# Sampling

#### for unsupervised language learning

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#### 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

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

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#### **Tasks**

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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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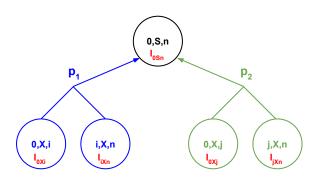
then,

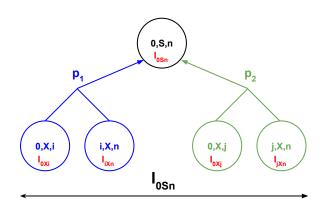
expectations trivial Inside-Outside run sampling trivial random tree traversal from start symbol

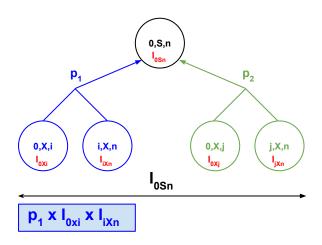
# Top-down sampling illustration

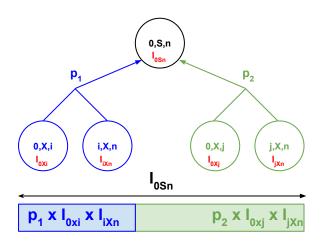


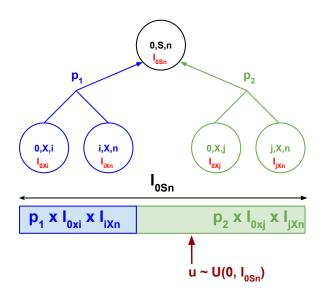
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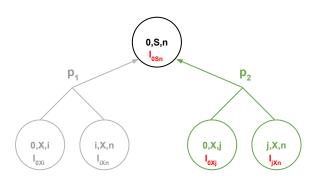


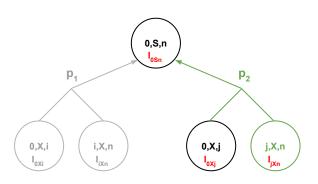


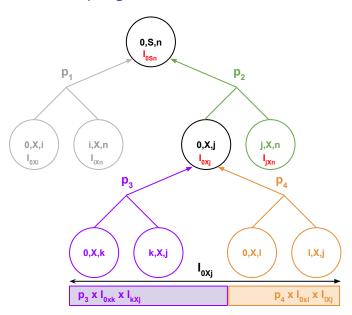


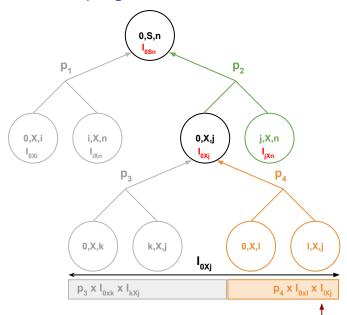












#### Sampling from chart

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

$$p(x) = \frac{p^*(x)}{Z_p} = \frac{p^*(x)}{\int_{\mathcal{X}} p^*(x) dx}$$

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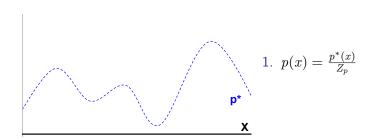
We could sample uniformly directly from the support  ${\mathcal X}$ 

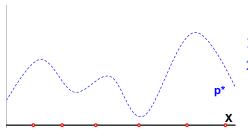
#### Estimating expectations

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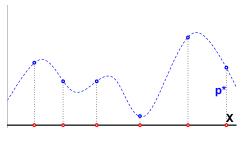
We could sample uniformly directly from the support  $\mathcal{X}$  approximating  $Z_p$  by how much of it we have seen

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



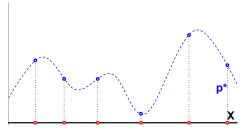


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$$\hat{\Phi} = \sum_{i=1}^{N} \phi(x^{(i)}) \hat{p}(x^{(i)})$$

Probability mass is often concentrated in a small region

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

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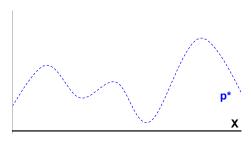
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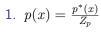
Suppose,  $10^3$  bits think of it as rules in a chart for  $|\mathbf{w}| = 10$ 

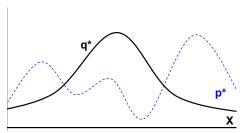
 $ho 2^{500} pprox 10^{150}$  trials square of the number of particles in the universe [MacKay, 1998]

