lecture 12: bayesian inference and monte carlo methods

STAT 545: Intro. to Computational Statistics

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Bayesian inference

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What if we believe *θ* is close to 0, is sparse, or is smooth? Encode this with a 'prior' probability *p*(*θ*).

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Point estimate discards information about uncertainty in *θ*

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Allows us to maintain and propagate uncertainty.

E.g. consider the likelihood $p(X|\theta) = N(X|\theta, 1)$

- What is a good prior over *θ*?
- What is a convenient prior over *θ*?
- The posterior distribution $p(\theta|X) \propto p(X|\theta)p(\theta)$ summarizes all new information about *θ* provided by the data
- In practice, these distributions are unwieldy.
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- In practice, these distributions are unwieldy.
- Need approximations.
- An exception: 'Conjugate priors' for exponential family distributions.

Let observations come from an exponential-family:

$$
p(x|\theta) = \frac{1}{Z(\theta)} h(x) \exp(\theta^\top \phi(x))
$$

= $h(x) \exp(\theta^\top \phi(x) - \zeta(\theta))$ with $\zeta(\theta) = \log(Z(\theta))$

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p(\theta|X) \propto \left(\prod_{i=1}^N h(x_i) \exp(\theta^\top \phi(x_i) - \zeta(\theta))\right) \eta(\theta) \exp(\theta^\top a - \zeta(\theta)b)
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$$

$$
\propto \eta(\theta) \exp\left(\theta^{\top} \left(a + \sum_{i=1}^{N} \phi(x_i)\right) - \zeta(\theta)(b+N)\right)
$$

Prior over *θ*: exp. fam. distribution with parameters (*a, b*). Posterior: same family with parameters $(a + \sum_{i=1}^{N} \phi(x_i), b + N)$. Rare instance where analytical expressions for posterior exists. In most cases a simple prior quickly leads to a complicated posterior, requiring Monte Carlo methods.

Prior over *θ*: exp. fam. distribution with parameters (*a, b*). Posterior: same family with parameters $(a + \sum_{i=1}^{N} \phi(x_i), b + N)$. Rare instance where analytical expressions for posterior exists. In most cases a simple prior quickly leads to a complicated posterior, requiring Monte Carlo methods. Note the conjugate prior is an entire family of distributions.

- Actual distribution is chosen by setting the parameters (*a, b*) (*a* has the same dimension as *ϕ*, *b* is a scalar)
- These might be set by e.g. talking to a domain expert.

Let $x_i \in \{0, 1\}$ indicate if a new drug works for subject *i*. The unknown probability of success is π : $x \sim \text{Bern}(\pi)$.

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p(x|\pi) = \pi^{1(x=1)}(1-\pi)^{1(x=0)}
$$

= exp (1(x = 1) log(\pi) + (1 - 1(x = 1)) log(1 - \pi))
= (1 - \pi) exp (1(x = 1) log \frac{\pi}{1 - \pi})
= $\frac{1}{1 + exp(\theta)}$ exp (\phi(x)\theta)

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 $\theta = \log \frac{\pi}{1-\pi}, \phi(x) = \mathbb{1}(x = 1), h(x) = 1, Z(\theta) = (1 + \exp(\theta)).$ Defining $\zeta(\theta) = \log Z(\theta)$ as in the previous slide,

$$
p(x|\theta) = \exp(\phi(x)\theta - \zeta(\theta))
$$

When *θ* = log *^π* 1*−π* is unknown, a Bayesian places a prior on it.

As before, define an exp. fam. prior with parameters \vec{a} :

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Then given data $X = (x_1, \ldots, x_N)$,

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p(\theta|\vec{a},X) \propto p(\theta,X|\vec{a})
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Thus, the posterior is in the same family as the prior, but with updated parameters $\left(a_1 + \sum_{i=1}^N \mathbb{1}(x_i = 1), a_2 - N\right)$.

Looking at the prior more carefully, we see:

$$
p(\theta|\vec{a}) \propto \exp(a_1\theta + a_2\zeta(\theta))
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\propto \exp\left(a_1\log\frac{\pi}{1-\pi} + a_2\log(1-\pi)\right)
$$

$$
\propto \pi^{a_1}(1-\pi)^{(a_2-a_1)}
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\n
$$
= \pi^{b_1-1}(1-\pi)^{(b_2-1)}
$$

This is just the Beta(b_1 , b_2) distribution, and you can check that the posterior is Beta $\left(b_1 + \sum_{i=1}^{N} 1\!\!1 (x_i = 1), b_2 + \sum_{i=1}^{N} 1\!\!1 (x_i = 0)\right)$.

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 b_1 and b_2 are sometimes called pseudo-observations, and capture our prior beliefs: before seeing any *x*'s our prior is as if we saw b_1 successes and b_2 failures. After seeing data, we factor actual observations into the pseudo-observations. $\frac{8}{299}$

monte carlo methods

What about the situation when the posterior $p(\theta|X)$ is no longer simple/available in closed form?

