MTH210: Statistical Computing

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1 Lecture-wise Summary

Lec No.	Date	Topic
1	Jan 5	FCH and Pseudorandom number generation
2	Jan 8	Pseudorandom numbers
3	Jan 9	Inverse Transform Method
4	Jan 12	Accept-Reject (discrete)
5	Jan 15	Accept-Reject (discrete)
6	Jan 16	Continuous: inverse transform and accept-reject
7	Jan 19	Accept-reject (Continuous)
8	Jan 23	Accept-reject (Continuous)
9	Jan 29	Quiz 1 and Accept-Reject
10	Jan 30	Box-Muller, Ratio-of-Uniforms
11	Feb 2	Ratio-of-Uniforms
12	Feb 5	Composition
13	Feb 6	Miscellaneous, Multivariate normal
14	Feb 10	Simple Monte Carlo
15	Feb 12	Quiz 2, Simple Importance Sampling
16	Feb 12	Simple Importance Sampling
17	Feb 13	Optimal proposals
18	Feb 16	Weighted Importance Sampling
19	Feb 26	Likelihood function, Maximum Likelihood Estimator
20	Feb 27	MLE and Linear regression
21	March 1	Linear regression and Penalized Regression
22	March 4	No-closed form solution, Newton-Raphson
23	March 5	Newton-Raphson algorithm
24	March 11	Gradient Ascent
25	March 12	Logistic Regression

2 Pseudorandom Number Generation

The building block of computational simulation is the generation of uniform random numbers. If we can draw from U(0,1), then we can draw from most other distributions. Thus the construction of sampling from U(0,1) requires special attention.

Computers can generate numbers between (0,1), which although are not exactly random (and in fact deterministic), but have the appearance of being U(0,1) random variables. These draws from U(0,1) are pseudorandom draws.

The goal in *pseudorandom* generation is to draw

$$X_1, \ldots, X_n \stackrel{\text{approx iid}}{\sim} U(0,1)$$
.

The resultant sample is as uniformly distributed as possible, and as independent as possible. We will learn about two different pseudorandom generators. These are very basic ones that are actually not really used in real life, but make our point well.

Note: After this lecture, we will always assume that all U(0,1) draws are exactly iid and perfectly random. We will forget that they are infact, pseudorandom. Pseudorandom generation is a whole field in itself; for more on this, checkout CS744 at IITK.

2.1 Multiplicative congruential method

A common algorithm to generate a sequence $\{x_n\}$ is the multiplicative congruential method:

- 1. Set seed x_0 , and positive integers a, m.
- 2. Obtain $x_t = a x_{t-1} \mod m$
- 3. Return sequence x_t/m for $t=1,\ldots,n$.

Since $x_t \in \{0, 1, ..., m-1\}$, $x_t/m \in (0, 1)$. Also note that after some finite number of steps < m, the algorithm will repeat itself, since when a seed x_0 is set, a deterministic sequence of numbers follows. Naturally, to allow for the sequence x_t to mimic uniform and random draws m should be large. Naturally, both a and m should be chosen to be large so as to avoid repetition. Typically m should be a large prime number.

Example 1. Set
$$a = 123$$
 and $m = 10$, and let $x_0 = 7$. Then $x_1 = 123 * 7 \mod 10 = 1$

```
x_2 = 123 * 1 \mod 10 = 3

x_3 = 123 * 3 \mod 10 = 9

x_4 = 123 * 9 \mod 10 = 7

x_5 = 123 * 7 \mod 10 = 1

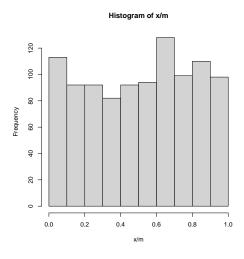
:
```

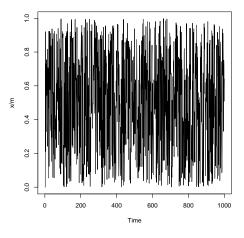
Thus, we see that the above choices of a, m, x_0 repeats itself. It is also recommended that a is large to ensure large jumps, and reduce "dependence" in the sequence. Based on the bits of your machine, it is recommended to set $m = 2^{31} - 1$ and $a = 7^5$. Notice that both are large.

```
m <- 2^(31) - 1
a <- 7^5
x <- numeric(length = 1e3)
x[1] <- 7

for(i in 2:1e3)
{
    x[i] <- (a * x[i-1]) %% m
}
par(mfrow = c(1,2))
hist(x/m) # looks close to uniformly distributed
plot.ts(x/m) # look like it's jumping around too</pre>
```

The histogram shows roughly "uniform" distribution of the samples and the trace plot shows the lack of dependence between samples.





Any pseudorandom generation method should satisfy:

- 1. for any initial seed, the resultant sequence has the "appearance" of being IID from Uniform(0,1).
- 2. for any initial seed, the number of values generated before repetition begins is large
- 3. the values can be computed efficiently.

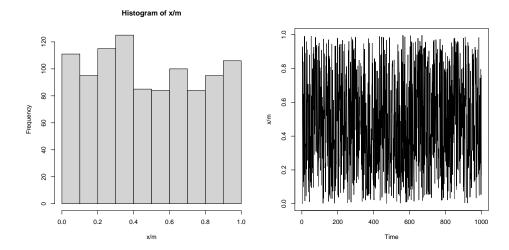
2.2 Mixed Congruential Generator

Notice that in the previous method, if we set the seed to be zero, the algorithms fails! To combat this, there is another method, the *mixed congruential generator*:

- 1. Set seed x_0 , and positive integers a, c, m.
- 2. $x_t = (a x_{t-1} + c) \mod m$
- 3. Return sequence x_t/m for t = 1, ..., n.

```
m <- 2^(31) - 1
a <- 7^5
c <- 2^(10) - 1
x <- numeric(length = 1e3)
x[1] <- 7

for(i in 2:1e3)
{
    x[i] <- (c + a * x[i-1]) %% m
}
par(mfrow = c(1,2))
hist(x/m) # looks close to uniformly distributed
plot.ts(x/m) # look like it's jumping around too</pre>
```

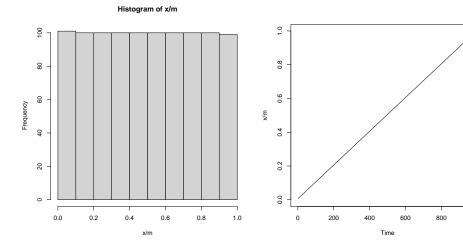


We must be cautious not to be happy with a just a histogram. A histogram shows that the empirical distribution of all samples is uniformly distributed. But we can still get a uniform looking histogram if we set a = 1, m = 1e3 and c = 1.

```
m <- 1e3
a <- 1
c <- 1
x <- numeric(length = 1e3)
x[1] <- 7

for(i in 2:1e3)
{
    x[i] <- (c + a * x[i-1]) %% m
}
par(mfrow = c(1,2))
hist(x/m) # looks VERY uniformly distributed
plot.ts(x/m) # Clearly "dependent" samples</pre>
```

Although a histogram shows an almost perfect uniform distribution, the trace plot shows that the draws don't behave like they are independent.



We could also use

$$x_n = (a_1 x_{n-1} + a_2 x_{n-2} + \dots + a_k x_{n-k} + c) \mod m$$

1000

but this requires more flops from the computer, and so is not as computationally viable.

We claim that these methods return "good" pseudosamples, in the sense of the three points stated above. There are statistical hypothesis tests, like the Kolmogorov-Smirnov test, one can do to test whether a sample is truly random: independent and identically distributed.

runif() in R uses the Mersenne-Twister generator by default (we will not go into this), but there are options to use other generators. After this, we will assume that runif() returns truly iid samples from U(0,1).

2.3 Generating U(a,b)

Suppose we can draw from U(a, b) for any $a, b \in \mathbb{R}$. But we only know how to draw from U(0, 1). Note that if $U \sim U(0, 1)$, then for any a, b,

$$(b-a)U + a \sim U(a,b) \quad .$$

That means, we can draw $U \sim U(0,1)$ and set X = (b-a)U + a. Then $X \sim U(a,b)$.

```
# Try for yourself

set.seed(1)
repeats <- 1e4
b <- 10
a <- 5
U <- runif(repeats, min = 0, max = 1)
X <- (b - a) * U + a #R is vectorized

hist(X)</pre>
```

Questions to think about

- Given a sample of pseudorandom draws from U(0,1) and perfectly IID draws from U(0,1), would you be able to tell the difference?
- Could we obtain uniform samples from \mathbb{R} ?

2.4 Exercises

1. (Using R) Consider the multiplicative congruential method. For a, m positive integers

$$x_n = ax_{n-1} \mod m$$
.

- (a) Set seed $x_0 = 5$, $m = 10^4$, a = 2. Generate $n = 10^4$ pseudorandom numbers using the above method. Does this look like a (pseudo) random sample from Uniform(0, 1)? Maybe plot a histogram to see the empirical distribution.
- (b) Now look at only the first 10 numbers: x_1, x_2, \ldots, x_{10} . What is the problem here?
- (c) How can you fix the problem noted in the previous step?

3 Generating Discrete Random Variables

Suppose X is a discrete random variable having probability mass function

$$\Pr(X = x_j) = p_j \quad j = 0, 1, \dots, \quad \sum p_j = 1.$$

Examples of such random variables are: Bernoulli, Poisson, Geometric, Negative Binomial, Binomial, etc. We will learn two methods to draw samples realizations of this discrete random variable:

- 1. Inverse transform method
- 2. The acceptance-rejection technique

3.1 Inverse transform method

Let's demonstrate the inverse transform method with an example first.

Example 2 (Bernoulli distribution). If $X \sim \text{Bern}(p)$, then

$$Pr(X = 1) = p$$
 and $Pr(X = 0) = 1 - p := q$.

Let $U \sim U(0,1)$. Define

$$X = \begin{cases} 0 & \text{if } U \le q \\ 1 & \text{if } q < U \le 1 \end{cases}.$$

Then $X \sim \text{Bern}(p)$.

