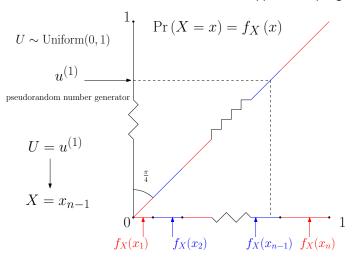
VE414 Lecture 9

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• Note for discrete random variables with a finite support, sampling is easy:



$$X = x_1, x_2, \ldots, x_n$$

Q: Suppose we can sample directly from any standard distributions, i.e. those in Appendix 2. How to sample from other distributions with a known cdf?

Theorem

Suppose $U \sim \mathrm{Uniform}(0,1)$ and F is a one-dimensional cumulative distribution, then $X = F^{-1}(U)$ has the distribution defined by F, where

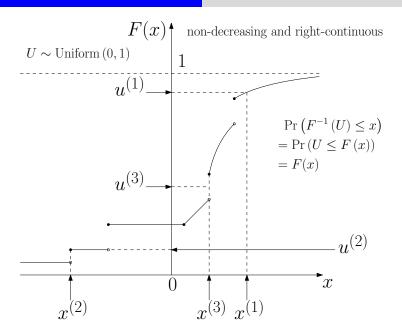
$$F^{-1}(u) = \inf\{x \colon F(x) \ge u\}$$

- Recall $\inf (S)$ denote the greatest lower bound of S.
- ullet For continuous and strictly increasing F, it is the value of x such that

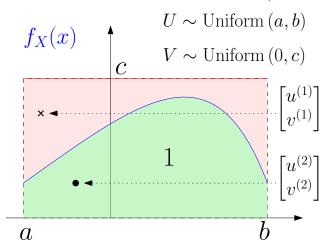
$$F(x) = u$$

• When there is discontinuity, it is the smallest "value" of x such that

$$F(x) \ge u$$

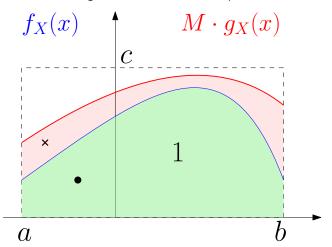


Q: How about a continuous distribution that we have its pdf but not its cdf?



Q: Why is this random sampling scheme more efficient when bc - ac is small?

• The smaller the red region the better since samples from it are wasted.



Q: How to reduce the rejections given a constant $M \in (1, \infty)$ and a pdf g_X ?

- Q: How about a distribution over any ${\cal S}$ that we have its pdf up to a constant?
 - Let $f_Y(y)$ be a pdf up to a multiplicative constant, i.e. A is unknown.

$$q_Y(y) = Af_Y(y)$$

where f_Y is the distribution from which we want to sample from.

 \bullet Suppose we have a constant $1 < M < \infty$ and a computable distribution

$$g_Y$$

that has the same support S as f_Y such that

$$q_Y(y) \le M \cdot g_Y(y)$$
 for all $y \in \mathcal{S}$

then we can sample from the distribution f_Y using rejection sampling via the distribution g_Y if we have a way to sample from g_Y . The distribution f_Y is known as the target distribution, g_Y is known as a proposal distribution.

• Let α denote the event that

$$V \le \frac{f_Y(U)}{M \cdot g_Y(U)}$$

where $U \sim g_Y$ and $V \sim \text{Uniform}(0,1)$, thus

$$f_{U,V}(u,v) = egin{cases} g_Y(u) \cdot 1, & ext{for } (u,v) \in \mathcal{S} imes [0,1], \\ 0, & ext{otherwise}. \end{cases}$$

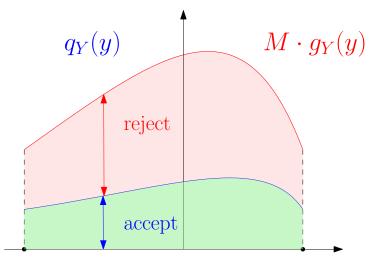
• Let U^* denote the random variable U when α has happened.

$$F_{U^*}(u^*) = \frac{\int_{-\infty}^{u^*} \int_{-\infty}^{q_Y(u)/(M \cdot g_Y(u))} f_{U,V}(u,v) \, dv \, du}{\frac{1}{M} \int_{-\infty}^{-\infty} q_Y(u) \, du} = \frac{\int_{-\infty}^{u^*} q_Y(u) \, du}{A}$$

$$\implies f_{U^*}(u^*) = f_Y(u^*)$$

from which we can conclude the random samples from this scheme follow the distribution defined by the pdf f_Y .

