

Lecture 3: Rejection Sampling

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MATH60047/70047 – Stochastic Simulation

October 17, 2022

**Imperial College
London**

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- ▶ Assignment
 - ▶ We will require a report written in Latex with figures (page limit to be announced with the assignment)

Recap

The Fundamental Theorem of Simulation

Rejection Sampling

Recap: We have seen

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- ▶ Uniform random variate generation

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However, those methods required a quite specific structure for us to be able to sample.

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What if we cannot evaluate $p(x)$ – only evaluate an unnormalised density $\bar{p}(x)$

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What if we cannot evaluate $p(x)$ – only evaluate an unnormalised density $\bar{p}(x)$

Can we still do *exact* sampling?

Is there a more general structure?

Theorem 1 (Theorem 2.2, Martino et al., 2018)

Drawing samples from one dimensional random variable X with a density $p(x)$ is equivalent to sampling uniformly on the two dimensional region defined by

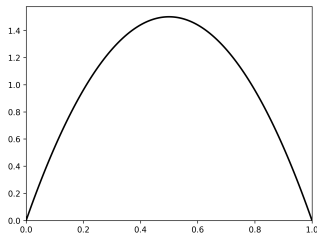
$$A = \{(x, y) \in \mathbb{R}^2 : 0 \leq y \leq p(x)\}. \quad (1)$$

In other words, if (x', y') is uniformly distributed on A , then x' is a sample from $p(x)$.

Let

$$p(x) = \text{Beta}(\alpha, \beta) = \frac{\Gamma(\alpha + \beta)}{\Gamma(\alpha) + \Gamma(\beta)} x^{\alpha-1} (1-x)^{\beta-1},$$

where $\Gamma(n) = (n-1)!$ for integers. For $\text{Beta}(2, 2)$:

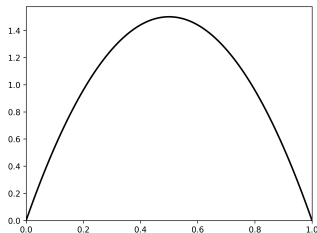


Its maximum is 1.5 in this specific case. Can we sample uniformly?

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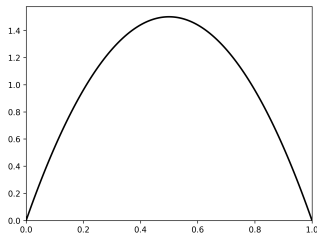
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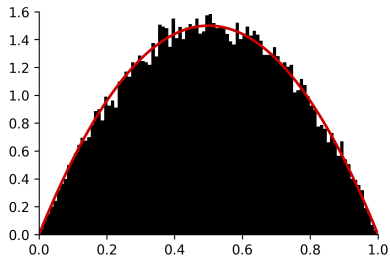
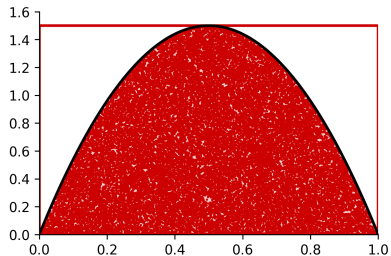


Its maximum is 1.5 in this specific case. Can we sample uniformly?

- ▶ Sample from the box $[0, 1] \times [0, 1.5]$ and keep the ones inside.
- ▶ Note though our aim is to 'test the x -marginal'

Sampling Beta density

Testing the theorem



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We will see that we can get away with an unnormalised expression of $p(x)$ (later)

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Rejection sampling

More than a box

Using a box wrapping the density is very inefficient:

- ▶ You need the maximum of the density

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which could be as hard as the sampling problem!

- ▶ For densities that are peaky, this could be wildly inefficient

Idea: Design a *proposal* density that tightly wraps the target density

Consider a (target) density $p(x)$ and a *proposal* density $q(x)$.

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For rejection sampling, we always choose a proposal such that

$$p(x) \leq Mq(x),$$

for $M \geq 1$.

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For rejection sampling, we always choose a proposal such that

$$p(x) \leq Mq(x),$$

for $M \geq 1$. Intuitively, the $Mq(x)$ curve should be **above** $p(x)$.