Proof. To show the result we only need to show that Pr(X = 1) = p and Pr(X = 0) = 1 - p. Recall that by the cumulative distribution function of U(0, 1), for any 0 < t < 1, $Pr(U \le t) = t$. Using this,

$$\Pr(X=0) = \Pr(U \le q) = q,$$

and also

$$\Pr(X = 1) = \Pr(q < U \le 1) = 1 - q = p.$$

Algorithm 1 Inverse transform algorithm for Bern(p)

- 1: Draw $U \sim U(0, 1)$
- 2: if U < q then X = 0 else X = 1
- 3: return X

Inverse transform method: The principles used in the above example can be extended to any generic discrete distribution. For a distribution with mass function

$$\Pr(X = x_j) = p_j$$
 for $j = 0, 1, ...$ with $\sum_{j=0} p_j = 1$.

Let $U \sim U(0,1)$. Set X to be

$$X = \begin{cases} x_0 & \text{if } U \le p_0 \\ x_1 & \text{if } p_0 < U \le p_0 + p_1 \\ x_2 & \text{if } p_0 + p_1 < U \le p_0 + p_1 + p_2 \\ \vdots \\ x_j & \text{if } \sum_{i=0}^{j-1} p_i < U \le \sum_{i=0}^{j} p_i \end{cases}$$

This works because

$$\Pr(X = x_j) = \Pr\left(\sum_{i=0}^{j-1} p_i < U \le \sum_{i=0}^{j} p_i\right) = \sum_{i=0}^{j} p_i - \sum_{i=0}^{j-1} p_i = p_j.$$

This method is called the <u>inverse transform method</u> since the algorithm is essentially looking at the inverse cumulative distribution function of the random variable.

Example 3 (Poisson random variables). The probability mass function for the Poisson random variable is

$$\Pr(X = i) = p_i = \frac{e^{-\lambda} \lambda^i}{i!} \quad i = 0, 1, 2, \dots,$$

Algorithm 2 Inverse transform for Poisson(λ)

```
1: Draw U \sim U(0, 1)
```

2: if
$$U \leq p_0$$
 then

3:
$$X = 0$$

4: else if
$$U \leq p_0 + p_1$$
 then

5:
$$X = 1$$

7: else if
$$U \leq \sum_{i=1}^{j} p_i$$
 then

8:
$$X = j$$

However, Algorithm 2 outlines a challenge in implementing this algorithm.

Q. What happens when λ is large?

A Poisson(λ) distribution with a large λ will yield p_j to be small when j is small. This implies Algorithm 2 can be quite slow here. A way to make it faster is the following. We know that most likely, a realization from Poisson will be closer to λ , so it will be beneficial to start from around λ . Set $I = |\lambda|$, and check whether

$$\sum_{i=0}^{I-1} p_i < U \le \sum_{i=0}^{I} p_i.$$

If it is, then return X = I. Else, if $U > \sum_{i=1}^{I} p_i$, then increase I, otherwise, decrease I and check again.

Questions to think about

- What other example can you think of where the inverse transform method could take a lot of time?
- Can you try and implement this for a Binomial random variable?

3.2 Accept-Reject for Discrete Random Variables

Although we can draw from any discrete distribution using the inverse transform method, you can imagine that for distributions on countably infinite spaces (like the

Poisson distribution), the inverse transform method may be very expensive. In such situations, and in some other situations as well, acceptance-rejection sampling may be more reliable.

Let $\{p_j\}$ denote the pmf of the <u>target</u> distribution with $Pr(X = a_j) = p_j$ and let $\{q_j\}$ denote the pmf of another distribution with $Pr(Y = a_j) = q_j$. Suppose you can efficiently draw from $\{q_j\}$ and you want to draw from $\{p_j\}$. Let c be a constant such that

$$\frac{p_j}{q_j} \le c < \infty$$
 for all j such that $p_j > 0$.

That is,

$$c \ge \sup_{j: p_j > 0} \frac{p_j}{q_j}.$$

If we can find such a $\{q_j\}$ and c, then we can implement an Acceptance-Rejection algorithm also known as Accept-Reject sampler. Here, distribution $\{q_j\}$ is called the proposal distribution. The idea is to draw samples from $\{q_j\}$ and accept these samples if they seem likely to be from $\{p_j\}$.

Note: When $\{p_j\}$ has a finite set of states, c is always finite (since the maximum exists). However, when target distribution does not have a finite set of states, then c need not be finite, and accept-reject is not possible.

Algorithm 3 Acceptance-Rejection sampler to draw 1 sample from $\{p_j\}$ using proposal $\{q_j\}$

- 1: Draw $U \sim U(0, 1)$
- 2: Simulate Y=y from proposal, independent of U. Let $q_y=\Pr(Y=y)$ and let $p_y=\Pr(X=y)$.
- 3: if $U \leq \frac{p_y}{cq_y}$ then
- 4: Return X = y and stop
- 5: **else**
- 6: Goto step 1

Theorem 1. When c is finite, the Accept-Reject method generates a random variable with probability

$$\Pr(X = a_j) = p_j.$$

Further, the number of iterations needed to generate an acceptance is distributed as Geometric(1/c).

Proof. First, we look at the second statement. We note that the number of iterations required to stop the algorithm is clearly geometrically distributed by the definition of the geometric distribution – the distribution of the number of Bernoulli trials needed to get one success (with support 1, 2, 3...).

We will show that the probability of success is 1/c. "Success" here is an acceptance. First, see that in any iteration of the algorithm we have

$$\Pr(Y = a_j, \text{accept}) = \Pr(Y = a_j) \Pr(\text{accept} \mid Y = a_j)$$
$$= q_j \Pr\left(U \le \frac{p_j}{cq_j} \mid Y = a_j\right)$$
$$= q_j \frac{p_j}{cq_j} = \frac{p_j}{c}.$$

Using this we can calculate the marginal pmf of accepting:

$$\Pr(\text{accept}) = \sum_{j} \Pr(Y = a_j, \text{accept}) = \sum_{j} \frac{p_j}{c} = \frac{1}{c}.$$

Thus, the second statement is proved. We will now use this to show the main statement. Note that

$$\Pr(X = a_j) = \sum_{n=1}^{\infty} \Pr(a_j \text{ accepted on iteration } n)$$

$$= \sum_{n=1}^{\infty} \Pr(\text{No acceptance until iteration } n - 1) \Pr(Y = a_j, \text{accept})$$

$$= \sum_{n=1}^{\infty} \left(1 - \frac{1}{c}\right)^{n-1} \frac{p_j}{c}$$

$$= p_j.$$

This completes the proof.

Note: Since the probability of acceptance in any loop is 1/c, the expected number of loops for one acceptance is c. The larger c is, the more expensive the algorithm.

One important thing to note is that within the support $\{a_j\}$ of $\{p_j\}$, the proposal distribution must always be positive. That is, for all a_j in the support of $\{p_j\}$, $\Pr(Y = a_j) = q_j > 0$. That is,

Example 4 (Sampling from Binomial using AR). The binomial distribution has pmf

$$\Pr(X = x) = \binom{n}{x} (1-p)^{n-x} p^x \quad \text{for } x = 0, 1, \dots, n.$$

We will use AR to simulate draws from Binomial(n, p). The first task is to choose a proposal distribution. We could use any of Poisson, negative-binomial, or geometric distributions. We cannot use Bernoulli, since the support of Bernoulli does not contain the support of Binomial.

We choose to use the geometric distribution, but we must be a little careful. We use the version of geometric distribution that is defined as the number of failures before the first success, so that the support of the geometric distribution has 0 in it. The pmf of the geometric distribution is

$$Pr(X = x) = (1 - p)^x p$$
 $x = 0, 1, \dots$

We will first find c. Note that

$$\frac{p(x)}{q(x)} = \frac{\binom{n}{x}(1-p)^{n-x}p^x}{(1-p)^xp}$$
$$= \binom{n}{x}(1-p)^{n-2x}p^{x-1}.$$

Set

$$c = \max_{x=0,1,\dots,n} \binom{n}{x} (1-p)^{n-2x} p^{x-1}.$$

For n = 10, p = 0.25, we yield c = 2.373...

To be safe (since we don't know all the decimal points), we may set c to be slightly larger (say c = 2.5) as c just needs to be an upper bound. Once c is known, the AR algorithm can be implemented simply as described. Now here, we would expect, on average, 2.5 values of Geometric random variables to be proposed until *one* acceptance.

Note, that c depends on both n and p. Particularly, if n is large, then c increases drastically. A way to understand this is then the mean of the target distribution (np) can be much larger than the mean of the proposal, (1-p)/p. In this case, this implies that the bulk of the mass of the target distribution is far away from the bulk of the

mass of the proposal distribution. This is not ideal. We want the pmf of the proposal and target to match each other as much as possible, so that c is close to 1. This suggests, that we may **not** want to choose the same p in the proposal distribution!

A possible fix, is to consider a Geometric (p^*) proposal where p^* is such that the mean of the target and the mean of the proposal matches:

$$np = \frac{1 - p^*}{p^*} \Rightarrow p^* = \frac{1}{np + 1}$$

In this case, we have

$$\frac{p(x)}{q(x)} = \frac{\binom{n}{x}(1-p)^{n-x}p^x}{(1-p^*)^x p^*}$$

the maximum over $\{0, 1, ..., n\}$ can be determined on the computer. For n = 100 and p = .25, the old bound is 1028.497 and the new one is 6.0455, which is much more efficient!

Example 5. (Geometric Random Variable) We consider the Geometric random variable with pmf (trails until x failures)

$$Pr(X = x) = (1 - p)^{x}p$$
 $x = 0, 1, 2, ...$

We cannot use Binomial as a proposal, but we can use Poisson. Let us consider the $Poisson(\lambda)$ proposal. The Poisson random variable has pmf

$$\Pr(X = x) = \frac{e^{-\lambda} \lambda^x}{x!}$$
 $x = 0, 1, 2,$

First step is to find c, if it exists

$$\frac{p(x)}{q(x)} = \frac{(1-p)^x p}{\frac{e^{-\lambda} \lambda^x}{x!}}$$
$$= \frac{p}{e^{-\lambda}} \left(\frac{1-p}{\lambda}\right)^x x!.$$

For small values of λ (< 1-p), the above clearly diverges as x increases, thus the maximum doesn't exist. This is true for large values of λ as well. To see, this (intuitively),

consider the Stirling's approximation of the factorial:

$$\log(x!) \approx x \log(x) - x \Rightarrow x! \approx e^{x \log x - x}$$
.