What information about *p*(*θ|X*) do we really need?

Typically, expectations of different functions *g*:

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What is *g* for to calculate 1) mean, 2) variance, 3) $p(\theta > 10|X)$?

Monte Carlo integration

Let us forget the posterior distribution *p*(*θ|X*), and consider some general probability distribution *p*(*x*). We want

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Monte Carlo approximation:

- Obtain points by sampling from *p*(*x*): *xⁱ ∼ p*
- Approximate integration with summation

$$
\hat{\mu} \approx \frac{1}{N} \sum_{i=1}^{N} g(x_i)
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If *xⁱ ∼ p*,

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\mathbb{E}_{p}[\hat{\mu}] = \frac{1}{N} \sum_{i=1}^{N} \mathbb{E}_{p}[g] = \mu
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E*p*[*g*] = *µ* Unbiased estimate

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 $\textsf{Var}_p[\hat{\mu}]=\frac{1}{N}$ $Var_p[g]$, Error = StdDev $\propto N^{-1/2}$

$$
\frac{1}{N}\sum_{i=1}^N f \to \mathbb{E}_p(g) = \mu \quad \text{ as } N \to \infty
$$

f → E*p*(*g*) = *µ* as *N → ∞* Consistent estimate (LLN)

Is this a good idea?

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error *∝ N −*4*/d*

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Independent of dimensionality!

- If unbiasedness is important to you.
- Very simple.
- Very modular: easily incorporated into more complex models (Gibbs sampling)

An aside: Monte Carlo should be your method of last resort!

Don't hesitate using numerical integration

• Numerical integration can be much faster and more accurate

Contrast

- *>* integrate(function(x) x *∗* exp(*−*x)*,* lower = 0*,* upper = Inf) with
- *>* mean(rexp(1000))

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- Easy to generate uniform RVs on a deterministic computer?

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- Careful with batch/parallel processing.

R has a bunch of random number generators. rnorm, rgamma, rbinom, rexp, rpoiss etc. What if we want samples from some other distribution?

Let *X* have pdf $p(x)$, and cdf $F(x) = P(X \le x) = \int_{-\infty}^{x} p(u) \, du$ Let:

> *X ∼ p*(*·*) $U = F(X)$

Inverse transform sampling Let *X* have pdf $p(x)$, and cdf $F(x) = P(X \le x) = \int_{-\infty}^{x} p(u) \, du$ Let:

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Equivalently, sample $U \sim \text{Unif}(0,1)$, and let $X = F^{-1}(U)$ Then *X ∼ p*(*·*) (see wikipedia for proof)

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Equivalently, sample $U \sim \text{Unif}(0,1)$, and let $X = F^{-1}(U)$ Then *X ∼ p*(*·*) (see wikipedia for proof)

E.g. *−* log(*U*) is Exponential(1). Usually hard to compute *F −*1 .

Let $p(x) = \frac{f(x)}{Z}$. Probability of a sample in $[x_0, x_0 + \Delta x] = p(x_0) \Delta x$.

If we sample points uniformly below the curve *Mf*(*x*):

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Let $p(x) = \frac{f(x)}{Z}$. Probability of a sample in $[x_0, x_0 + \Delta x] = p(x_0) \Delta x$.

If *Mf*(*x*) *≤ Nq*(*x*) *∀x* for constant *N* and distribution *q*(*·*) Sample points uniformly under *Nq*(*x*). (sample $x_0 \sim q(\cdot)$, and assign it a uniform height in [0, $Nq(x_0)$]

Let $p(x) = \frac{f(x)}{Z}$. Probability of a sample in $[x_0, x_0 + \Delta x] = p(x_0) \Delta x$.

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Let $p(x) = \frac{f(x)}{Z}$. Probability of a sample in $[x_0, x_0 + \Delta x] = p(x_0) \Delta x$.

Equivalent algorithm: (convince yourself)

- Propose *x [∗] ∼ q*(*·*)
- Accept with probability *Mf*(*x ∗*)*/Nq*(*x ∗*)

Let $p(x) = \frac{f(x)}{Z}$. Probability of a sample in $[x_0, x_0 + \Delta x] = p(x_0) \Delta x$.

We need a bound on *f*(*x*).

A loose bound leads to lots of rejections. Probability of acceptance = $\frac{MZ}{N}$.

A probability density takes the form $p(x) = \frac{f(x)}{Z}$

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However, rejection sampling doesn't need *Z* or *p*(*x*)

Example 1:

$$
p(x) \propto \exp(-x^2/2)|\sin(x)|
$$

Example 2 (truncated normal):

$$
p(x) \propto \exp(-x^2/2)1_{\{x>c\}}
$$

What is *M* for each case? What can we say about efficiency?

