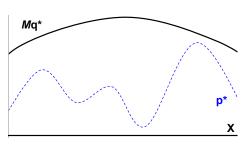
$$\hat{w}(x^{(i)}) = \frac{w^*(x^{(i)})}{\sum_{j=1}^{N} w^*(x^{(j)})}$$

Introduces an instrumental distribution q(x)

- a better guess than sampling uniformly from the state space
- ightharpoonup q(x) is such that sampling from it is trivial
- the variance of the estimate becomes a q(x)

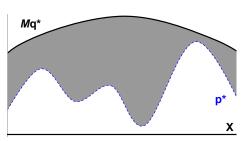


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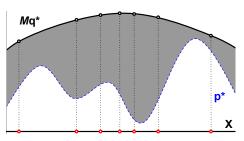
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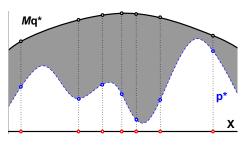
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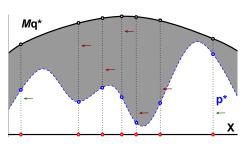


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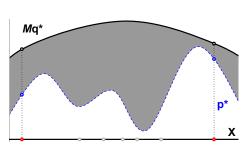
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Accepted x's make an exact sample from p(x)

$$\hat{\Phi} = \sum_{i=1}^{N} \phi(x^{(i)})$$

Introduces an upperbound $Mq^*(x) \ge p^*(x)$

- 1. sample (x,u) uniformly distributed over the (d+1)-dimensional surface under $Mq^*(x)$
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Problem

- low acceptance rate
- lacktriangle in high dimensional spaces, M is typically huge the ratio $rac{Z_p}{MZ_q}
 ightarrow 0$

Consider the integration of a parser and a $2\mathrm{nd}$ order HMM tagger

$$p(\mathbf{t}) = p_G(\mathbf{t}) p_{H_2}(h(\mathbf{t}))$$

where $h(\mathbf{t})$ is the sequence of tags

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the parser alone makes the instrumental distribution

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Rejection sampling

replace p_{H_2} by a lower-order upperbound (e.g. 0-order HMM)

$$q(\mathbf{t}) = p_G(\mathbf{t}) q_{H_0}(h(\mathbf{t}))$$

Markov chain Monte Carlo

A Markov chain that leaves the desired distribution invariant

- unlike MC, samples are not independent
- ▶ in the limit of an infinite chain, the state of the chain converges to the target distribution
- we typically discard the beginning of the chain (i < k) to reduce dependency on starting conditions

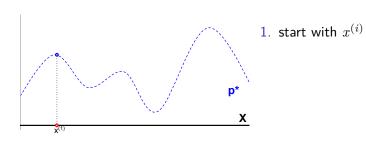
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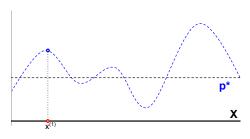
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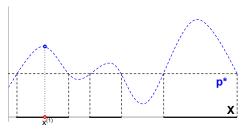
Samples and expectation

- 1. $\{x^{(i)}\}_{i=k}^{N}$
- 2. $\hat{\Phi} = \sum_{i=k}^{N} \phi(x^{(i)})$

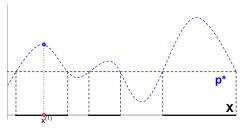




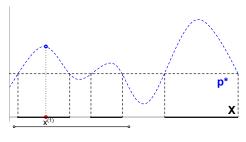
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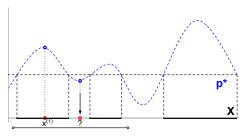
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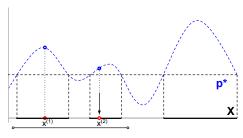
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- ightharpoonup draw x' uniformly from I
- lacktriangleright make $x^{(i+1)}=x'$ if $x'\in S$, that is, $p^*(x')>y$

An attempt to get a "black box" sampler

- form of auxiliary variable sampling
- no need for proxy distributions
- requires assessing p for a given sample and for the boundaries of an interval I
- ightharpoonup finding I can be hard

Task sample from the joint $p(\mathbf{x} = x_1, \dots, x_n)$

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Method

repeatedly sample from the conditional for each x_{k}

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- can be done when we know how to sample from all the required conditional distributions
- running the sampler for a sufficiently long time produces a samples of values for x from close to the target distribution

MCMC pros and cons

Cons

- 1. slow mixture (particularly Gibbs)
- 2. hard to diagnose convergence

Pros

- 1. enable inference when $p(\boldsymbol{x})$ is just too complex for dynamic programming
- estimates can always be improved by increasing the number of samples

In synchronous parsing we recognise pairs of strings (\mathbf{x},\mathbf{y})

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MLE via EM requires Inside-Outside $O(|\mathbf{x}|^3|\mathbf{y}|^3)$ prohibitive!

We introduce an auxiliary variable per chart cell

chart

$$S = \{ \langle A, i, j, k, l \rangle : 0 \le i < j \le |\mathbf{x}|, 0 \le k < l \le |\mathbf{y}|, A \in V \}$$

slice variables

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The slice variables act as cutoff on the probabilities of the rules considered in each cell

▶ rule applications r_s with $\theta_{r_s} \leq u_s$ are pruned from the dynamic program

Sampling $p(\mathbf{u}|\mathbf{t})$

 $ightharpoonup u_s$ are conditionally independent

$$u_s \sim p(u_s|\mathbf{t}) = egin{cases} U(u_s;0, heta_{r_s}) & ext{if } r_s \in \mathbf{t} \\ eta(u_s;a,1) & ext{otherwise} \end{cases}$$

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Slice sampling for synchronous parsing

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The hyperparameter a controls the degree of pruning Blunsom and Cohn [2010]

Bayesian inference

Bayesian MAP inference

$$p(\boldsymbol{\theta}|\mathcal{W}) \propto p_G(\mathcal{W}|\boldsymbol{\theta})p(\boldsymbol{\theta})$$

- W data (set of strings)
- $p_G(\mathcal{W}|\boldsymbol{\theta})$ likelihood of data given model $\boldsymbol{\theta}$
- $p(\theta)$ prior (if uniform we get MLE)

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$$\boldsymbol{\theta} \sim p_D(\boldsymbol{\theta}; \boldsymbol{\alpha})$$

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Posterior is also a Dirichlet

▶ $p_D(\theta|\mathcal{T}; \alpha) = p_D(\theta|\mathbf{f}(\mathcal{T}) + \alpha)$ "updates the prior conditioning on evidence"

Let's rewrite the posterior in terms of a joint distribution

$$p(\boldsymbol{\theta}|\mathcal{W}; \boldsymbol{\alpha}) = \sum_{\mathcal{T} \in G(\mathcal{W})} p(\mathcal{T}, \boldsymbol{\theta}|\mathcal{W}; \alpha)$$

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there exists efficient techniques to sample from a Dirichlet Johnson et al. [2007]

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- approximate distributions in general



References I

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