

Advanced Simulation - Lecture 3

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From a statistical problem to a sampling problem

- From a statistical model you get a likelihood function and a prior on the parameters.
- Applying Bayes rule, you are interested in

$$\pi(\theta \mid \text{observations}) = \frac{\mathcal{L}(\text{observations}; \theta)p(\theta)}{\int_{\Theta} \mathcal{L}(\text{observations}; \theta)p(\theta)d\theta}.$$

- Inference \equiv integral w.r.t. posterior distribution.
- Integrals can be approximated by Monte Carlo.
- For Monte Carlo you need samples.
- Today: inversion, transformation, composition, rejection.

- Consider a real-valued random variable X and its associated cumulative distribution function (cdf)

$$F(x) = \mathbb{P}(X \leq x) = F(x).$$

- The cdf $F : \mathbb{R} \rightarrow [0, 1]$ is
 - increasing; i.e. if $x \leq y$ then $F(x) \leq F(y)$,
 - right continuous; i.e. $F(x + \varepsilon) \rightarrow F(x)$ as $\varepsilon \rightarrow 0^+$,
 - $F(x) \rightarrow 0$ as $x \rightarrow -\infty$ and $F(x) \rightarrow 1$ as $x \rightarrow +\infty$.
- We define the generalised inverse

$$F^{-}(u) = \inf \{x \in \mathbb{R}; F(x) \geq u\}$$

also known as the quantile function.

Inversion Method

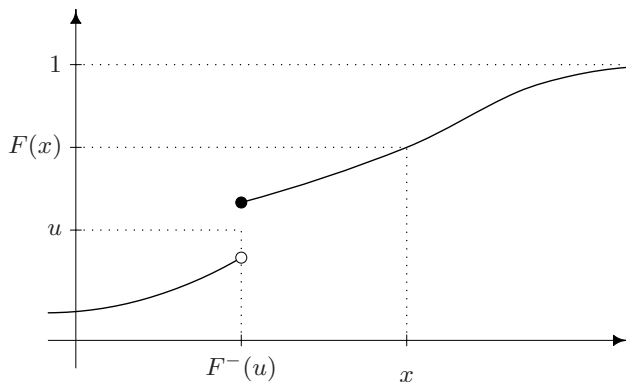


Figure: Cumulative distribution function F and representation of the inverse cumulative distribution function.

- **Proposition.** Let F be a cdf and $U \sim \mathcal{U}_{[0,1]}$. Then $X = F^{-1}(U)$ has cdf F .
- In other words, to sample from a distribution with cdf F , we can sample $U \sim \mathcal{U}_{[0,1]}$ and then return $F^{-1}(U)$.
- *Proof.* $F^{-1}(u) \leq x \Leftrightarrow u \leq F(x)$ so for $U \sim \mathcal{U}_{[0,1]}$, we have

$$\mathbb{P}(F^{-1}(U) \leq x) = \mathbb{P}(U \leq F(x)) = F(x).$$

- **Exponential distribution.** If $F(x) = 1 - e^{-\lambda x}$, then $F^-(u) = F^{-1}(u) = -\log(1 - u) / \lambda$.

Thus when $U \sim \mathcal{U}_{[0,1]}$, $-\log(1 - U) / \lambda \sim \mathcal{Exp}(\lambda)$ and $-\log(U) / \lambda \sim \mathcal{Exp}(\lambda)$.

- **Discrete distribution.** Assume X takes values $x_1 < x_2 < \dots$ with probability p_1, p_2, \dots so

$$F(x) = \sum_{x_k \leq x} p_k,$$

$$F^-(u) = x_k \text{ for } p_1 + \dots + p_{k-1} < u \leq p_1 + \dots + p_k.$$

Transformation Method

- Let $Y \sim q$ be a \mathbb{Y} -valued random variable that we can simulate (e.g., by inversion)
- Let $X \sim \pi$ be \mathbb{X} -valued, which we wish to simulate.
- It may be that we can find a function $\varphi : \mathbb{Y} \rightarrow \mathbb{X}$ with the property that if we simulate $Y \sim q$ and then set $X = \varphi(Y)$ then we get $X \sim \pi$.
- Inversion is a special case of this idea.

- **Gamma distribution.** Let $Y_i, i = 1, 2, \dots, \alpha$, be i.i.d. with $Y_i \sim \text{Exp}(1)$ and $X = \beta^{-1} \sum_{i=1}^{\alpha} Y_i$ then $X \sim \text{Ga}(\alpha, \beta)$.