The rejection sampler:

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▶ $x' \sim q(x),$

Rejection sampling

The algorithm

The rejection sampler:

- ▶ $x' \sim q(x)$,
- ▶ Accept the sample x' with probability

$$a(x') = \frac{p(x')}{Mq(x')} \leq 1.$$

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How does this relate to the Fundamental Theorem of Simulation?

Rejection sampling

The algorithm: A closer look

To implement the method:

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This is how it is generally implemented!

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Rejection sampling

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- ▶ Accept

$$u' \leq p(x'),$$

This would give us (x', u') uniformly under the curve! (hence x' samples would be distributed w.r.t. $p(x)$)

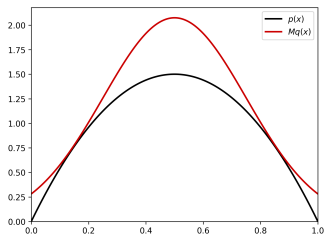
Rejection sampling

Examples: Same Beta(2, 2), better proposal

Choose

$$q(x) = \mathcal{N}(0.5, 0.25),$$

with $M = 1.3$.



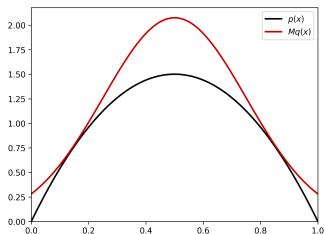
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Simulation.

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- ▶ Z is called the normalising constant
 - ▶ It is a super important quantity for many other purposes
- ▶ We write $p(x) \propto \bar{p}(x)$ to say p is proportional to $\bar{p}(x)$ but normalised to integrate (or sum) to one.

Rejection sampling

What do we do?

The fundamental theorem of simulation is better than it looks. It works with unnormalised densities (recap, unnormalised version)

Theorem 2 (Theorem 2.2, Martino et al., 2018)

Drawing samples from one dimensional random variable X with a density $p(x) \propto \bar{p}(x)$ is equivalent to sampling uniformly on the two dimensional region defined by

$$A = \{(x, y) \in \mathbb{R}^2 : 0 \leq y \leq \bar{p}(x)\}. \quad (2)$$

In other words, if (x', y') is uniformly distributed on A , then x' is a sample from $p(x)$.

Rejection sampling

The algorithm: A closer look

To implement, choose M and q such that $\bar{p}(x) \leq Mq(x)$

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Exactly same – \bar{p} used instead of p provided that $\bar{p}(x) \leq Mq(x)$

Unnormalised densities

Why would you have them?

Easy to imagine the discrete case: Imagine you want to obtain the probability of observing something discrete

- ▶ People with black jumpers: 530
- ▶ People with red jumpers: 403
- ▶ People with yellow jumpers: 304

In order to talk about 'probability of seeing a black jumper', you'd normalise these numbers (normalisation is easy to compute here).

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for some function f (which is generally called a *potential*).

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f usually comes from a rule which determines how probability mass should be spread . Ising model:

$$p_{\beta}(\sigma) = \frac{e^{-\beta H(\sigma)}}{Z_{\beta}},$$

where $H(\sigma)$ defines the energy of a spin configuration of a magnetic material. The normalising constant is

$$Z_{\beta}(\sigma) = \sum_{\sigma} e^{-\beta H(\sigma)}.$$

All configurations for atomic spins!

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In Bayesian statistics (what we care about), we are given

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$$p(y) = \int p(y|x)p(x)dx,$$

we do not know.

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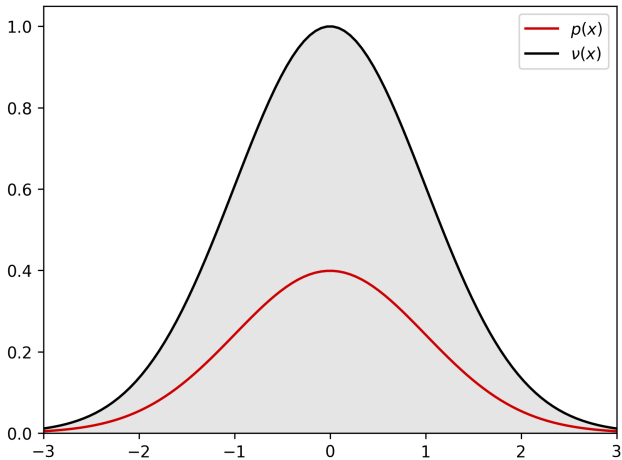
$$p(y) = \int p(y|x)p(x)dx,$$

we do not know.