Using this:

$$\frac{p(x)}{q(x)} = \frac{p}{e^{-\lambda}} \left(\frac{1-p}{\lambda}\right)^x x!$$

$$\approx \frac{p}{e^{-\lambda}} \left(\frac{1-p}{\lambda}\right)^x e^{x \log x - x}$$

$$= \frac{p}{e^{-\lambda}} \left(\frac{(1-p)e^{\log(x)}}{e\lambda}\right)^x$$

Thus, no matter how large λ is, eventually as x increases $e^{\log(x)}$ will be larger than λ and the ratio will diverge. Thus, this proposal does not allow an AR for the Geomtric distribution.

Question to think about

- Why is c always greater than 1?
- What happens when c is large or small?

3.3 Exercises

1. Show that if $U \sim U(0,1)$, then for any a,b,

$$(b-a)*U+a \sim U(a,b)$$

2. Use the inverse transform method to sample from a geometric distribution, where for 0 and <math>q = 1 - p,

$$\Pr(X = i) = pq^{i-1}, \quad i \ge 1, \quad \text{where } q = 1 - p.$$

3. We want to draw a sample from the random vector $(X,Y)^{\top}$, that follows the distribution with joint probability mass function

$$P(X = i, Y = j) = \theta_{i,j}$$
 where $i, j \in \{0, 1\}$.

Here $\sum_{i,j} \theta_{i,j} = 1$. Write an inverse-transform algorithm to draw realizations of $(X,Y)^{\top}$.

- 4. List as many appropriate proposal distributions as you can think of for the following target distributions:
 - Binomial
 - Bernoulli
 - Geometric
 - Negative Binomial
 - Poisson
- 5. (Using R) Draw 10,000 draws from a Binomial(20, .75) distribution using an accept-reject sampler.
- 6. In an accept-reject algorithm, we need to find c such that

$$\frac{p_j}{q_i} \le c$$
 for all j for which $p_i > 0$.

And, the probability of accepting in any iteration is 1/c. Why is c guaranteed to be more than 1?

- 7. Simulate from a Negative Binomial(n, p) using the inverse transform and accept-reject methods. Implement in R with n = 10 successes and p = .30.
- 8. Simulate from the following "truncated Poisson distribution" with pmf:

$$\Pr(X=i) = \frac{e^{-\lambda} \lambda^i / i!}{\sum_{j=0}^m e^{-\lambda} \lambda^j / j!} \quad i = 0, 1, 2, \dots, m.$$

Implement in R with m = 30 and $\lambda = 20$.

9. Suppose we want to obtain samples from a discrete distribution with pmf $\{p_i\}$. We use accept-reject with proposal distribution with pmf $\{q_i\}$, such that for some $\alpha \in \mathbb{R}$:

$$\frac{p_i}{q_i} \propto i^{\alpha} \qquad i = 1, 2, \dots, ,$$

For what values of α would this AR algorithm work?

- 10. Suppose we want to obtain samples from a discrete distribution with pmf $\{p_i\}$. Two possible proposal distributions are $\{q_i^{(1)}\}$ and $\{q_2^{(2)}\}$, yielding AR bounds c_1 and c_2 such that $c_1 > c_2$. Which proposal distribution is better?
- 11. Implement a an algorithm to sample from a Zero Inflated Binomial distribution. Can you think of an application of such a distribution?

4 Generating continuous random variables

Similar to generating discrete random variables, there are various methods for generating continuous random variables. We will discuss three main methods:

- 1. Inverse transform
- 2. The accept-reject method
- 3. Ratio of uniforms

We will also discuss a few special samplers.

4.1 Inverse transform

The principles of the inverse transform method for discrete distributions, apply similarly to continuous random variables. Consider a random variable X with probability density function f(x) so that $f(x) \geq 0$, $\int_{-\infty}^{\infty} f(x) = 1$ with distribution function

$$F(x) = \int_{-\infty}^{x} f(x) \, dx \, .$$

The following theorem will be the foundation for the inverse transform method.

Theorem 2. Let $U \sim U(0,1)$. For any continuous distribution F, a random variable $X = F^{-1}(U)$ has distribution F.

Proof. Let F_X be the distribution function of $X = F^{-1}(U)$. We need to show that $F_X = F$. Note that for any $x \in \mathbb{R}$,

$$F_X(x) = \Pr(X \le x)$$

$$= \Pr(F^{-1}(U) \le x)$$

$$= \Pr(F(F^{-1}(U)) \le F(x)) \qquad \text{(Since } F \text{ is non-decreasing)}$$

$$= \Pr(U \le F(x))$$

$$= F(x).$$

The above theorem then implies that if we can invert the CDF function, then we can obtain random draws from that random variable.

Example 6. Exponential(1): For the Exponential(1) distribution, the cdf is $F(x) = 1 - e^{-x}$. Thus,

$$F^{-1}(u) = -\log(1 - u).$$

To generate $X \sim \text{Exp}(1)$ we can thus use the following algorithm:

Algorithm 4 Exponential(1) Inverse transform

- 1: Generate $U \sim U(0,1)$
- 2: Set $X = -\log(1 U) \sim \text{Exp}(1)$

Similarly, we can draw from an Exponential (λ) distribution.

Example 7. Cauchy distribution: Cauchy distribution has pdf

$$f(x) = \frac{1}{\pi} \frac{1}{(1+x^2)},$$

and

$$u = F(x) = \int_{-\infty}^{x} f(y)dy = \frac{1}{\pi}\arctan(x) + \frac{1}{2}.$$

So, $F^{-1}(u) = \tan(\pi(u - .5))$.

Algorithm 5 Cauchy distribution

- 1: Generate $U \sim U(0,1)$
- 2: Set $X = \tan(\pi(U .5)) \sim \text{Cauchy}$

Example 8. Gamma distribution: The CDF of a Gamma (n, λ) distribution is

$$F(x) = \int_0^x \frac{\lambda e^{-\lambda y} (\lambda y)^{n-1}}{\Gamma(n)} dy.$$

Here, we don't know the CDF in closed form and thus cannot analytically find the inverse. This is an example where the inverse transform method cannot work in practice (even though it works theoretically). Thus, unlike the discrete case, this genuinely motivates the need for another method to sample from a distribution.

Questions to think about

- The CDF F(x) is a deterministic function, so how is $F^{-1}(U)$ a random quantity?
- Can we use the inverse transform method to generate sample from a normal distribution?

4.2 Accept-reject method

Suppose we cannot generate from distribution F with pdf f(x), like the Gamma distribution example in inverse transform. We can use accept-reject in a similar way as the discrete case. That is, we choose an appropriate proposal distribution with density g(x), and accept or reject it based on certain probabilities.

Let the support of F be \mathcal{X} and choose a proposal distribution G with density g(x) whose support is larger or the same as the support of F. That is, if \mathcal{Y} is the support of G then, $\mathcal{X} \subseteq \mathcal{Y}$. If we can fine c such that

$$\sup_{x \in \mathcal{X}} \frac{f(x)}{g(x)} \le c,$$

then an accept-reject sampler can be implemented.

Algorithm 6 Accept-reject for continuous random variables

- 1: Draw $U \sim U(0,1)$
- 2: Draw proposal $Y \sim G$, independently
- 3: if $U \leq \frac{f(Y)}{c \ q(Y)}$ then
- 4: Return X = Y
- 5: else
- 6: Go to Step 1.

Theorem 3. Algorithm 6 returns $X \sim F$. Further, the number of loops the AR algorithm takes to return X is distributed Geometric (1/c).

Proof. Let F_X denote the CDF of the random variable draw returned by the algorithm. For an arbitrary $x \in \mathcal{X}$. We will show that

$$F_X(x) = F(x) .$$

First, we consider the probability of acceptance:

$$\begin{aligned} & \text{Pr}(\text{accept}) = \text{Pr}\left(U \leq \frac{f(Y)}{cg(Y)}\right) \\ & = \text{E}\left[I\left(U \leq \frac{f(Y)}{cg(Y)}\right)\right] \end{aligned} \\ & = \text{E}\left[E\left[I\left(U \leq \frac{f(Y)}{cg(Y)}\right) \mid Y\right]\right] \qquad \text{using iterated expectations} \\ & = \text{E}\left[\text{Pr}\left(U \leq \frac{f(Y)}{cg(Y)} \mid Y\right)\right] \\ & = \text{E}\left[\frac{f(Y)}{cg(Y)}\right] \\ & = \int_{\mathcal{V}} \frac{f(y)}{cg(y)}g(y)dy \\ & = \frac{1}{c}\int_{\mathcal{V}} f(y)dy \\ & = \frac{1}{c}\int_{\mathcal{X}} f(y)dy + \frac{1}{c}\int_{\mathcal{V}/\mathcal{X}} f(y)dy \qquad \text{since } \mathcal{Y} \subseteq \mathcal{X} \\ & = \frac{1}{c} \; . \end{aligned}$$

Now that we have this established, consider

$$F_X(x) = \Pr(X \le x) = \Pr(Y \le x \mid \text{accept})$$

$$= \frac{\Pr\left(Y \le x, U \le \frac{f(Y)}{cg(Y)}\right)}{\Pr(\text{accept})}$$

$$= c \cdot \operatorname{E}\left[\operatorname{E}\left[I\left(Y \le x, U \le \frac{f(Y)}{cg(Y)}\right) \mid Y\right]\right]$$

$$= c \cdot \operatorname{E}\left[I\left(Y \le x\right) \operatorname{E}\left[I\left(U \le \frac{f(Y)}{cg(Y)}\right) \mid Y\right]\right]$$

$$= c \cdot \operatorname{E}\left[I\left(Y \le x\right) \frac{f(Y)}{cg(Y)}\right]$$

$$= c \cdot \int_{-\infty}^{x} \frac{f(y)}{cg(y)} g(y) dy$$

$$= \int_{-\infty}^{x} f(y) dy$$

$$= F(x).$$

From the proof, we know that Pr(accept) = 1/c, and so, just like the discrete example, the number of attempts it takes to generate an acceptance is distributed Geometric (1/c). Thus

Expected number of loops for an acceptance is = c.