#### 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)

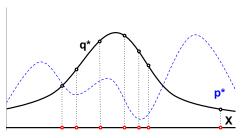






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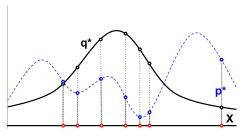
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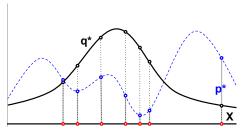
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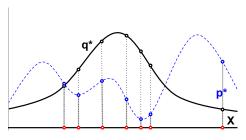


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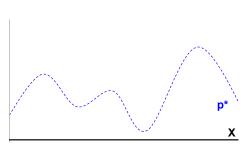
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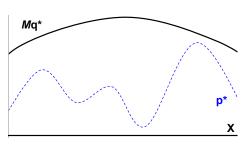
$$\hat{w}(x^{(i)}) = \frac{w^*(x^{(i)})}{\sum_{j=1}^{N} w^*(x^{(j)})}$$

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)

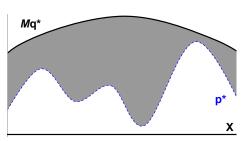


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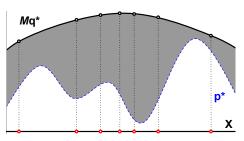
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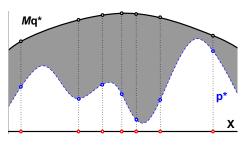
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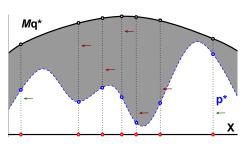


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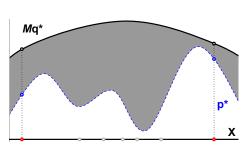
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Accepted x's make an exact sample from p(x)

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

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

- 1. sample (x,u) uniformly distributed over the (d+1)-dimensional surface under  $Mq^*(x)$
- 2. retain only points uniformly distributed under  $p^*(x)$

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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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### 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

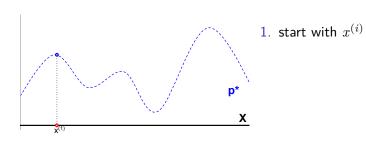
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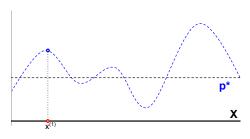
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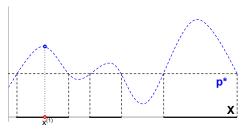
#### Samples and expectation

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

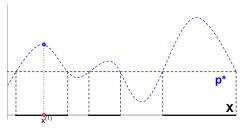




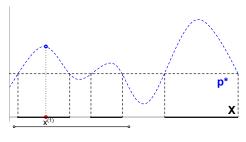
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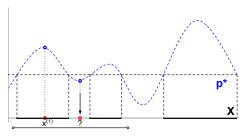
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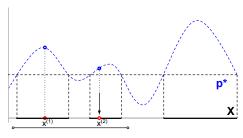
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- ightharpoonup draw x' uniformly from I
- lacktriangleright make  $x^{(i+1)}=x'$  if  $x'\in S$ , that is,  $p^*(x')>y$

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

#### Method

repeatedly sample from the conditional for each  $\mathit{x}_{\mathit{k}}$ 

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Conditioning greatly reduces dimensionality

- 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  $(\mathbf{x},\mathbf{y})$ 

In synchronous parsing we recognise pairs of strings (x, y)Consider a binary ITG (a special case of SCFG)

- ▶ start symbol *S*
- ightharpoonup and a single nonterminal X which can be rewritten
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We introduce an auxiliary variable per chart cell

chart

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

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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

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$$\boldsymbol{\theta}_A \sim p_D(\boldsymbol{\theta}_A|\mathbf{f}_A(\mathcal{T}) + \boldsymbol{\alpha})$$

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

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One can sample in order to

- compute expectations (e.g. when performing MLE)
- marginalise latent variables

Sampling provides a powerful inference framework

- when the distribution is too complex to be represented
- also for Bayesian inference where complex posteriors do not factorise conveniently as the likelihood typically does

People tend to associate MC/MCMC with Bayesian inference

they are orthogonal

One can sample in order to

- compute expectations (e.g. when performing MLE)
- marginalise latent variables
- approximate distributions in general



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