Therefore, we can evaluate the posterior density only in unnormalised way:

$$p(x|y) \propto p(y|x)p(x).$$

An example of an unnormalised density.



Rejection sampling

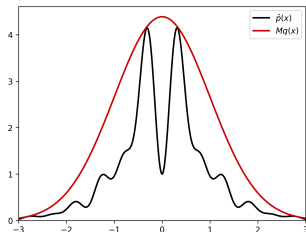
Examples: Sampling complex things

Consider (Robert and Casella, 2004)

$$p(x) \propto \exp(-x^2/2)(\sin^2(6x) + 3 \cos^2(x) \sin^2 4x + 1)$$

and

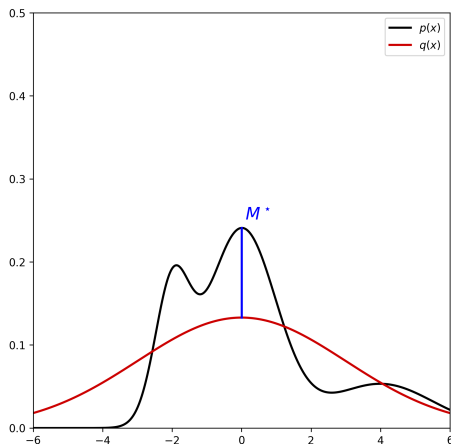
$$q(x) = \mathcal{N}(0, 1) = \exp(-x^2/2)/\sqrt{2\pi}.$$



$M = 11$.

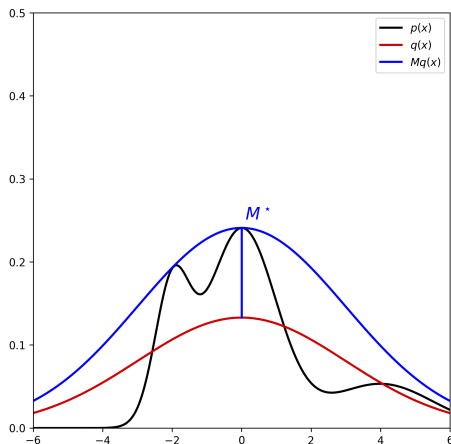
A standard choice for M is

$$M^* = \sup_x \frac{p(x)}{q(x)}.$$



A standard choice for M is

$$M^{\star} = \sup_x \frac{p(x)}{q(x)}.$$



An important quantity for a rejection sampler is *the acceptance rate*. For the normalised case, it can be analytically derived:

$$\hat{a} = \frac{1}{M}.$$

For the unnormalised case, similarly,

$$\hat{a} = \frac{Z}{M}.$$

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For the unnormalised case, similarly,

$$\hat{a} = \frac{Z}{M}.$$

We often would like to optimise our proposal q , such that the acceptance rate is maximised. \implies Minimize M !

Rejection sampling

Example: Optimising rejection sampling

Assume that we would like to sample from

$$X \sim \Gamma(\alpha, 1),$$

for $\alpha > 1$. The density is given by

$$p(x) = \frac{x^{\alpha-1}e^{-x}}{\Gamma(\alpha)}, \quad \text{for } x > 0,$$

where $\Gamma(\alpha)$ is the Gamma function. $\Gamma(n) = (n-1)!$

Rejection sampling

Example: Optimising rejection sampling

Choose as a *proposal*:

$$q_{\lambda}(x) = \text{Exp}(x; \lambda) = \lambda e^{-\lambda x}, \quad x > 0,$$

with $0 < \lambda < 1$.

Rejection sampling

Example: Optimising rejection sampling

Choose as a *proposal*:

$$q_{\lambda}(x) = \text{Exp}(x; \lambda) = \lambda e^{-\lambda x}, \quad x > 0,$$

with $0 < \lambda < 1$. How do we ensure that $p(x) \leq Mq(x)$?