Accept-reject method: intuition

At a proposed value y:

- if f(y) is large but g(y) is small means this value will not be proposed often and is a good value to accept for f, so higher probability of accepting it.
- if f(y) is small but g(y) is large, then this value will be proposed often but is unlikely for f, so accept this value less often.

We can choose any g we want as long its support is larger than other the support of f, and the resulting c is finite. However, some gs will be better than other gs, based on the expected number of iterations, c.

Example 9. Beta distribution: Consider the beta distribution Beta(4,3), where

$$f(x) = \frac{\Gamma(7)}{\Gamma(4)\Gamma(3)} x^{4-1} (1-x)^{3-1} \quad 0 < x < 1; \quad .$$

Consider a uniform proposal distribution. So that G = U(0,1) and

$$g(x) = 1$$
 for $x \in (0, 1)$.

Note that, $\mathcal{X} = \mathcal{Y}$ in this case. For this choice of g,

$$\sup_{x \in (0,1)} \frac{f(x)}{g(x)} = \sup_{x \in (0,1)} f(x)$$

We can show that maximum of f(x) occurs at x = 3/5 and

$$\sup_{x \in (0,1)} \frac{f(x)}{g(x)} = \sup_{x \in (0,1)} f(x) = 60 \left(\frac{3}{5}\right)^3 \left(\frac{2}{5}\right)^2 = 2.0736 = c.$$

Algorithm 7 Accept-reject for Beta(4,3)

1: Draw $U \sim U(0, 1)$

2: Draw proposal $Y \sim U(0,1)$

3: if $U \leq \frac{f(Y)}{c g(Y)}$ then

4: Return X = Y

5: else

6: Go to Step 1.

In order to choose a good proposal distribution (that yields a finite c), it is important to choose a g so that it has "fatter tails" than f. This ensures that as $x \to \pm \infty$, g dominates f, so that $c \to 0$ in the extremes, rather than blows up.

Example 10. Normal distribution

The target density function is

$$f(x) = \frac{1}{\sqrt{2\pi}} e^{-x^2/2} \,.$$

We know that the t-distribution has the right support and fatter tails and the "fattest" t distribution is with degrees of freedom 1, which is Cauchy. The pdf of a Cauchy distribution is

$$g(x) = \frac{1}{\pi} \frac{1}{1 + x^2} \,.$$

(We know we can sample from Cauchy using inverse transform, so that is easy.) We will need to find the supremum of the ratio of the densities. Consider

$$\frac{f(x)}{g(x)} = \frac{\pi}{\sqrt{2\pi}} (1 + x^2) e^{-x^2/2}.$$

When $x \to \infty$, $-\infty$, $e^{-x^2/2}$ decreases more rapidly than x^2 increases, the ratio tends to zero. This can be shown more formally using L'Hopital's rule.

Taking a derivative of the above ratio and setting it to 0 (and checking the second

derivative condition), yields that the supremum above occurs at x = -1, 1, so

$$\sup_{x \in \mathbb{R}} \frac{f(x)}{g(x)} = \frac{f(1)}{g(1)} = \sqrt{2\pi}e^{-1/2} \approx 1.746 \Rightarrow c = 1.80.$$

The actual algorithm can now be implemented similarly as before.

Note: Consider if the target distribution was Cauchy, and the proposal was N(0,1)? The ratio would clearly diverge as $x \to -\infty, \infty$, and thus an accept-reject sampler would not be possible.

Example 11. Sampling from a uniform circle Consider a unit circle centered at (0,0):

We are interested in sampling uniformly from within this circle. Since the area of the circle is π , the target density is

$$f(x,y) = \frac{1}{\pi}I(x^2 + y^2 < 1)$$
.

Consider the uniform distribution on the square as a proposal distribution

$$g(x,y) = \frac{1}{4}I(-1 < x < 1)I(-1 < y < 1)$$
.

First, we will find c. For x, y such that $x^2 + y^2 < 1$,

$$\frac{f(x,y)}{g(x,y)} = \frac{4}{\pi}I(x^2 + y^2 < 1) \le \frac{4}{\pi} := c.$$

Next, note that

$$\frac{f(x,y)}{cg(x,y)} = \frac{4}{\pi}I(x^2 + y^2 < 1)\frac{\pi}{4} = I(x^2 + y^2 < 1).$$

So for any (x, y) drawn from within the square, the ratio will be either 1 or 0, thus, no need to draw a uniform at all!

Note: How do we draw uniformly from within the box? Note that

$$g(x,y) = \left[\frac{1}{2} I(-1 < x < 1) \right] \left[\frac{1}{2} I(-1 < y < 1) \right] = g_1(x) \cdot g_1(y),$$

where g_1 is a density of a U(-1,1) random variable. Thus, with two independent draws $U_1, U_2 \stackrel{\text{iid}}{\sim} U(-1,1)$, we have $U_1 \times U_2$ being a draw from the uniform box.

Algorithm 8 Accept-reject for Uniform distribution on a circle

- 1: Draw proposal $(U_1, U_2) \sim U(-1, 1) \times U(-1, 1)$
- 2: **if** $U_1^2 + U_2^2 \le 1$ **then**
- 3: Return $(X, Y) = (U_1, U_2)$
- 4: else
- 5: Go to Step 1.

Questions to think about

- In A-R, do we want c to be large or small?
- Why is c guaranteed to be more than 1?
- How can you decide whether one proposal distribution is better than another proposal distribution?
- Try implementing the circle/square example in 3 dimension, 4 dimensions, and a general p dimensions. What happens to c?
- Can a similar A-R algorithm be implemented for Beta(m,n) for all $m,n\in\mathbb{Z}$?

4.2.1 Choosing a proposal

Sometimes it is difficult to find a good proposal or even one that works! That is, for a target density f(x) it can sometimes be challenging to find a proposal density g(x) such that

$$\sup_{x} \frac{f(x)}{g(x)} < \infty$$

Here are certain examples of when it may be difficult / impossible to implement accept-reject.

Example 12 (Beta). Consider a Beta(m, n)

$$f(x) = \frac{\Gamma(m+n)}{\Gamma(m)\Gamma(n)} x^{m-1} (1-x)^{n-1}$$

Depending on m and n, the Beta distribution can behave quite differently. Particularly, note that when both m, n < 1 the Beta density function is unbounded!

When m, n < 1, if we use a uniform proposal distribution

$$\sup_{x \in (0,1)} \frac{f(x)}{g(x)} = \sup_{x \in (0,1)} \frac{\Gamma(m+n)}{\Gamma(m)\Gamma(n)} x^{m-1} (1-x)^{n-1} = \infty.$$

So a Uniform distribution will not work. In fact, any proposal distribution with a bounded density will not work. So this is an example of a distribution where it is difficult to find a good proposal distribution.

However, when say $n \geq 1$, then

$$f(x) = \frac{\Gamma(m+n)}{\Gamma(m)\Gamma(n)} x^{m-1} (1-x)^{n-1}$$
$$\leq \frac{\Gamma(m+n)}{\Gamma(m)\Gamma(n)} x^{m-1}.$$

If we look at the upper bound, the function x^{m-1} on $x \in (0,1)$ can define a valid distribution if normalized. So, consider $g(x) = mx^{m-1}$, which is a proper density on $0 \le x \le 1$, and

$$\frac{f(x)}{g(x)} \le \frac{1}{m} \frac{\Gamma(m+n)}{\Gamma(m)\Gamma(n)} =: c$$

This Accept-Reject sampler can be implemented easily. Similarly if $m \ge 1$. Thus, an AR sampler is easier to implement here if one of m or n is more than (or equal to) 1.

Example 13 (Accept-Reject for Cauchy target). Consider the target density

$$f(x) = \frac{1}{\pi} \frac{1}{1+x^2} \quad x \in \mathbb{R},$$

The Cauchy distribution is known to have "fat tails" so that as $x \to \pm \infty$, the density function reduces to zero slowly. This means that it is very challenging to find g(x) that "dominates" the density in the tails.

For example, let the proposal be N(0,1). As discussed before, we will see the ratio of

the densities be

$$\frac{f(x)}{g(x)} = \frac{2\pi}{\pi} \frac{e^{x^2/2}}{1+x^2} \to \infty \text{ as } x \to \pm \infty!$$

In fact, as far we know, there are no possible standard accept-reject algorithms possible here.

4.2.2 Choosing parameters for a fixed proposal family

If you have chosen a family of proposal distributions that you know gives a finite c, it may be unclear what the best parameters for that proposal distribution is. That is, if the target f(x) and the proposal density is $g(x \mid \theta)$ (where θ is a parameter you can change, to change the behaviour of the proposal, then you want to find a value of the parameter θ so that the resulting proposal is the "best".

Notice that the upper bound will be a function of θ , so that

$$\sup_{x} \frac{f(x)}{g(x|\theta)} \le c(\theta) .$$

The value $c(\theta)$ is the expected number of loops for the accept-reject algorithm. Since we want this to be small, the best proposal density within this family would be the one that minimizes $c(\theta)$, so set

$$\theta^* := \arg\min_{\theta} c(\theta)$$
.

Example 14 (Gamma distribution). Consider the target distribution Gamma(α, β)

$$f(x) = \frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha - 1} e^{-\beta x}.$$

Further, suppose we want to use an $\text{Exp}(\lambda)$ proposal. Then

$$g(x|\lambda) = \lambda e^{-\lambda x}.$$

We can now find $c(\lambda)$,

$$\frac{f(x)}{g(x)} = \frac{\beta^{\alpha}}{\Gamma(\alpha)} \frac{x^{\alpha - 1} e^{-\beta x}}{\lambda e^{-\lambda x}}$$

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

First note that no matter what λ is, if $0 < \alpha < 1$, then $f(x)/g(x) \to \infty$ as $x \to 0$. So accept-reject with this proposal won't work!

However, when $\alpha \geq 1$, then $x^{\alpha-1}$ increases, so we want to choose λ such that $e^{-x(\beta-\lambda)}$ decreases (since exponential decay is more powerful than polynomial increase) (of course, you should show this more mathematically). Thus we want $\beta > \lambda$!