Proof. The moment generating function of X is

$$\mathbb{E}\left(e^{tX}\right) = \prod_{i=1}^{\alpha} \mathbb{E}\left(e^{\beta^{-1}tY_i}\right) = (1 - t/\beta)^{-\alpha}$$

which is the MGF of the gamma density $\pi(x) \propto x^{\alpha-1} \exp(-\beta x)$ of parameters α, β .

- **Beta distribution.** See Lecture Notes.

- **Gaussian distribution.** Let $U_1 \sim \mathcal{U}_{[0,1]}$ and $U_2 \sim \mathcal{U}_{[0,1]}$ be independent and set

$$R = \sqrt{-2 \log(U_1)}, \quad \vartheta = 2\pi U_2.$$

We have

$$\begin{aligned} X &= R \cos \vartheta \sim \mathcal{N}(0, 1), \\ Y &= R \sin \vartheta \sim \mathcal{N}(0, 1). \end{aligned}$$

- Indeed $R^2 \sim \text{Exp}\left(\frac{1}{2}\right)$ and $\vartheta \sim \mathcal{U}_{[0,2\pi]}$ so

$$q(r^2, \theta) = \frac{1}{2} \exp(-r^2/2) \frac{1}{2\pi}.$$

- Bijection:

$$(x, y) = \left(\sqrt{r^2} \cos \theta, \sqrt{r^2} \sin \theta \right) \\ \Leftrightarrow (r^2, \theta) = \left(x^2 + y^2, \arctan(y/x) \right)$$

so

$$\pi(x, y) = q(r^2, \theta) \left| \det \frac{\partial (r^2, \theta)}{\partial (x, y)} \right|$$

where

$$\left| \det \frac{\partial (r^2, \theta)}{\partial (x, y)} \right|^{-1} = \left| \det \begin{pmatrix} \frac{\cos \theta}{2r} & -r \sin \theta \\ \frac{\sin \theta}{2r} & r \cos \theta \end{pmatrix} \right| = \frac{1}{2}.$$

- Hence we have

$$\pi(x, y) = \frac{1}{2\pi} \exp \left(- \left(x^2 + y^2 \right) / 2 \right).$$

Transformation Method - Multivariate Normal

- Let $Z = (Z_1, \dots, Z_d) \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, 1)$. Let L be a real invertible $d \times d$ matrix satisfying $L L^T = \Sigma$, and $X = LZ + \mu$. Then $X \sim \mathcal{N}(\mu, \Sigma)$.
- We have indeed $q(z) = (2\pi)^{-d/2} \exp\left(-\frac{1}{2}z^T z\right)$ and

$$\pi(x) = q(z) |\det \partial z / \partial x|$$

where $\partial z / \partial x = L^{-1}$ and $\det(L^{-1}) = \det(\Sigma)^{-1/2}$.
Additionally,

$$\begin{aligned} z^T z &= (x - \mu)^T (L^{-1})^T L^{-1} (x - \mu) \\ &= (x - \mu)^T \Sigma^{-1} (x - \mu). \end{aligned}$$

- In practice, use a Cholesky factorization $\Sigma = L L^T$ where L is a lower triangular matrix.

Sampling via Composition

- Assume we have a joint pdf $\bar{\pi}$ with marginal π ; i.e.

$$\pi(x) = \int \bar{\pi}_{X,Y}(x, y) dy$$

where $\bar{\pi}(x, y)$ can always be decomposed as

$$\bar{\pi}_{X,Y}(x, y) = \bar{\pi}_Y(y) \bar{\pi}_{X|Y}(x|y).$$

- It might be easy to sample from $\bar{\pi}(x, y)$ whereas it is difficult/impossible to compute $\pi(x)$.
- In this case, it is sufficient to sample

$$Y \sim \bar{\pi}_Y \text{ then } X|Y \sim \bar{\pi}_{X|Y}(\cdot|Y)$$

so $(X, Y) \sim \bar{\pi}_{X,Y}$ and hence $X \sim \pi$.

Finite Mixture of Distributions

- Assume one wants to sample from

$$\pi(x) = \sum_{i=1}^p \alpha_i \cdot \pi_i(x)$$

where $\alpha_i > 0$, $\sum_{i=1}^p \alpha_i = 1$ and $\pi_i(x) \geq 0$, $\int \pi_i(x) dx = 1$.

