

Importance Sampling and Variance Reduction

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Computational Methods in Statistics and Data Science (Stats 406)

AR Reject Rates

Recall two different **candidate distributions** in order to sample from a **truncated standard Normal** on $[0,1]$.

- Standard uniform rejected about **15% of the candidates**.
- The density $g(y) = (2/3)(2 - y)$ rejected about **4%**

Could we do even better and reject 0%?

Changing variables

The reason we usually need collections of X is that we want to estimate:

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Remember the trick we used for integrating arbitrary functions:

$$\int_{-\infty}^{\infty} h(x)f(x) dx = \int_{-\infty}^{\infty} h(x)f(x)\frac{g(x)}{g(x)} dx = E\left(h(Y)\frac{f(Y)}{g(Y)}\right)$$

for random variable Y with density $g(y)$.

Example: Tail probabilities

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Ex.: For $Z \sim N(0, 1)$, what is $P(Z \geq 4.5) = E(I(Z \geq 4.5))$?

```
> k <- 100000
```

```
> sum(rnorm(k) >= 4.5)
```

```
[1] 1
```

Using a shifted exponential

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```
> ys <- rexp(k) + 4.5  
> ratios <- dnorm(ys) / (dexp(ys - 4.5))  
> mean(ratios)  
  
[1] 3.416e-06  
  
> (truep <- pnorm(4.5, lower.tail = FALSE))  
  
[1] 3.398e-06
```

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Estimated variance of the importance sampling version:

```
> var(ratios) / k
```

```
[1] 1.951e-16
```

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Note: The importance weights $f(Y)/g(y)$ are very similar to the ratios we computed for the accept-rejection algorithm.

Picking Envelope Distribution

Recall that we had the strict requirement for accept-reject sampling:

$$\frac{f(x)}{c g(x)} \leq 1, \quad \text{for some } c > 0 \text{ and all } x$$

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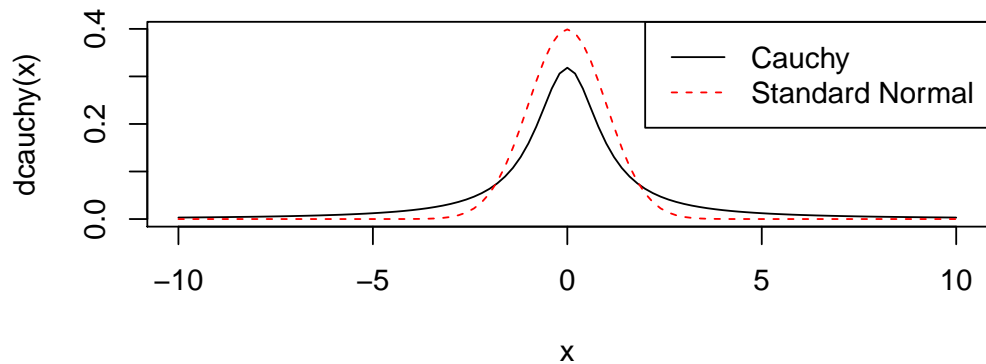
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In particular, we need the importance sampling estimator to have finite variance for the law of large numbers and the CLT to hold.

$$E_Y \left(h(Y)^2 \frac{f(Y)^2}{g(Y)^2} \right) = \int_{-\infty}^{\infty} h(y)^2 \frac{f(y)^2}{g(y)^2} g(y) dy = E_X \left(h(X)^2 \frac{f(X)}{g(X)} \right) < \infty$$

Example: Targeting the Cauchy distribution with the Normal

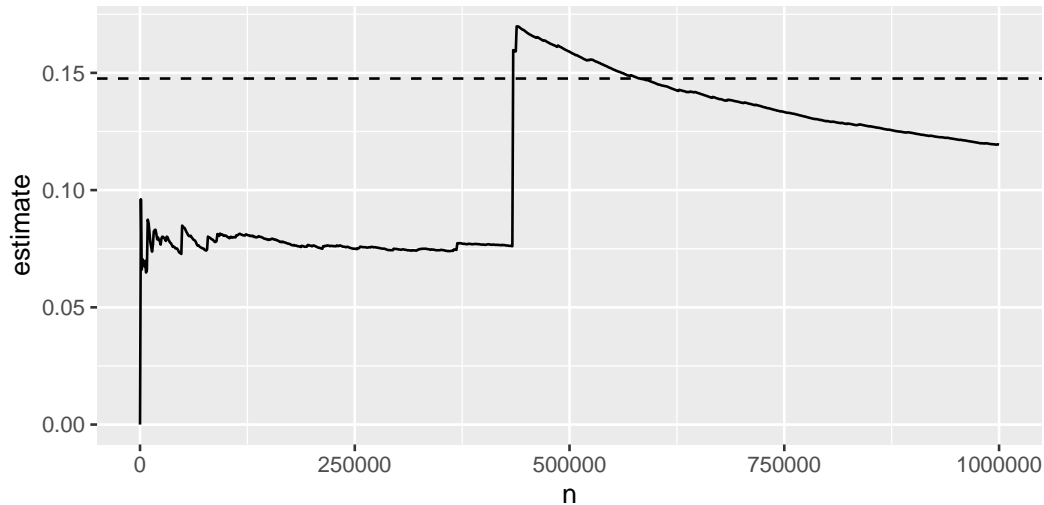


Cauchy from Normal

Let $C \sim \text{Cauchy}(0)$. Let's estimate $P(C \geq 2)$:

```
> k <- 1000000 # one million samples  
> ys <- rnorm(k)  
> iweights <- dcauchy(ys) / dnorm(ys)  
> estimates <- cumsum(iweights * (ys >= 2)) / (1:k)
```

Plotting estimate vs. number of samples



Avoiding degenerate envelopes

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Specifically, if the quantity

$$h(x) \frac{f_X}{g_X} < c, \quad \text{for some } c$$

then you should be ok.

The cases we'll consider in this class will be safe, but keep this in mind when using this a final project, e.g.

Efficiency of our estimators

Suppose we need to estimate $\theta = E(h(X))$ and we have identified **two distributions that we can sample from**:

$$E(h(X)) = E\left(h(Y)\frac{f(Y)}{g(Y)}\right) = E\left(h(W)\frac{f(W)}{d(W)}\right)$$

(where g is the PDF of Y and d is the PDF of W).

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Notice that in large samples:

$$\hat{\theta}_1 \sim N\left(\theta, \text{Var}\left(\hat{\theta}_1\right)\right), \quad \hat{\theta}_2 \sim N\left(\theta, \text{Var}\left(\hat{\theta}_2\right)\right)$$

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They only differ in the variance terms!

Variance terms

Recall that for **IID data**, the variance of the sample mean is

$$\text{Var}(\hat{\theta}_1) = \text{Var}\left(\frac{1}{n} \sum_{i=1}^n h(Y_i) \frac{f(Y_i)}{g(Y_i)}\right) = \frac{1}{n} \text{Var}\left(h(Y) \frac{f(Y)}{g(Y)}\right)$$

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Notice that for any function $t(x)^2 = |t(x)| |t(x)|$.

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Picking a $g(y)$ to minimize variance

Observe that that because $|h(x)|f(x) > 0$, the following is a proper density:

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and if $h(y) > 0$ for all y where $f(y) > 0$, then the **variance would be zero!**

Using minimum variance g

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Why is this helpful? We can try to pick $g(y) \approx c|h(y)|f(y)$

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Specifically, we want to estimate $E(X)$. We could pick many bounded distributions on $(0, 1)$. Which would be best?

- Uniform(0,1)
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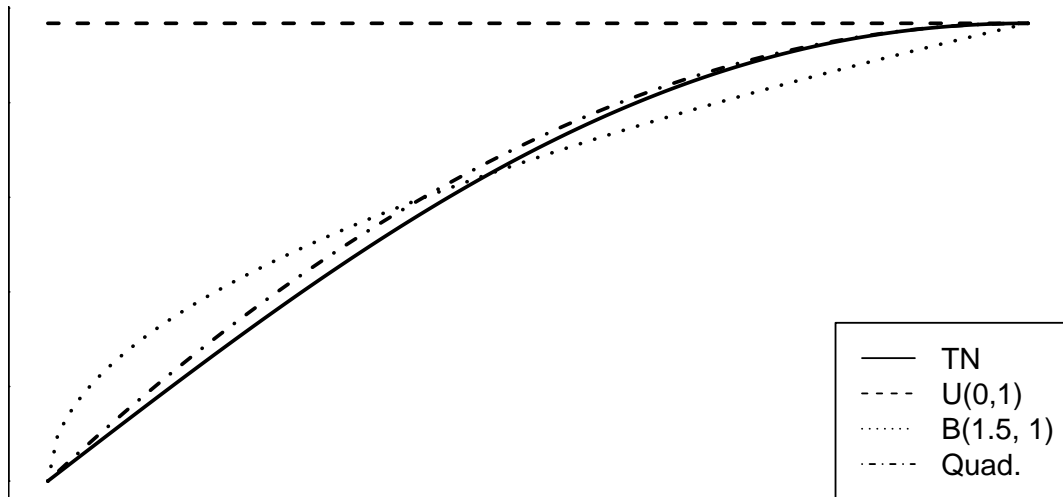
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Goal: we want a density $\propto |x|\phi(x)$.

Graphing $|h(x)|f(x)/g(x)$



Using Beta(1.5, 1) and the Quadratic Density