Thus, we restrict attention to $\alpha \geq 1$, $\lambda < \beta$,

$$c(\lambda) = \sup_{x} \frac{f(x)}{g(x)} = \sup_{x>0} \frac{\beta^{\alpha}}{\lambda \Gamma(\alpha)} x^{\alpha-1} e^{-x(\beta-\lambda)}$$

which you can show, occurs at

$$x = \frac{\alpha - 1}{\beta - \lambda},$$

for which

$$c(\lambda) = \frac{\beta^{\alpha}}{\lambda \Gamma(\alpha)} \left(\frac{\alpha - 1}{\beta - \lambda} \right)^{\alpha - 1} e^{1 - \alpha},$$

which is minimized for

$$\lambda = \beta/\alpha$$
.

Thus, the optimal exponential proposal for the Gamma $(\alpha, \beta), \alpha > 1$ is $\text{Exp}(\beta/\alpha)$.

Questions to think about

• How would you implement accept-reject for $Gamma(\alpha, \beta)$ for $0 < \alpha < 1$?

4.3 The Box-Muller transformation for N(0,1).

A classical method to generate samples from N(0,1) is the Box-Muller transformation method. Here, we will draw random variables (R^2, Θ) from a certain distribution in the polar coordinate system, and then use a transformation h, so that $h(R^2, \Theta) \sim N(0, 1)$. To find the h, we will need some theory for this.

Let X and $Y \stackrel{\text{iid}}{\sim} N(0,1)$. The joint density of (X,Y) is

$$f(x,y) = \frac{1}{2\pi} e^{-x^2/2} e^{-y^2/2} \quad x \in \mathbb{R}, y \in \mathbb{R}.$$

Let (R^2,Θ) denote the polar coordinates of (X,Y) so that, $X=R\cos\Theta$ and Y=

 $R \sin \Theta$; here the support of R is $(0, \infty)$ and the support of Θ is $(0, 2\pi)$. Then,

$$R^2 = X^2 + Y^2 \qquad \tan \Theta = \frac{Y}{X} \,.$$

Notationally, we denote a realization from (R^2, Θ) as (d, θ) and find the joint density of $f(d, \theta)$. Thus, let $d = x^2 + y^2$ and $\theta = \tan^{-1}(y/x)$. We know that the density for (d, θ) can be found by

$$f(d,\theta) = |J|f(x,y)$$
 where $J = \begin{vmatrix} \frac{\partial x}{\partial d} & \frac{\partial y}{\partial d} \\ \frac{\partial x}{\partial \theta} & \frac{\partial y}{\partial \theta} \end{vmatrix}$

Solving for J,

$$J = \begin{vmatrix} \frac{\partial \sqrt{d} \cos \theta}{\partial d} & \frac{\partial \sqrt{d} \sin \theta}{\partial d} \\ \frac{\partial \sqrt{d} \cos \theta}{\partial \theta} & \frac{\partial \sqrt{d} \sin \theta}{\partial \theta} \end{vmatrix} = \begin{vmatrix} \frac{1}{2} \frac{\cos \theta}{\sqrt{d}} & \frac{1}{2} \frac{\sin \theta}{\sqrt{d}} \\ -\sqrt{d} \sin \theta & \sqrt{d} \cos \theta \end{vmatrix} = \frac{1}{2}.$$

Since $d = x^2 + y^2$, the joint density of (R^2, Θ) is $f(d, \theta)$ with

$$f(d,\theta) = \frac{1}{2} \frac{1}{2\pi} e^{-d/2} \qquad 0 < d < \infty, 0 < \theta < 2\pi$$

$$= \underbrace{\frac{1}{2\pi} I(0 < \theta < 2\pi)}_{U(0,2\pi)} \quad \underbrace{\frac{1}{2} e^{-d/2} I(0 < d < \infty)}_{\text{Exp(2)}}$$

This is a separable density, so R^2 and Θ are independent, and $\Theta \sim U[0, 2\pi]$ and $R^2 \sim \text{Exp}(2)$.

To generate from Exp(2), we can use an inverse transform method. If $U \sim U(0,1)$, then by the inverse transform method, $-2 \log U \sim \text{Exp}(2)$ (verify for yourself). To generate from $U(0,2\pi)$, we know if $U \sim U(0,1)$, then $2\pi U \sim U(0,2\pi)$. The Box-Muller algorithm then is given in Algorithm 9 which produces X and Y from N(0,1) indendently.

Algorithm 9 Box-Muller algorithm for N(0,1)

- 1: Generate U_1 and U_2 from U(0,1) independently
- 2: Set $R^2 = -2 \log U_1$ and $\Theta = 2\pi U_2$
- 3: Set $X = R\cos(\Theta) = \sqrt{-2\log U_1}\cos(2\pi U_2)$
- 4: and $Y = R \sin(\Theta) = \sqrt{-2 \log U_1} \sin(2\pi U_2)$.

4.4 Ratio-of-Uniforms

Ratio-of-uniforms is a powerful, however not so popular method to generate samples for a continuous random variables. When it works, it can work really well. The method is based critically on the following theorem.

Theorem 4. Let f(x) be a target density with support \mathcal{X} and distribution function F. Define the set

$$D = \left\{ (u, v) : 0 \le u \le \sqrt{f\left(\frac{v}{u}\right)} \right\}.$$

If D bounded, let (U, V) be uniformly distributed over the set D; then $V/U \sim F$.

Proof. We will show that the density of Z = V/U is f(z). Note that by definition, the joint density of (U, V) is

$$g_{(U,V)}(u,v) = \frac{1}{\int \int_D du \, dv} I\{(u,v) \in D\}$$
.

Consider transformation $(U, V) \mapsto (U, Z)$ with Z = V/U. Then U = U and V = UZ. It's easy to see that the Jacobian for this transformation is U. So

$$g_{(U,Z)}(u,z) = \frac{u}{\int \int_D du \, dv} I\left\{0 \le u \le f^{1/2}(z)\right\}.$$

Now that we have the joint distribution of (U, Z), all we need to show is that the marginal distribution of Z is F. Finding the marginal density of Z = V/U, we integrate out U,

$$g_Z(z) = \int \frac{u}{\int \int_D du \, dv} I\left\{0 \le u \le f^{1/2}(z)\right\} \, du$$
$$= \frac{1}{\int \int_D du \, dv} \int_0^{f^{1/2}(z)} u \, du$$
$$= \frac{f(z)}{2 \int \int_D du \, dv} \, .$$

Since $g_Z(z)$ and f(z) are both densities, this implies that

$$1 = \int g_Z(z)dz = \frac{\int f(z)dz}{2 \int \int_D du \, dv} = \frac{1}{2 \int \int_D du \, dv} \Rightarrow \int \int_D du \, dv = \frac{1}{2}$$

This implies $f_Z(z) = f(z)$. Thus, Z = V/U has the desired distribution.

So if we can draw $(U, V) \sim \text{Unif}(D)$, then $V/U \sim F$. But D looks quite complicated, so how do we uniformly draw from D?

Think back to the AR technique used to draw uniformly from a circle! If D is a bounded set, then if we enclose D in a rectangle, we can use accept-reject to draw uniform draws from D! So, the task is to find $[0, a] \times [b, c]$ such that

$$0 \le u \le a$$
 $b \le v \le c$ for all $(u, v) \in D$.

We just need to find any such a, b, c. First, note that if $\sup_x f^{1/2}(x)$ exists, then

$$0 \le u \le f^{1/2} \left(\frac{v}{u}\right) \le \sup_{x \in \mathcal{X}} f^{1/2}(x) =: a.$$

Note now that inside D, if $x = v/u \Rightarrow v/x = u \le f^{1/2}(x)$. This implies that

$$\frac{v}{x} \le f^{1/2}(x) .$$

Now for:

$$x \ge 0: \quad v \le x f^{1/2}(x) \le \sup_{x \in \mathcal{X}} x f^{1/2}(x) =: c$$

$$x \le 0: \quad v \ge x f^{1/2}(x) \ge \inf_{x \in \mathcal{X}} x f^{1/2}(x) =: b.$$

Note that if $\sqrt{f(x)}$ or $x^2f(x)$ are unbounded, then D is unbounded, and the method cannot work. Now that we have found the rectangle: $[0, a] \times [b, c]$, we can propose from the rectangle, check if the proposed value is in the region D; if it is, we accept it and return V/U. This leads to the following algorithm:

Algorithm 10 Ratio-of-Uniforms

- 1: Generate $(U,V) \sim U[0,a] \times U[b,c]$
- 2: If $U \leq \sqrt{f(V/U)}$, then set X = V/U.
- 3: Else go to 1.

Steps 1 and 2 in Algorithm 10 are implementing an Accept-Reject to sample uniformly from D. To understand how effective this algorithm will be, we can calculate the

probability of acceptance for the AR. First, note that

$$\sup_{(u,v)\in D} \frac{f(u,v)}{g(u,v)} = \sup_{(u,v)\in D} \frac{\frac{I((u,v)\in D)}{\int_C dudv}}{\frac{1}{a*(c-b)}} = 2a(c-b)$$

Thus,

$$\Pr\left(\text{Accepting for AR in RoU}\right) = \frac{1}{2a(c-b)}.$$

So if a is large and/or (c - b) is large, the probability is small, and thus the algorithm will take a large number of loops to yield one acceptance.

Example 15 (Exponential(1)).

$$f(x) = e^{-x} \quad x \ge 0$$

Here,

$$D = \{(u, v) : 0 \le u \le e^{-v/2u}\}.$$

Since $e^{-x/2}$ is a decreasing function, $a = \sup_x e^{-x/2} = 1$. Additionally,

$$b = \inf_{x \le 0} x e^{-x/2} = 0$$
 (since support is $x \ge 0$)

and

$$c = \sup_{x \ge 0} x e^{-x/2} \Rightarrow c = 2e^{-1}$$
 (show for yourself).

So we sample from $U[0,1] \times [0,2/e]$ and then implement accept-reject.

Example 16 (Normal(θ, σ^2)). The target density is:

$$f(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-(x-\theta)^2/2\sigma^2}$$
.