- We can introduce $Y \in \{1, \dots, p\}$ and

$$\bar{\pi}_{X,Y}(x, y) = \alpha_y \times \pi_y(x).$$

- To sample from $\pi(x)$, first sample Y from a discrete distribution such that $\mathbb{P}(Y = k) = \alpha_k$ then

$$X | (Y = y) \sim \pi_y.$$

Rejection Sampling

Basic idea: Sample from a proposal q different from the target π and correct through rejection step to obtain a sample from π .

Algorithm (Rejection Sampling). Given two densities π, q with $\pi(x) \leq M q(x)$ for all x , we can generate a sample from π by

- 1 Draw $X \sim q$, draw $U \sim \mathcal{U}_{[0,1]}$.
- 2 Accept $X = x$ as a sample from π if

$$U \leq \frac{\pi(x)}{M q(x)},$$

otherwise go to step 1.

- **Proposition.** The distribution of the samples accepted by rejection sampling is π .

Proof. We have for any (measurable) set A

$$\mathbb{P}(X \in A | X \text{ accepted}) = \frac{\mathbb{P}(X \in A, X \text{ accepted})}{\mathbb{P}(X \text{ accepted})}$$

where

$$\begin{aligned}\mathbb{P}(X \in A, X \text{ accepted}) &= \int_{\mathbb{X}} \int_0^1 \mathbb{I}_A(x) \mathbb{I}\left(u \leq \frac{\pi(x)}{M q(x)}\right) q(x) du dx \\ &= \int_{\mathbb{X}} \mathbb{I}_A(x) \frac{\pi(x)}{M q(x)} q(x) dx \\ &= \int_{\mathbb{X}} \mathbb{I}_A(x) \frac{\pi(x)}{M} dx = \frac{\pi(A)}{M}.\end{aligned}$$

So

$$\mathbb{P}(X \text{ accepted}) = \mathbb{P}(X \in \mathbb{X}, X \text{ accepted}) = \frac{\pi(\mathbb{X})}{M} = \frac{1}{M}$$

and

$$\mathbb{P}(X \in A | X \text{ accepted}) = \pi(A).$$

- Rejection sampling produces samples from π . It requires to be able to evaluate the density of π point-wise, and an upper bound M on $\pi(x)/q(x)$.

Rejection Sampling

- In most practical scenarios, we only know π and q up to some normalising constants; i.e.

$$\pi = \tilde{\pi}/Z_\pi \text{ and } q = \tilde{q}/Z_q$$

where $\tilde{\pi}, \tilde{q}$ are known but $Z_\pi = \int_{\mathbb{X}} \tilde{\pi}(x) dx$, $Z_q = \int_{\mathbb{X}} \tilde{q}(x) dx$ are unknown.

- If Z_π, Z_q are unknown but you can upper bound:

$$\tilde{\pi}(x)/\tilde{q}(x) \leq \tilde{M},$$

then using $\tilde{\pi}, \tilde{q}$ and \tilde{M} in the algorithm is correct.

- Indeed we have

$$\frac{\tilde{\pi}(x)}{\tilde{q}(x)} \leq \tilde{M} \Leftrightarrow \frac{\pi(x)}{q(x)} \leq \tilde{M} \frac{Z_q}{Z_\pi} = M.$$

Rejection Sampling

- Let T denote the number of pairs (X, U) that have to be generated until X is accepted for the first time.
- **Lemma.** T is geometrically distributed with parameter $1/M$ and in particular $\mathbb{E}(T) = M$.
- In the unnormalised case, this yields

$$\mathbb{P}(X \text{ accepted}) = \frac{1}{M} = \frac{Z_\pi}{\widetilde{M}Z_q},$$

$$\mathbb{E}(T) = M = \frac{Z_q \widetilde{M}}{Z_\pi},$$

and it can be used to provide unbiased estimates of Z_π/Z_q and Z_q/Z_π .

Examples

- **Uniform density on a bounded subset of \mathbb{R}^p .**

Consider the problem of sampling uniformly over $B \subset \mathbb{R}^p$, a bounded subset of \mathbb{R}^p :

$$\pi(x) \propto \mathbb{I}_B(x).$$

Let R be a rectangle with $B \subset R$ and

$$q(x) \propto \mathbb{I}_R(x).$$

- Then we can use $\widetilde{M} = 1$ and

$$\widetilde{\pi}(x) / \left(\widetilde{M}' \widetilde{q}(x) \right) = \mathbb{I}_B(x).$$

- The probability of accepting a sample is then Z_π / Z_q .