```
> tn <- function(x) { dnorm(x) / (pnorm(1) - pnorm(0)) }  
> k <- 10000  
> yg <- qx(runif(k))  
> iwg <- tn(yg) / gx(yg) ## h(y) f(y) / g(y)  
> mean(yg * iwg)  
  
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```
> yr <- rbeta(k, 1.5, 1)  
> iwr <- tn(yr) / dbeta(yr, 1.5, 1)  
> mean(yr * iwr)
```

```
[1] 0.4587
```

Comparing Variance

```
> varg <- var(yg * iwg) # variance of single  $h(Y) f(Y) / g(Y)$  term  
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Relative CI width:

```
> diff(t.test(yg * iwg)$conf.int) / diff(t.test(yr * iwr)$conf.int)
```

```
[1] 0.2177
```


Importance Sampling Resampling

Generating samples

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Using **importance sampling re-sampling** (ISRS) we use the **importance weights** to pick samples.

Generate m samples Y_i . Pick a number J between 1 and m with probability*

$$\frac{1}{m} \frac{f(Y_i)}{g(Y_i)}, \quad i = 1, \dots, m$$

and set $X = Y_J$ (notice: all Y are random and J is random).

Proof

The distribution of Y_J is the same as X . We need to show $P(Y_J \leq t) = P(X \leq x)$.
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$$= \int_{-\infty}^t f(y) dy = \Pr(X \leq t) \quad f \text{ is density of } X$$

Normalizing the weights

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The resulting Y_J will not have exactly the same distribution as X , but when m is large, the difference can be very small. This bias is the tradeoff for accepting all samples.

Example: Drawing from a truncated distribution of Z

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We'll use the $\text{Exp}(1)$ samples and their importance weights to estimate the conditional mean.

```
> ys <- rexp(k) + 4.5
> imp_weights <- dnorm(ys) / (dexp(ys - 4.5))
> omega <- imp_weights / sum(imp_weights) ## the 1/m term gets canceled
> xs <- sample(ys, replace = TRUE, prob = omega)
> median(xs)

[1] 4.646
```

Conclusion: Standard Normal tails go to zero really fast! 50% of all Z larger than 4.5 are within .15 of 4.5.

Densities Known to a Constant

Unnormalized Densities

Recall from **importance sampling re-sampling** we used **normalized weights**:

$$\omega_i = \frac{f(Y_i)/g(Y_i)}{\sum_{j=1}^n f(Y_j)/g(Y_j)}$$

Unnormalized Densities

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$$\omega_i = \frac{f(Y_i)/g(Y_i)}{\sum_{j=1}^n f(Y_j)/g(Y_j)}$$

Suppose **did not** actually know f , but only knew

$$f^*(x) \propto f(x) \quad \text{i.e., } f(x) = cf^*(x), \quad \text{s. t. } \int_{-\infty}^{\infty} f^*(x) dx = c$$

Unnormalized Densities

Recall from **importance sampling re-sampling** we used **normalized weights**:

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Implication: we can use ISRS to draw from f^* (essentially, we'll estimate c).

Previously we saw ISRS for **generating samples**. Here we will use it for **Monte Carlo estimation**.