The set D is

$$D = \left\{ (u, v) : 0 \le u \le \left(\frac{1}{2\pi\sigma^2} \right)^{1/4} e^{-(v - u\theta)^2/4\sigma^2 u^2} \right\}$$

In order to draw the region later, we need to rearrange the bound above, which gives us (by taking log):

$$(v - \theta u)^2 \le -4\sigma^2 u^2 \left(\log u + \frac{1}{4} \log(2\pi\sigma^2) \right).$$

The above defines the region D. Now, in order to bound the region D, we find the limits a, b, c:

$$a = \sup_{x \in \mathbb{R}} (2\pi\sigma^2)^{-1/4} e^{-(x-\theta)^2/4\sigma^2} = (2\pi\sigma^2)^{-1/4}$$

$$b = \inf_{x \le 0} \left(\frac{1}{2\pi\sigma^2} \right)^{1/4} x e^{-(x-\theta)^2/4\sigma^2} \quad \text{and} \quad c = \sup_{x \ge 0} \left(\frac{1}{2\pi\sigma^2} \right)^{1/4} x e^{-(x-\theta)^2/4\sigma^2}$$

First, we find b and then c will follow similarly. Note that b will be non-positive, and thus, to find the infimum, we first take negative and then log. That is, let for x < 0, let

$$A(x) = \left(\frac{1}{2\pi\sigma^2}\right)^{1/4} (-x)e^{-(x-\theta)^2/4\sigma^2}$$

Then A(x) is non-negative, and we want to find the supremem of A(x) for $x \leq 0$. Taking log:

$$\log(A(x)) = -\frac{1}{4}\log(2\pi\sigma^2) + \log(-x) - \frac{(x-\theta)^2}{4\sigma^2}$$

$$\Rightarrow \frac{d\log(A(x))}{dx} = \frac{1}{x} - \frac{(x-\theta)}{2\sigma^2} \stackrel{set}{=} 0$$

$$\Rightarrow x = \frac{\theta \pm \sqrt{\theta^2 + 8\sigma^2}}{2}$$

Now, we need to decide which of \pm would be choose. Note that $\sqrt{\theta^2 + 8\sigma^2} > \theta$. Hence, since we are taking $\sup_{x\geq 0} A(x)$, we obtain

$$x_b := \frac{\theta - \sqrt{\theta^2 + 8\sigma^2}}{2}$$

Thus,

$$b = x_b f^{1/2}(x_b)$$

Similarly, we obtain

$$x_c := \frac{\theta + \sqrt{\theta^2 + 8\sigma^2}}{2}$$

with

$$c = x_c f^{1/2}(x_c) .$$

All that needs to be done now is to implement Algorithm 10 with these values of a, b, c,

given the values of θ and σ^2

Questions to think about

1. Construct a similar RoU sampler for Cauchy distribution.

2. Why does RoU fail when D is unbounded?

3. For N(0,1) between RoU and AR using Cauchy proposal, which is more efficient, in terms of the expected number of uniforms required for one acceptance?

4.5 The Composition Method

We have now learned many algorithm for sampling distributions. For certain special distributions, it is easier to use a *composition method* for sampling.

Suppose we have an efficient way of simulating random variables from two pmfs $\{p_j^{(1)}\}\$ and $\{p_j^{(2)}\}\$, and we want to simulate from

$$\Pr(X = j) = \alpha p_j^{(1)} + (1 - \alpha) p_j^{(2)} \quad j \ge 0 \text{ where } 0 < \alpha < 1.$$

First you should note that the above composition pmf is a valid pmf since $\sum_{j} \Pr(X = j) = 1$. How would we sample in such a situation?

Let $X_1 \sim P^{(1)}$ and $X_2 \sim P^{(2)}$. Set

$$X = \begin{cases} X_1 & \text{with probability } \alpha \\ X_2 & \text{with probability } 1 - \alpha \end{cases}.$$

Algorithm 11 Composition method

1: Draw $U \sim U(0, 1)$

2: if $U \leq \alpha$ then simulate $X_1 \sim P^{(1)}$ else simulate X_2 and stop

Proof. Consider

$$\Pr(X=j)$$
 = $\Pr(X=j, U \le \alpha) + \Pr(X=j, \alpha < U \le 1)$ (by law of total probability)

$$= \Pr(X = j \mid U \leq \alpha) \Pr(U \leq \alpha) + \Pr(X = j \mid \alpha < U \leq 1) \Pr(\alpha < U \leq 1)$$

$$= \Pr(X_1 = j \mid U \leq \alpha) \Pr(U \leq \alpha) + \Pr(X_2 = j \mid \alpha < U \leq 1) \Pr(\alpha < U \leq 1)$$

$$= \Pr(X_1 = j) \Pr(U \leq \alpha) + \Pr(X_2 = j) \Pr(\alpha < U \leq 1)$$
 (by independence of U and X_1, X_2)
$$= \alpha p_j^{(1)} + (1 - \alpha) p_j^{(2)}.$$

We can set this up more generally for k different distributions. In general, F_i , i = 1, ..., k are distribution functions, and α_i are such that $0 < \alpha_i < 1$ for all i and $\sum_i \alpha_i = 1$. The composition (or mixture) distribution is

$$F(x) = \sum_{i=1}^{k} \alpha_i F_i(x) .$$

If each of the F_j are continuous distributions with densities f_j , then the composition or mixture density is

$$f(x) = \sum_{i=1}^{k} \alpha_i f_i(x) .$$

Let $X_i \sim F_i$. To simulate from the composition F, set

$$X = \begin{cases} X_1 & \text{with probability } \alpha_1 \\ X_2 & \text{with probability } \alpha_2 \\ \vdots & & \\ X_k & \text{with probability } \alpha_k \end{cases}.$$

Example 17 (Zero inflated Poisson distribution). A Poisson(λ) distribution usually has a small mass at 0. But sometimes, we need a counting distribution with large mass at 0. For example, consider the random variable X being the number of COVID-19 patients tested positive every hour. Many hours of the day this number may be 0, and then this number can be quite high for some hours.

In such a case, we may use the zero inflated Poisson distribution (ZIP). Recall that if $X \sim \text{Poisson}(\lambda)$

$$Pr(X = k) = e^{-\lambda} \frac{\lambda^k}{k!} \quad k = 0, 1, \dots$$

If $X \sim \text{ZIP}(\delta, \lambda)$ for $\delta > 0$

$$\Pr(X = k) = \begin{cases} \delta + (1 - \delta)e^{-\lambda} & \text{if } k = 0\\ (1 - \delta)e^{-\lambda}\frac{\lambda^k}{k!} & \text{if } k = \{1, 2, \dots\} \end{cases}.$$

Note that the mean of a ZIP is $(1 - \delta)\lambda < \lambda$ since more mass is given at 0. We will use the composition method to sample from the ZIP distribution. To sample from a ZIP, first $p_j^{(1)}$ be defined as

$$Pr(X_1 = 0) = 1$$
 and $Pr(X_1 \neq 0) = 0$,

and let $X_2 \sim \text{Poisson}(\lambda)$. Define the pmf:

$$\Pr(X = k) = \delta p_k^{(1)} + (1 - \delta) p_k^{(2)}.$$

Then $X \sim \text{ZIP}(\delta, \lambda)$. To see this, plug in k = 0 and $k = 1, 2, \ldots$ above:

Algorithm 12 Zero inflated Poisson distribution

- 1: Draw $U \sim U(0,1)$
- 2: if $U \leq \delta$ then X = 0 else simulate $X \sim \text{Poisson}(\lambda)$

Other composition or mixture distributions are also possible. Think about Zero-inflated Binomial, Zero-inflated Geometric, 2-inflated Poisson, etc.

Example 18 (Mixture of normals). Consider two normal distributions $N(\mu_1, \sigma_1^2)$ and $N(\mu_1, \sigma_2^2)$. For some 0 , the mixture density is

$$f(x) = pf_1(x; \mu_1, \sigma_1^2) + (1 - p)f_2(x; \mu_2, \sigma_2^2)$$

$$= p\frac{1}{\sqrt{2\pi\sigma_1^2}} \exp\left\{-\frac{1}{2} \left(\frac{x - \mu_1}{\sigma_1}\right)^2\right\} + (1 - p)\frac{1}{\sqrt{2\pi\sigma_2^2}} \exp\left\{-\frac{1}{2} \left(\frac{x - \mu_2}{\sigma_2}\right)^2\right\}$$

Mixture distributions are particularly useful for clustering problems and we will come back to them again in the data analysis part of the course. If we want to sample from this distribution

Algorithm 13 Sampling from a Gaussian mixture

- 1: Generate $U \sim U[0, 1]$
- 2: If U < p, generate $N(\mu_1, \sigma_1^2)$
- 3: Otherwise, generate $N(\mu_2, \sigma_2^2)$.

Example 19 (Zero-inflated gamma distribution). Just like the zero-inflated Poisson distribution, there are zero-inflated normal and Gamma distributions. Let's motivate the zero-inflated Gamma distribution:

Suppose you are an auto-insurance company and you want to study the cost of claims associated with each customer. That is, each customer, if they have an accident, will come to you and claim insurance money reimbursement for the accident. So

Let X = insurance money asked for by a customer in a month.

However, most customers will not enter into any accidents, so they will claim Rs 0. But when they do, they will claim reimbursement for some amount of money that, say, will follow a Gamma distribution.

The density function can be defined as follows for 0

$$f(x) = p\mathbb{I}(x=0) + (1-p)\frac{\beta^{\alpha}}{\Gamma(\alpha)}x^{\alpha-1}e^{-x\beta}.$$

Algorithm 14 Sampling from a zero-inflated Gamma

- 1: Generate $U \sim U[0,1]$
- 2: If U < p, return X = 0
- 3: Otherwise, generate $X \sim \text{Gamma}(\alpha, \beta)$.

4.6 Miscellaneous methods in sampling

4.6.1 Known relationships

It is always useful to remember the relationships between different distributions.

1. **Binomial distribution**: We know that if $Y_1, Y_2, \ldots, Y_n \stackrel{iid}{\sim} \text{Bern}(p)$, then

$$X = Y_1 + Y_2 + \dots Y_n \sim \operatorname{Bin}(n, p).$$

So, we can simulate n Bernoulli variables, add them up, and we have a realization from a Binomial(n, p).

2. **Negative binomial distribution**: Number of failures until the rth success. So possibly related to geometric! If $Y_1, Y_2, \ldots, Y_r \stackrel{iid}{\sim} \text{Geom}(p)$ (on failures), then

$$X = Y_1 + Y_2 + \cdots + Y_r \sim NB(r, p).$$

3. Beta distribution If $X \sim \text{Gamma}(a, 1)$ and $Y \sim \text{Gamma}(b, 1)$, then

$$\frac{X}{X+Y} \sim \text{Beta}(a,b)$$
.