Rejection sampling

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with $0 < \lambda < 1$. How do we ensure that $p(x) \leq M q_\lambda(x)$? Choose

$$M_\lambda = \sup_x \frac{p(x)}{q_\lambda(x)}.$$

Find M_λ for fixed λ first:

$$\frac{p(x)}{q_\lambda(x)} = \frac{x^{\alpha-1} e^{(\lambda-1)x}}{\lambda \Gamma(\alpha)}.$$

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Maximise this w.r.t. x ?

Rejection sampling

Example: Optimising rejection sampling

How to compute a maximum? It is useful to take log:

$$\arg \max_x f(x) = \arg \max_x \log f(x).$$

Take the log of

$$\frac{p(x)}{q_\lambda(x)} = \frac{x^{\alpha-1} e^{(\lambda-1)x}}{\lambda \Gamma(\alpha)}.$$

So we want to optimise

$$G(x) = \log \frac{p(x)}{q_\lambda(x)} = (\alpha - 1) \log x + (\lambda - 1)x - \log \lambda \Gamma(\alpha).$$

Set $\frac{dG(x)}{dx} = 0$.

Rejection sampling

Example: Optimising rejection sampling

The derivative

$$\frac{dG(x)}{dx} = \frac{\alpha - 1}{x} + (\lambda - 1) = 0$$

which implies

$$x^* = \frac{\alpha - 1}{1 - \lambda}.$$

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which implies

$$x^* = \frac{\alpha - 1}{1 - \lambda}.$$

How do we understand if this is a maximum? Compute

$$\frac{d^2G(x)}{dx^2} = -\frac{\alpha - 1}{x^2},$$

plug x^* into this

$$\frac{d^2G(x^*)}{dx^2} = -\frac{(\alpha - 1)(1 - \lambda)^2}{(\alpha - 1)^2} < 0,$$

as $\alpha > 1$ and $0 < \lambda < 1$.

Rejection sampling

Example: Optimising rejection sampling

Therefore,

$$\begin{aligned} M_\lambda &= \frac{p(x^\star)}{q_\lambda(x^\star)}, \\ &= \frac{x^{\star\alpha-1} e^{(\lambda-1)x^\star}}{\lambda\Gamma(\alpha)}, \\ &= \frac{\left(\frac{\alpha-1}{1-\lambda}\right)^{\alpha-1} e^{(\lambda-1)\frac{\alpha-1}{1-\lambda}}}{\lambda\Gamma(\alpha)} \\ &= \frac{\left(\frac{\alpha-1}{1-\lambda}\right)^{\alpha-1} e^{-(\alpha-1)}}{\lambda\Gamma(\alpha)}. \end{aligned}$$

Rejection sampling

Example: Optimising rejection sampling

Recall that, we are interested in the acceptance probability (or maximising it)

$$\frac{p(x)}{M_\lambda q_\lambda(x)} = \left(\frac{x(1-\lambda)}{\alpha-1} \right)^{\alpha-1} e^{(\lambda-1)x+\alpha-1}.$$

Now, the task is to minimise M_λ w.r.t. λ so we get the *optimal* proposal ($\hat{a} = 1/M_\lambda$ would be maximised).

Rejection sampling

Example: Optimising rejection sampling

Recall

$$M_{\lambda} = \frac{\left(\frac{\alpha-1}{1-\lambda}\right)^{\alpha-1} e^{-(\alpha-1)}}{\lambda \Gamma(\alpha)}.$$

Compute

$$\begin{aligned} \log M_{\lambda} &= (\alpha - 1) \log(\alpha - 1) - (\alpha - 1) \log(1 - \lambda) \\ &\quad - (\alpha - 1) - \log \lambda - \log \Gamma(\alpha). \end{aligned}$$