• Again the rejection-acceptance ratio depends on how small the red region is.



Algorithm 1: REJECTION SAMPLING

```
: functions g_Y(y), q_Y(y), constant M, number of samples n
  Output: sample array [y_i]
1 Function rejection(g_Y(y), q_Y(y) M, and n):
        i \leftarrow 0:
        while i \neq n do
             v \sim \text{Uniform}(0,1);
                                                                                   /* draw uniform */
                                                                                  /* draw from q_Y */
             y \sim g_Y;
           \begin{array}{l} \text{if } v \leq \frac{q_Y(y)}{Mg_Y(y)} \text{ then} \\ \mid i \leftarrow i+1; \\ y_i \leftarrow y; \end{array}
             end if
        end while
        return [y_i];
                                                                                           /* samples */
```

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11 12 **end** Q: How to sample from a truncated normal distribution, truncated at c=0,

$$f_Y(y) \propto I_{y>c} \phi_Y(y)$$

where $\phi_Y(y)$ is the standard normal pdf and $I_{y>c}$ is the indicator function.

- Assuming we can sample from ϕ_Y , we take a sample from ϕ_Y and simply retain the portion of the sample that is bigger than zero.
- ullet However, it will become inefficient if the truncation c is pushed to the right.
- ullet It is better to use a distribution over $(0,\infty)$ that has long tail if c is big, e.g.

$$g_Y(y) = \lambda \exp(-\lambda y)$$

- Q: Which exponential distribution should we use? And how about M?
 - ullet We want to choose the smallest M such that

$$\frac{\phi(y)}{1 - \Phi(c)} \le M\lambda e^{\lambda y} \qquad \text{for all } y \ge c$$

once M is available λ should be chosen to maximum $\Pr(\alpha)$.

• Suppose we are interested in the following expectation,

$$\mu = \mathrm{E}[h(Y)] = \int_{-\infty}^{\infty} h(y) f_Y(y) \, dy = \int_{\mathcal{D}} h(y) f_Y(y) \, dy$$

where the target distribution $f_Y(y)$ is cannot be sampled directly.

- Q: Given what we have learn so far, what would you do to estimate μ ?
 - Suppose we have a distribution

$$g_Y(y)$$

from what we can sample from over \mathcal{D} , then

$$\mu = \int_{\mathcal{D}} h(y) f_Y(y) \, dy = \int_{\mathcal{D}} h(y) f_Y(y) \frac{g_Y(y)}{g_Y(y)} \, dy$$
$$= \int_{\mathcal{D}} h(y) \frac{f_Y(y)}{g_Y(y)} g_Y(y) \, dy = \mathbb{E} \left[h(y) \frac{f_Y(y)}{g_Y(y)} \right]$$

Q: Why is this useful?

• The importance sampling estimate of

$$\mu = \mathrm{E}\left[h(Y)\right]$$

is given by the following evaluated at samples $Y_i \sim g_Y$,

$$\hat{\mu}_g = \frac{1}{n} \sum_{i=1}^n h(y_i) \frac{f_Y(y_i)}{g_Y(y_i)}$$

which means h(y) and $f_Y(y)$ as well as $g_Y(y)$ must be computable.

- Q: What happens if $g_Y(y) = 0$ for some $y \in \mathcal{D}$?
 - ullet The proposal distribution g_Y does not have to be positive everywhere in \mathcal{D} ,

$$g_Y(y) > 0$$
 whenever $h(y)f_Y(y) \neq 0$.

is sufficient for it to work.

• In practice, the value y^* will not occur in our sample if $g_Y(y^*) = 0$.