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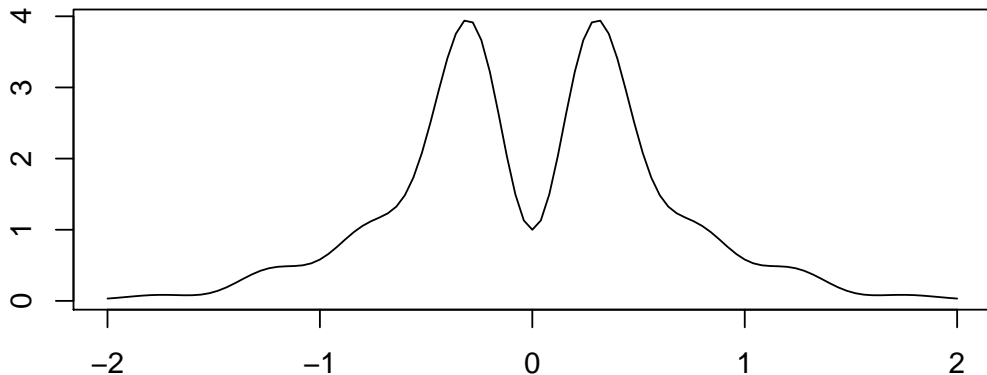
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$$\tilde{\theta} = \sum_{i=1}^n h(Y_i) \omega_i = \frac{1}{n} \sum_{i=1}^n h(Y_i) (n\omega_i)$$

Example: “Rabbit” distribution

$$f(x) \propto \exp(-x^2/2) [\sin(6x)^2 + 3 \cos(x)^2 \sin(4x)^2 + 1] = f^*(x), \quad -\infty < x < \infty$$



Estimating the variance of the rabbit distribution

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Using a standard normal as the envelope:

```
> k <- 10000
> ys <- rnorm(k)
> as <- fstar(ys) / dnorm(ys)
> omegas <- as / sum(as)
> reweighted_ys2 <- ys^2 * (k * omegas)
> mean(reweighted_ys2)

[1] 0.387
```

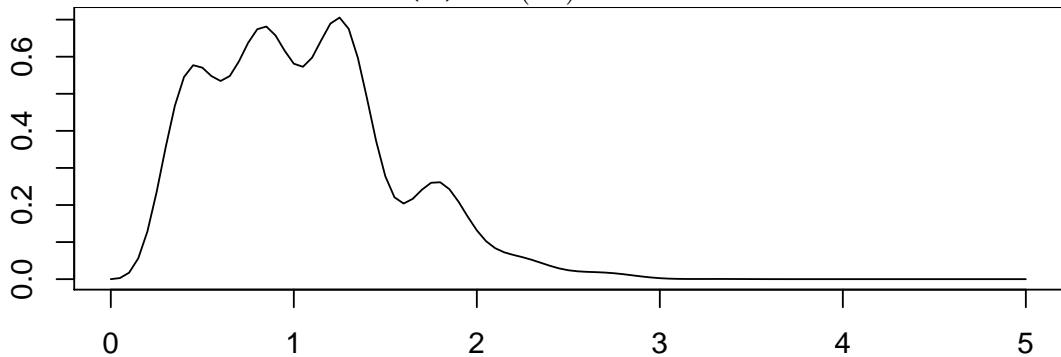
Rabbit distribution: Plotting $|h(x)|f(x)$

While achieving a **variance of zero is probably impossible**, we can tune our envelope by making it as **close to $|h(x)|f(x)$** as possible.

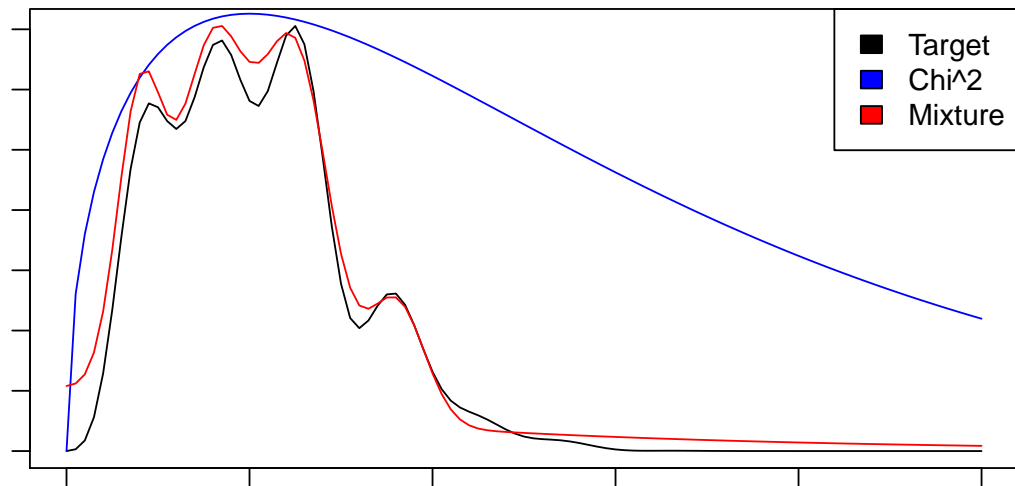
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While achieving a **variance of zero is probably impossible**, we can tune our envelope by making it as **close to $|h(x)|f(x)$** as possible.