Rejection sampling

Example: Optimising rejection sampling

$$\frac{d \log M_\lambda}{d\lambda} = \frac{\alpha - 1}{1 - \lambda} - \frac{1}{\lambda} = 0,$$

which implies

$$\lambda\alpha - \lambda = 1 - \lambda,$$

therefore

$$\lambda^* = \frac{1}{\alpha}.$$

Finally we get the optimal M by computing

$$M_{\lambda^*} = \frac{\alpha^\alpha e^{-(\alpha-1)}}{\Gamma(\alpha)}.$$

Rejection sampling

Example: Optimising rejection sampling

In order to sample from $\Gamma(\alpha, 1)$, we perform

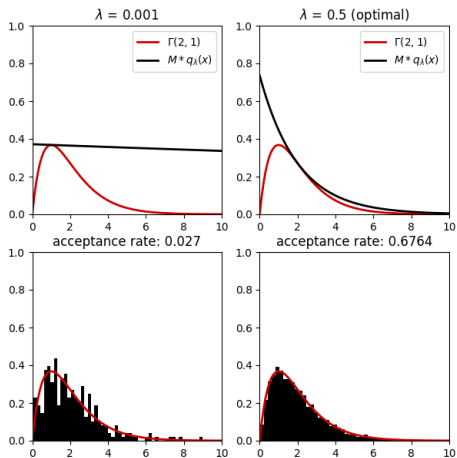
- ▶ Sample $X' \sim \text{Exp}(1/\alpha)$ and $U \sim \text{Unif}(0, 1)$
- ▶ If

$$U \leq (x/\alpha)^{\alpha-1} e^{(1/\alpha-1)x+\alpha-1},$$

accept X' , otherwise start again.

Rejection sampling

Example: Optimising rejection sampling



Rejection Sampling

Example: Sampling truncated distributions

Given $\mathcal{N}(x; 0, 1)$, suppose we are interested in sampling this density between $[-a, a]$. We can write this truncated normal density as

$$p(x) = \frac{\bar{p}(x)}{Z} = \frac{\mathcal{N}(x; 0, 1) \mathbf{1}_{\{|x| \leq a\}}(x)}{\int_{-a}^a \mathcal{N}(y; 0, 1) dy}.$$

We can choose $q(x) = \mathcal{N}(x; 0, 1)$ anyway, and we have $\bar{p}(x) \leq q(x)$ (i.e. we can take $M = 1$). The resulting algorithm is extremely intuitive: All you need is to sample from $q(x) = \mathcal{N}(x; 0, 1)$ and reject if this sample is out of bounds $[-a, a]$.

Example: Beta(2, 2)

$$p(x) = \text{Beta}(\alpha, \beta) = \frac{\Gamma(\alpha + \beta)}{\Gamma(\alpha) \Gamma(\beta)} x^{\alpha-1} (1-x)^{\beta-1},$$

Choose

$$\bar{p}(x) = x^{\alpha-1} (1-x)^{\beta-1},$$

Choose

$$q(x) = \text{Unif}(0, 1)$$

What is

$$M = \sup_x \frac{\bar{p}(x)}{q(x)}$$

Example: Beta(2, 2)

Compute

$$\log \bar{p}(x)/q(x) = (\alpha - 1) \log x + (\beta - 1) \log(1 - x)$$

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The derivative

$$\frac{d \log \bar{p}(x)/q(x)}{dx} = \frac{\alpha - 1}{x} + \frac{1 - \beta}{1 - x}$$

Example: Beta(2, 2)

Compute

$$\log \bar{p}(x)/q(x) = (\alpha - 1) \log x + (\beta - 1) \log(1 - x)$$

The derivative

$$\frac{d \log \bar{p}(x)/q(x)}{dx} = \frac{\alpha - 1}{x} + \frac{1 - \beta}{1 - x}$$

The maximum is

$$x^* = \frac{\alpha - 1}{\alpha + \beta - 2}.$$

Example: Beta(2, 2)

Since

$$M = \frac{\bar{p}(x^*)}{q(x^*)}$$

We obtain

$$M = \frac{(\alpha - 1)^{\alpha-1}(\beta - 1)^{\beta-1}}{(\alpha + \beta - 2)^{\alpha+\beta-2}}.$$

Rejection Sampling

Example: $\text{Beta}(2, 2)$

The algorithm:

- ▶ Sample $X' \sim q(x) = \text{Unif}(0, 1)$
- ▶ Sample $U \sim \text{Unif}(0, 1)$
- ▶ If $U \leq \bar{p}(X')/Mq(X')$,
 - ▶ Accept X'

See you tomorrow!

- ① Martino, Luca, David Luengo, and Joaquín Míguez (2018). *Independent random sampling methods*. Springer.
- ① Robert, Christian P and George Casella (2004). *Monte Carlo statistical methods*. Springer.