- Q: How to use importance sampling if $f_Y(y)$ is only known up to a constant?
- If $f_Y(y)$ is not computable, i.e. A is not available

$$q_Y(y) = Af_Y(y)$$

then we have to reply on the following ratios

$$\mu = \frac{\int_{\mathcal{D}} h(y) q_Y(y) \, dy}{\int_{\mathcal{D}} q_Y(y) \, dy} = \frac{\int_{\mathcal{D}} h(y) \frac{q_Y(y)}{g_Y(y)} g_Y(y) \, dy}{\int_{\mathcal{D}} \frac{q_Y(y)}{g_Y(y)} g_Y(y) \, dy}$$

which can be estimated according to the basic concept of Monte Carlo using

$$\frac{\frac{1}{n} \sum_{i=1}^{n} h(y_i) \frac{q_Y(y_i)}{g_Y(y_i)}}{\frac{1}{n} \sum_{i=1}^{n} \frac{q_Y(y_i)}{g_Y(y_i)}} = \frac{\sum_{i=1}^{n} h(y_i) w_i}{\sum_{i=1}^{n} w_i} \quad \text{where} \quad w_i = \frac{q_Y(y_i)}{g_Y(y_i)}$$

are called importance ratios.

• If $g_Y(y)$ is chosen so that the following function is roughly constant,

$$h(y)\frac{q_Y(y)}{g_Y(y)}$$

then obtaining fairly precise estimates requires fewer samples than otherwise.

Importance sampling is not reliable if the importance ratios vary a lot

$$w_i = \frac{q_Y(y_i)}{g_Y(y_i)}$$

- The worst case is when w_i and $g_Y(y_i)$ go in opposite direction, if so, we should really try some other proposal distribution.
- Plotting $\ln(w_i)$ with $g_Y(y_i)$, and comparing individual w_i with its average are traditional methods of assessing whether the estimates are poor.
- Avoid using the importance sampling estimates with a large number of small w_i with a few really big w_i .

 Importance sampling and rejection sampling are quite similar ideas. Both of them distort a sample from one distribution in order to sample from another.

$$\begin{array}{ll} \text{Rejection sampling:} & V \leq \frac{q_{Y}\left(Y\right)}{Mg_{Y}\left(Y\right)} & \text{Explicitly} \\ \\ \text{Importance sampling:} & w\left(Y\right) = \frac{q_{Y}\left(Y\right)}{g_{Y}\left(Y\right)} & \text{Implicitly} \end{array}$$

- The difference can be best understood in terms of the trade-off and the type of problems that the two methods are usually used and are good at.
- Although rejection sampling provides more, it becomes inefficient when the target distribution becomes complicated, especially as the dimension grows.
- ullet Let $\mathbf{Y} = \{Y_1, Y_2, \ldots\}$ be a random process of potentially ∞ -dimension with

$$f_{\mathbf{Y}}\left(\mathbf{Y}\right) = \prod_{j\geq 1}^{k} f_{j}\left(y_{j} \mid y_{1}, \dots, y_{j-1}\right)$$
 where $f_{1}(y_{1})$

where K is a random variable over positive integers.

Even given some appropriate proposal distribution that we can sample from

$$g_{\mathbf{Y}}\left(\mathbf{Y}
ight) = \prod_{j\geq 1}^{k} g_{j}\left(y_{j} \mid y_{1}, \ldots, y_{j-1}
ight) \qquad ext{where} \quad g_{1}(y_{1})$$

it is not clearly what M should we use in the rejection sampling

$$V \le \frac{f_{\mathbf{Y}}\left(\mathbf{Y}\right)}{Mg_{\mathbf{Y}}\left(\mathbf{Y}\right)}$$

since the exact pdfs are not simple, despite being simple enough to compute.

• Using importance sampling, we simply estimate

$$\mu = \mathbb{E}\left[h\left(\mathbf{Y}\right)\right]$$
 by $\hat{\mu}_g = \frac{1}{n} \sum_{i=1}^n h\left(\mathbf{y}_i\right) w_{k_i}$

where the set of vectors $\{\mathbf{y}_1,\cdots,\mathbf{y}_n\}$ is a sample from $g_{\mathbf{Y}}$ and

$$w_{k_i} = \prod_{j=1}^{k_i} \frac{f_j(y_j \mid y_1, \dots, y_{j-1})}{g_j(y_j \mid y_1, \dots, y_{j-1})}$$