4. **Dirichlet distribution**: The Dirichlet distribution is a distribution over pmf.

$$f(x_1, x_2, \dots, x_k) = \frac{\Gamma(\alpha_1 + \dots + \alpha_k)}{\prod_{i=1}^k \Gamma(\alpha_i)} \prod_{i=1}^k x_i^{\alpha_i - 1} \quad 0 \le x_i \le 1, \sum_{i=1}^k x_i = 1.$$

The Dirichlet distribution is a generalization of the Beta distribution. Similarly,

$$Y_1 \sim \text{Gamma}(\alpha_1, 1)$$

$$Y_2 \sim \text{Gamma}(\alpha_2, 1)$$

:

$$Y_k \sim \text{Gamma}(\alpha_k, 1)$$

Let

$$X_i = \frac{Y_i}{\sum_{i=1}^k Y_i} \,.$$

Then $(X_1, \ldots, X_k) \sim \text{Dir}(\alpha_1, \alpha_2, \ldots, \alpha_k)$.

5. Chi-squared distribution: If $Y_1, Y_2, \dots, Y_k \stackrel{iid}{\sim} N(0, 1)$, then

$$X = Y_1^2 + Y_2^2 + \dots + Y_k^2 \sim \chi_k^2$$
.

This way we can simulate χ^2 distributions with integer degrees of freedom.

6. t-distribution Let $Z \sim N(0,1)$ and $Y \sim \chi_k^2$, then

$$X = \frac{Z}{\sqrt{\frac{Y}{k}}} \sim t_k \,.$$

7. Location-scale family: Let F be a distribution in the location-scale family. Then, if Z has CDF $F_Z(z)$ in the sense that it doesn't have any parameters. Then for $\mu \in \mathbb{R}$ and $\sigma > 0$,

$$Y = \mu + \sigma Z$$
 has CDF $F_Y(y) = F_Z\left(\frac{y - \mu}{\sigma}\right)$.

If Z has pdf f(z) then Y has pdf $\sigma^{-1}f((y-\mu)/\sigma)$.

So, if
$$Z \sim N(0,1)$$
, then $Y = \mu + \sigma Z \sim N(\mu, \sigma^2)$.

4.6.2 Multidimensional target

We have almost entirely focused on univariate densities, but most often interest is in multivariate/multidimensional target distribution.

• Conditional Distribution: Consider a variable $\mathbf{X} = (X_1, X_2, \dots, X_k)$, with a joint pdf

$$f(\mathbf{x}) = f(x_1, x_2, \dots, x_k).$$

We can use conditional distribution properties:

$$f(\mathbf{x}) = f_{X_1}(x_1) f_{X_2|X_1}(x_2) \dots f_{X_k|X_1,\dots,X_{k-1}}(x_k)$$
.

Algorithm 15 Sampling X using conditional distributions

- 1: Generate $X_1 \sim f_{X_1}(x_1)$
- 2: Generate $X_2 \sim f_{X_2|X_1}(x_2)$
- 3: Generate $X_3 \sim f_{X_3|X_2,X_1}(x_3)$
- 4: :
- 5: Generate $X_k \sim f_{X_k|X_{k-1},...,X_1}(x_k)$
- 6: Return $\mathbf{X} = (X_1, \dots, X_k)$
 - Multivariate normal: Consider sampling from a $N_k(\mu, \Sigma)$ where Σ is positive definite. Then for $|\cdot|$ denoting determinant,

$$f_{\mathbf{X}}(\mathbf{x}) = \left(\frac{1}{2\pi}\right)^{k/2} |\Sigma|^{-1/2} \exp\left\{-\frac{(\mathbf{x}-\mu)^T \Sigma^{-1} (\mathbf{x}-\mu)}{2}\right\},$$

is the density of a multivariate normal distribution with mean μ and covariance Σ . First, note that since Σ is a positive-definite (symmetric) matrix, we can use the eigenvalue decomposition

$$\Sigma = Q\Lambda Q^{-1}$$

where Q is the matrix of eigenvectors and since Σ is symmetric, Q is guaranteed to be an orthongal matrix so that $Q^{-1} = Q^T$ and Λ is a diagonal matrix of eigenvalues. Then, we can define the *square-root* of Σ as

$$\Sigma^{1/2} := Q \Lambda^{1/2} Q^{-1}$$
,

so that

$$\Sigma^{1/2}\Sigma^{1/2} = Q\Lambda^{1/2}Q^{-1}Q\Lambda^{1/2}Q^{-1} = Q\Lambda Q^{-1}.$$

Similarly, the inverse square-root is

$$\Sigma^{-1/2} = Q\Lambda^{-1/2}Q^{-1} \,,$$

Set $\mathbf{Z} = \Sigma^{-1/2}(\mathbf{X} - \mu)$. Then

$$\mathbf{Z} \sim N_k(0, I_k)$$
.

That is, **Z** is a k-dimensional multivariate normal distribution with an identity covariance matrix. Which implies if $\mathbf{Z} = (Z_1, \dots, Z_k)$, then $Cov(Z_i, Z_j) = 0$ for

all $i \neq j$.

For the normal distribution, if the covariance is zero, then the random variables are independent! This isn't true in general but is true for normal random variables.

So, to sample from $N_k(\mu, \Sigma)$, we can sample $Z_1, Z_2, \ldots, Z_k \stackrel{iid}{\sim} N(0, 1)$, and set $\mathbf{Z} = (Z_1, \ldots, Z_k)$. Then

$$\mathbf{X} := \mu + \Sigma^{1/2} \mathbf{Z} \sim N_k(\mu, \Sigma)$$
.

Then $\mathbf{X} \sim N_k(\mu, \Sigma)$.

Questions to think about

• Can you construct a zero-inflated normal distribution and find a suitable application of it?

4.7 Exercises

- 1. Using the inverse transform method, simulate from $\text{Exp}(\lambda)$ for any $\lambda > 0$. Implement this for $\lambda = 5$.
- 2. Use the inverse transform method to obtain samples from the Weibull(α, λ)

$$f(x) = \alpha \lambda x^{\alpha - 1} e^{-\lambda x^{\alpha}}, \quad x > 0.$$

3. (Ross 5.1) Give a method for generating a random variable having density function

$$f(x) = \frac{e^x}{e - 1} \qquad 0 \le x \le 1.$$

4. (Ross 5.2) Give a method for generating a random variable having density function

$$f(x) = \begin{cases} \frac{x-2}{2} & \text{if } 2 \le x \le 3\\ \frac{2-x/3}{2} & \text{if } 3 \le x \le 6 \end{cases}$$

5. (Ross 5.3) Use the inverse transform method to generate a random variable having

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distribution function

$$F(x) = \frac{x^2 + x}{2}$$
 $0 \le x \le 1$.

6. Sample following the following distribution using two different methods:

$$f(x) = \begin{cases} \frac{3}{4}(1 - x^2) & \text{if } x \in (-1, 1) \\ 0 & \text{otherwise} \end{cases}$$

7. (Ross 5.6) Let X be an Exp(1). Provide an efficient algorithm for simulating a random variable whose distribution is the conditional distribution of X given that X < 0.05. That is, its density function is

$$f(x) = \frac{e^{-x}}{1 - e^{-0.05}}$$
 $0 < x < 0.05$.

Using R generate 1000 such random variables and use them to estimate $\mathrm{E}[X\mid X<0.05].$

8. (Ross 5.7) Suppose it is relatively easy to generate random variables from any of the distributions F_i , i = 1, ..., k. How could we generate a random variable from the distribution function

$$F(x) = \sum_{i=1}^{n} p_i F_i(x),$$

where $p_i \geq 0$ and $\sum p_i = 1$.

9. (Ross 5.8) Using the previous exercise, provide algorithms for generating random variables from the following distributions:

(a)
$$F(x) = \frac{x+x^3+x^5}{3}, 0 \le x \le 1.$$

(b)
$$F(x) = \begin{cases} \frac{1 - e^{-2x} + 2x}{3} & \text{if } x \in (0, 1) \\ \frac{3 - e^{-2x}}{3} & \text{if } x \in [1, \infty) \end{cases}$$

10. (Ross 5.9) Give a method to generate a random variable with distribution function

$$F(x) = \int_0^\infty x^y e^{-y} dy \qquad 0 \le x \le 1$$

11. (Ross 5.15) Give two methods for generating a random variable with density function

$$f(x) = xe^{-x}, 0 \le x < \infty.$$

12. (Ross 5.18) Give an algorithm for generating a random variable having density function

$$f(x) = 2xe^{-x^2}, \qquad x > 0.$$

13. (Ross 5.19) Show how to generate a random variable who distribution function is

$$F(x) = \frac{x + x^2}{2}, \qquad 0 \le x \le 1$$

using the inverse transform, accept-reject, composition method.

14. (Ross 5.20) Use the AR method to find an efficient way to generate a random variable having density function

$$f(x) = \frac{(1+x)e^{-x}}{2}$$
 $0 < x < \infty$.

- 15. (Ross 5.21) Consider the target density to be a truncated $Gamma(\alpha, 1)$, $\alpha < 1$ defined on (a, ∞) for some a > 0. Suppose the proposal distribution is a truncated exponential (λ) , defined on the same (a, ∞) . What is the best λ to use?
- 16. (Using R)
 - (a) Implement an accept-reject sampler to sample uniformly from the circle $\{x^2 + y^2 \le 1\}$ and obtain 10000 samples and estimate the probability of acceptance. Does it approximately equal $\pi/4$?
 - (b) Now consider sampling uniformly from a p-dimensional sphere (a circle is p=2). Consider a p-vector $\mathbf{x}=(x_1,x_2,\ldots,x_p)$ and let $\|\cdot\|$ denote the Euclidean norm. The pdf of this distribution is

$$f(\mathbf{x}) = \frac{\Gamma\left(\frac{p}{2} + 1\right)}{\pi^{p/2}} I\{\|\mathbf{x}\| \le 1\}.$$

Use a uniform p-dimensional hypercube to sample uniformly from this sphere. Implement this for p = 3, 4, 5, and 6. What happens as p increases?