Recall our goal of estimating $\text{Var}(X) = E(X^2)$ for the “rabbit distribution”.



A few choices



$\chi^2(3)$ and Mixture of truncated Normals and Exponential

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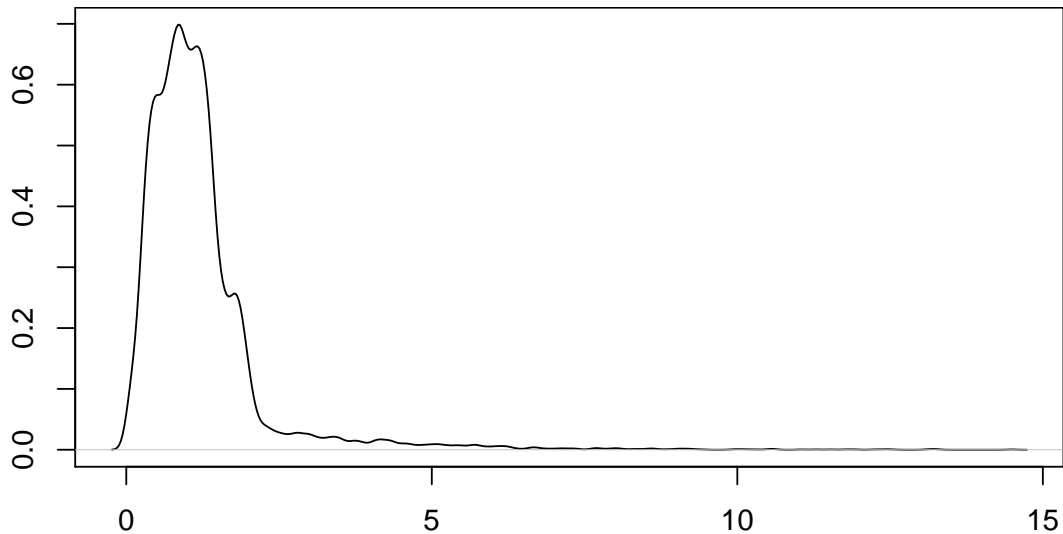
Candidate two is a mixture of truncated Normals and an Exponential:

$$\begin{aligned} &0.15 N_{[0,\infty)} \left(\frac{2}{5}, \frac{1}{8^2} \right) + 0.28 N_{[0,\infty)} \left(\frac{4}{5}, \frac{3^2}{16^2} \right) + \\ &0.28 N_{[0,\infty)} \left(\frac{5}{4}, \frac{3^2}{16^2} \right) + 0.08 N_{[0,\infty)} \left(\frac{9}{5}, \frac{5^2}{32^2} \right) + \\ &0.21 \text{Exp} \left(\frac{1}{2} \right) \end{aligned}$$

Mixture (unnormalized) Density

```
function(x) {  
  0.15 * (x >= 0) * dnorm(x, mean = 2/5, sd = 1/8) +  
  0.28 * (x >= 0) * dnorm(x, mean = 4/5, sd = 3/16) +  
  0.28 * (x >= 0) * dnorm(x, mean = 5/4, sd = 3/16) +  
  0.08 * (x >= 0) * dnorm(x, mean = 9/5, sd = 5/32) +  
  0.21 * dexp(x, 1/2)  
}  
<bytecode: 0x55f8666698e8>
```

Validating Mixture Sampler




```
> chi3 <- rchisq(k, df = 3)
> chi3_ratios <- fstar(chi3) / dchisq(chi3, df = 3)
> chi3_omegas <- chi3_ratios / sum(chi3_ratios)
> chi3_x2 <- chi3^2 * (k * chi3_omegas) #
> (chi3_est <- mean(chi3_x2))

[1] 0.3904
```

Mixture estimator

```
> mixs <- rmix(k)
> mixs_ratios <- fstar(mixs) / dmix_star(mixs)
> mixs_omegas <- mixs_ratios / sum(mixs_ratios)
> mixs_x2 <- mixs^2 * (k * mixs_omegas)
> (mixs_est <- mean(mixs_x2))

[1] 0.3893
```

Stacking up the variances

We've already seen method using a **reweighted standard Normal**.