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$$
\mathbb{E}_{p}[g] = \int g(x)p(x)dx = \int g(x)\frac{p(x)}{q(x)}q(x)dx = \mathbb{E}_{q}\left[\frac{g(x)p(x)}{q(x)}\right]
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Use Monte Carlo approximation to the latter expectation:

• Draw proposal *x* from $q(\cdot)$ and calculate weight $w(x) = \frac{p(x)}{q(x)}$.

$$
\int g(x)p(x)dx \approx \frac{1}{N}\sum_{s=1}^N w(x_s)g(x_s)
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Since $w(x) = p(x)/q(x) = \frac{f(x)}{Zq(x)}$:

- We don't need a bounding envelope.
- We need normalizn constant *Z* (but see later).

Importance sampling vs simple Monte Carlo

Simple Monte Carlo/MCMC (left) uses sampling approximation Importance sampling (right) weights the samples

When does this make sense? Sometimes it's easier to simulate from *q*(*x*) than *p*(*x*). When does this make sense? Sometimes it's easier to simulate from *q*(*x*) than *p*(*x*).

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To reduce variance. E.g. rare event simulation.

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To reduce variance. E.g. rare event simulation. Let *x ∼* (0*,* 1)

• What is *P*(*X >* 5)?

Let $X = (x_1, ..., x_{100})$ be a hundred dice. What is $p(\sum x_i \ge 550)$?
Importance sampling:

Let $X = (x_1, ..., x_{100})$ be a hundred dice. What is $p(\sum x_i \ge 550)$?

Rejection sampling (from *p*(*x*)) leads to high rejection.

Importance sampling:

Let $X = (x_1, ..., x_{100})$ be a hundred dice. What is $p(\sum x_i \ge 550)$?

Rejection sampling (from *p*(*x*)) leads to high rejection.

A better choice might be to bias the dice.

E.g. *q*(*x_i* = *v*) ∞ *v* (for *v* ∈ {1, ...6})

Define
$$
S_X = \sum X_i
$$

$$
p(S \ge 550) = \sum_{y \in all \text{ configs of 100 dice}} \delta(\sum y \ge 550)p(y)
$$

$$
= \sum_{y \in all \text{ configs of 100 dice}} \frac{p(y)}{q(y)} \delta(\sum y \ge 550)q(y)
$$

For a proposal *X [∗] ∼ q*,

$$
w(X^*) = \frac{p(X^*)}{q(X^*)} = \frac{(1/6)^{100}}{\prod_i q(X_i^*)}
$$

Use approximation $p(S \ge 550) \approx \sum_{j=1}^{N} w(X_j) \delta(\sum x_i^j \ge 550)$

$$
\text{Var}[\mu_{imp}] = \mathbb{E}[\mu_{imp}^2] - \mu^2
$$

=
$$
\mathbb{E}\left[\left(\frac{1}{N}\sum_{i=1}^N w_i g(x_i)\right)^2\right] - \mu^2
$$

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&= \int_{\mathcal{X}} q(x) \left(\frac{p(x)g(x)}{q(x)}\right)^2 dx - \mu^2 \\
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&\geq \left(\int_{\mathcal{X}} q(x) \frac{p(x)g(x)}{q(x)} dx\right)^2 - \mu^2 \\
&= 0\n\end{aligned}
$$

We achieve this lower bound when $q(x) \propto p(x)g(x)$. A slightly useless result, because

$$
q(x) = \frac{p(x)g(x)}{\int_{\mathcal{X}} p(x)g(x)dx}
$$

requires solving the integral we care about.

We want a small variance in the weights *w*(*xⁱ*). Easy to check at $\mathbb{E}_q[w(x)] = 1$.

$$
\begin{aligned} \n\text{Var}_q[w(x)] &= \mathbb{E}_q[w(x)^2] - \mathbb{E}_q[w(x)]^2 \\ \n&= \int_{\mathcal{X}} \left(\frac{p(x)}{q(x)}\right)^2 q(x) \, \mathrm{d}x - 1 \qquad = \int_{\mathcal{X}} \frac{p(x)^2}{q(x)} \, \mathrm{d}x - 1 \n\end{aligned}
$$

Can be unbounded. E.g. $p = \mathcal{N}(0, 2)$ and $q = \mathcal{N}(0, 1)$.

A popular diagnosis statistic: effective sample size (ESS).

$$
ESS = \frac{\left(\sum_{i=1}^{N} w(x_i)\right)^2}{\sum_{i=1}^{N} w(x_i)^2}
$$

Small ESS *→* Large variability in *w*'s *→* bad estimate. Large ESS promises you nothing!

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Reuse samples from the proposal distribution *q*(*x*):

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\hat{Z} = \frac{1}{N} \sum_{i=1}^{N} \frac{f(x_i)}{q(x_i)} = \frac{1}{N} \sum_{i=1}^{N} \tilde{w}(x_i)
$$

Can use to approximate importance sampling weights $w(x_i)$:

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w(x_i) = \frac{p(x_i)}{q(x_i)} = \frac{f(x_i)}{Zq(x_i)} \approx \frac{1}{2}\tilde{w}(x_i)
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Use $\tilde{w}(x)$ instead of $w(x)$ in the Monte Carlo approximation. Is biased for finite *N*, but consistent as *N → ∞*.