- **Normal density.** Let $\tilde{\pi}(x) = \exp\left(-\frac{1}{2}x^2\right)$ and $\tilde{q}(x) = 1/(1+x^2)$. We have

$$\frac{\tilde{\pi}(x)}{\tilde{q}(x)} = (1+x^2) \exp\left(-\frac{1}{2}x^2\right) \leq 2/\sqrt{e} = \tilde{M}$$

which is attained at ± 1 . The acceptance probability is

$$\mathbb{P}\left(U \leq \frac{\tilde{\pi}(X)}{\tilde{M}\tilde{q}(X)}\right) = \frac{Z_{\pi}}{\tilde{M}Z_q} = \frac{\sqrt{2\pi}}{\frac{2}{\sqrt{e}}\pi} = \sqrt{\frac{e}{2\pi}} \approx 0.66,$$

and the mean number of trials to success is approximately $1/0.66 \approx 1.52$.

Examples: Genetic linkage model

- We observe

$$(Y_1, Y_2, Y_3, Y_4) \sim \mathcal{M} \left(n; \frac{1}{2} + \frac{\theta}{4}, \frac{1}{4} (1 - \theta), \frac{1}{4} (1 - \theta), \frac{\theta}{4} \right)$$

where \mathcal{M} is the multinomial distribution and $\theta \in (0, 1)$.

- The likelihood of the observations is thus

$$\begin{aligned} p(y_1, \dots, y_4; \theta) &= \frac{n!}{y_1! y_2! y_3! y_4!} \left(\frac{1}{2} + \frac{\theta}{4} \right)^{y_1} \left(\frac{1}{4} (1 - \theta) \right)^{y_2 + y_3} \left(\frac{\theta}{4} \right)^{y_4} \\ &\propto (2 + \theta)^{y_1} (1 - \theta)^{y_2 + y_3} \theta^{y_4}. \end{aligned}$$

- Bayesian approach where we select $p(\theta) = \mathbb{I}_{[0,1]}(\theta)$ and are interested in

$$p(\theta | y_1, \dots, y_4) \propto (2 + \theta)^{y_1} (1 - \theta)^{y_2 + y_3} \theta^{y_4} \mathbb{I}_{[0,1]}(\theta).$$

Examples: Genetic linkage model

- Rejection sampling using a proposal $q(\theta) = \tilde{q}(\theta) = p(\theta)$ to sample from $p(\theta | y_1, \dots, y_4)$.
- To use accept-reject, we need to upper bound

$$\frac{\tilde{\pi}(\theta)}{\tilde{q}(\theta)} = \tilde{\pi}(\theta) = (2 + \theta)^{y_1} (1 - \theta)^{y_2 + y_3} \theta^{y_4}$$

- Maximum of $\tilde{\pi}$ can be found using standard optimization procedure to perform rejection sampling. For a realisation of (Y_1, Y_2, Y_3, Y_4) equal to $(69, 9, 11, 11)$ obtained with $n = 100$ and $\theta^* = 0.6$, results shown in following figure.

Examples: Genetic linkage model

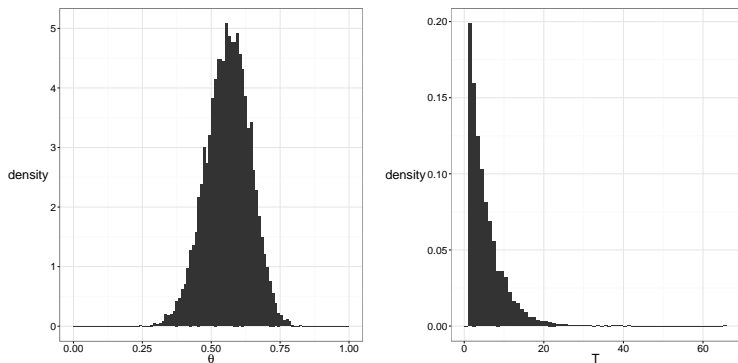


Figure: Histogram of 10,000 samples drawn from posterior obtained by rejection sampling (left); and histogram of waiting time distribution before acceptance (right).