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```
> var(reweighted_ys2)
```

```
[1] 0.05506
```

```
> var(chi3_x2)
```

```
[1] 0.2351
```

```
> var(mixs_x2)
```

```
[1] 0.01227
```

So the *Normal* envelope beats the χ^2 envelope, but the mixture beats both.

Other Variance Reduction Techniques

Additional techniques

Importance sampling is a very powerful tool, but it is not the only method of **reducing the variance of estimates**.

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We will briefly look at three others:

- Antithetic variables
- Control variates
- Stratified sampling

Antithetic variables

So far, we have been generating, **independent, identically distributed** collections of random variables.

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With **antithetic variables**, we introduce **dependence** in way to **decrease the variance** of the estimator.

The trick is to find a way to generate the dependence in a very particular way.

Thinking more about dependence and variance

Suppose we have an estimator for θ that is a sample mean of T_i : $\hat{\theta} = m^{-1} \sum_{i=1}^m T_i$
(e.g., $T_i = h(Y_i)f(Y_i)/g(Y_i)$).

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The variance of this estimator is

$$\text{Var}(\hat{\theta}) = \frac{1}{m^2} \left[\sum_{i=1}^m \text{Var}(T_i) + \sum_{i \neq j} \text{Cov}(T_i, T_j) \right]$$

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When T_i is independent of T_j is the covariance term is zero.

What if we could generate **negatively correlated** T_i and T_j ?

Example: Uniform random variables

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You proved that $U' = 1 - U$ has the same distribution as $U \sim U(0, 1)$ and

$$\text{Cov}(U, U') = E(U(1 - U)) - (1/4) = E(U) - E(U^2) - 1/4 = 1/4 - 1/3 = -1/12$$

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Since Q_X (the quantile function for X) is **monotonic**, $Q_X(U)$ and $Q_X(1 - U)$ are also **negatively correlated**.

Example continued

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In order to reduce the variance, we'll use the following identity:

$$\hat{\theta} = \frac{1}{m} \left[\sum_{i=1}^{m/2} Q(U_i) + \sum_{i=1}^{m/2} Q(1 - U_i) \right] = \frac{1}{m/2} \sum_{i=1}^{m/2} (Q(U_i) + Q(1 - U_i))/2$$

```
> iids <- runif(10000)^(1/3)
> mean(iids)

[1] 0.7518

> (est_var_iid <- var(iids) / 10000)

[1] 3.721e-06
```

Antithetic variables

To keep the sample size the same, let's only generate 5000 uniforms, then supplement those with 5000 copies of $1 - U$.

```
> tmp <- runif(5000)
> antis <- (tmp^(1/3) + (1 - tmp)^(1/3)) / 2
> mean(antis)
```

```
[1] 0.7499
```

```
> (est_var_anti <- var(antis) / 5000)
```

```
[1] 4.76e-07
```

Percent variance reduction

```
> 1 - (est_var_anti / est_var_iid)
```

```
[1] 0.8721
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An 86% reduction in variance (and we only had to generate 1/2 as many random variables).

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Note: it is not always obvious how to generate the antithetic variable pairs, but when you can, they are a very powerful tool.

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Observe that

$$E(\hat{\theta}_c) = \theta + c(\mu - \mu) = \theta \quad (\text{unbiased})$$

Variance

What is the variance of $\hat{\theta}_c$?

$$\text{Var}(\hat{\theta}_c) = \text{Var}(g(X_1)) + c^2 \text{Var}(f(X_1)) + 2c \text{Cov}(g(X_1), f(X_1))$$

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We don't have time for an example, but see **sec. 5.5 of SCR**.

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- Estimate $E(I(X \in A_i)h(X))$ using m_i samples ($\hat{\theta}_i$)
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If the variances of the individual portions are smaller than $\text{Var}(\hat{\theta})$ on average, the overall variance will be smaller for $\hat{\theta}_k$ than $\hat{\theta}$.

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- The key is picking a good envelope distribution that matches $|h(x)|f(x)$
- Control variates are a very powerful approach when possible, but requires more knowledge of the problem.
- These are often combined in practice.
- Stratified estimation is also relatively easy to implement and complements importance sampling.