17. (Using R)

(a) Using accept-reject and a standard normal proposal, obtain samples from a truncated standard normal distribution with pdf:

$$f(x) = \frac{1}{\Phi(a) - \Phi(-a)} \frac{1}{\sqrt{2\pi}} e^{-x^2/2} I(-a < x < a),$$

where $\Phi(\cdot)$ is the CDF of a standard normal distribution. Run for a=4 and a=1. What are the differences between the two settings.

(b) Now consider a multivariate truncated normal distribution, where for $\mathbf{x} = (x_1, x_2, \dots, x_p)$, the pdf is

$$f(\mathbf{x}) = \left(\frac{1}{\Phi(a) - \Phi(-a)}\right)^p \left(\frac{1}{\sqrt{2\pi}}\right)^p e^{-\mathbf{x}^T \mathbf{x}/2} I(-a < \mathbf{x} < a).$$

Implement an accept-reject sampler with proposal distribution $N_p(0, I)$ with a = 4 and p = 3, 10 and with a = 1 and p = 3, 10. Describe the differences between these settings.

- 18. Implement an accept-reject sampler to draw from a Gamma(α , 1) for $\alpha > 1$. Using the above method, can you draw samples from Gamma(α , β), for any β ?
- 19. In accept-reject sampling, why is $c \ge 1$?
- 20. Use ratio-of-uniforms method to sample from a truncated exponential distribution with density

$$f(x) = \frac{e^{-x}}{1 - e^{-a}} \quad 0 < x < a.$$

How efficient is this algorithm?

21. Use ratio-of-uniforms method to sample from the distribution with density

$$f(x) = \frac{1}{x^2} \quad x \ge 1.$$

- 22. Use ratio-of-uniforms method to draw samples from a t_{ν} distribution for $\nu \geq 1$.
- 23. (Zero-inflated Gamma distribution) Suppose you are an auto-insurance company and you want to study the cost of claims associated with each customer. That is, each customer, if they have an accident, will come to you and claim insurance money reimbursement for the accident. So

Let X = insurance money asked for by a customer in a month.

However, most customers will not enter into any accidents, so they will claim Rs 0. But when they do, they will claim reimbursement for some amount of money that, say, will follow a Gamma distribution.

The density function can be defined as follows for 0

$$f(x) = p\mathbb{I}(x=0) + (1-p)\frac{\beta^{\alpha}}{\Gamma(\alpha)}x^{\alpha-1}e^{-x\beta}.$$

Provide an algorithm to sample this random variable.

5 Importance Sampling

We have so far learned many (many!) ways of sampling from different distributions. These sampling methodologies are particularly useful when we want to estimate characteristic of F. Using computer simulated samples from F to estimate characteristics of F is broadly termed as $Monte\ Carlo$

5.1 Simple Monte Carlo

Suppose F is a distribution with density f. We are interested in estimating the expectation of a function $h: \mathcal{X} \to \mathbb{R}$ with respect to F. That is, we want to estimate

$$\theta := \mathrm{E}_F[h(X)] = \int_{\mathcal{X}} h(x) f(x) \, dx < \infty,$$

we assume that θ is finite. We also assumed that

$$\sigma^2 = \operatorname{Var}_F(h(X)) < \infty$$
.

Note: there is no "data" here, there is just an integral! We are just interested in estimating an annoying integral.

Note: notation $E_F[X]$ means the expectation is with respect to F. From now on, it is very important to keep track of what the expectation is with respect to.

Suppose we can draw iid samples $X_1, \ldots, X_N \stackrel{\text{iid}}{\sim} F$ (this we can do using the many methods we have learned). Then, by the weak law of large numbers, as $N \to \infty$,

$$\hat{\theta} = \frac{1}{N} \sum_{t=1}^{N} h(X_t) \stackrel{p}{\to} \theta.$$

In addition, we can find the variance of the estimator:

$$\operatorname{Var}(\hat{\theta}) = \operatorname{Var}\left(\frac{1}{N} \sum_{t=1}^{N} h(X_t)\right)$$

$$= \frac{1}{N^2} \sum_{t=1}^{N} \operatorname{Var}_F(h(X_t)) \quad \text{because of independence}$$

$$= \frac{\operatorname{Var}_F(h(X_1))}{N} \quad \text{because of identical}$$

$$=\frac{\sigma^2}{N}$$
.

Naturally, a central limit theorem also holds if $\sigma^2 < \infty$, so that as $N \to \infty$

$$\sqrt{N}(\hat{\theta} - \theta) \stackrel{d}{\to} N(0, \sigma^2)$$
,

This central limit theorem gives us an expected behavior of $\hat{\theta}$ for large values of n.

Q. But is there a way we can obtain a better estimator of θ ?

A. Possibly by using importance sampling.

5.2 Simple importance sampling

Our goal is the same. For $h: \mathcal{X} \to \mathbb{R}$, we want to estimate $\theta = \mathrm{E}_F[h(X)]$. Similar to the the accept-reject sampler, we will choose a proposal distribution. Let G be a distribution with density g defined on \mathcal{X} so that,

$$E_{F}[h(X)] = \int_{\mathcal{X}} h(x)f(x)dx$$

$$= \int_{\mathcal{X}} \frac{h(x)f(x)}{g(x)}g(x) dx$$

$$= E_{G} \left[\frac{h(Z)f(Z)}{g(Z)} \right], \qquad Z \sim G$$

If $Z_1, \ldots, Z_N \stackrel{\text{iid}}{\sim} G$, then an estimator of θ is

$$\hat{\theta}_g = \frac{1}{N} \sum_{t=1}^N \frac{h(Z_t) f(Z_t)}{g(Z_t)}.$$

The estimator $\hat{\theta}_g$ is the *importance sampling estimator*, the method is called *importance sampling* and G is the *importance distribution*.

Let

$$w(Z_t) = \frac{f(Z_t)}{g(Z_t)}$$

be the weights assigned to each point Z_t . Then $\hat{\theta}_g$ is a weighted average of $h(Z_t)$. Intuitively, this means that depending on how likely a sampled value is for f and g, a weight is assigned to that value.

Example 20 (Moments of Gamma distribution). Suppose we want to estimate the kth moment of a Gamma distribution. That is, let F be the density of a Gamma(α, β) distribution. Then

$$\theta = \int_0^\infty x^k \frac{\beta^\alpha}{\Gamma(\alpha)} x^{\alpha - 1} e^{-\beta x} dx.$$

Suppose we set G to be also an Exponential(λ) distribution. Let $Z_1, \ldots, Z_N \sim \text{Exp}(\lambda)$

$$\hat{\theta}_g = \frac{1}{N} \sum_{t=1}^N \left[\frac{h(Z_t) f(Z_t)}{g(Z_t)} \right]$$

$$= \frac{1}{N} \sum_{t=1}^N \left[\frac{\beta^{\alpha}}{\Gamma(\alpha)} \frac{Z_t^k Z_t^{\alpha - 1} e^{-\beta Z_t}}{\lambda e^{-\lambda Z_t}} \right].$$

So we have now constructed an alternative estimator of θ . In fact, a different choice of G will yield a different estimator. It is now important to study the properties of this importance sampling estimator. We study a sequence of properties.

Theorem 5 (Unbiasedness). The importance sampling estimator $\hat{\theta}_g$ is unbiased for θ .

Proof. To show an estimator is unbiased, we need to show that $E[\hat{\theta}_g] = \theta$. Consider

$$E\left[\hat{\theta}_g\right] = E_G \left[\frac{1}{N} \sum_{t=1}^N \frac{h(Z_t) f(Z_t)}{g(Z_t)}\right]$$

$$= \frac{1}{N} \sum_{t=1}^N E_G \left[\frac{h(Z_t) f(Z_t)}{g(Z_t)}\right]$$

$$= \frac{1}{N} \sum_{t=1}^N E_G \left[\frac{h(Z_1) f(Z_1)}{g(Z_1)}\right]$$

$$= \int_{\mathcal{X}} \frac{h(z) f(z)}{g(z)} g(z) dz$$

$$= \int_{\mathcal{X}} h(z) f(z) dz$$

$$= \theta.$$

Theorem 6. The importance sampling estimator is consistent for θ . That is, as $N \to \infty$,

$$\hat{\theta}_q \stackrel{p}{\to} \theta$$
.

Proof. Note that $\hat{\theta}_g$ is just a sample average:

$$\hat{\theta}_g = \frac{1}{N} \sum_{t=1}^N \frac{h(Z_t) f(Z_t)}{g(Z_t)}.$$

The law of large numbers applies to any sample average whose expectation is finite. So by the law of large numbers, as $N \to \infty$,

$$\hat{\theta}_g \stackrel{p}{\to} \mathrm{E}[\hat{\theta}_g] = \theta$$
.

This means that as we get more and more samples from G, our estimator will get increasingly closer to the truth.

However, we should never be happy with a point estimator!

It is essential to quantify the variability in our estimator $\hat{\theta}_g$ in order to ascertain how "erratic" or "stable" the estimator is. We also want to establish expected behavior for $\hat{\theta}_g$, but does a central limit theorem hold?. Notice that a simple Monte Carlo is just a sample average, so we should be able to directly apply the CLT result, if the variance is finite. Note that, the variance of $\hat{\theta}_g$ is

$$\operatorname{Var}(\hat{\theta}_g) = \operatorname{Var}_g\left(\frac{1}{N}\sum_{t=1}^N \frac{h(Z_t)f(Z_t)}{g(Z_t)}\right) = \frac{1}{N}\operatorname{Var}_g\left(\frac{h(Z_1)f(Z_1)}{g(Z_1)}\right) =: \frac{\sigma_g^2}{N}.$$

A central limit theorem will hold if $\sigma_g^2 = \operatorname{Var}_g\left(\frac{h(Z_1)f(Z_1)}{g(Z_1)}\right) < \infty$.

So the question is, when is this finite?

The following theorem provides a sufficient condition.

Theorem 7. Suppose $\sigma^2 = \operatorname{Var}_F(h(X)) < \infty$. If g is chosen such that

$$\sup_{z \in \mathcal{X}} \frac{f(z)}{g(z)} \le M < \infty$$

then

$$\sigma_g^2 < \infty$$
 .

Proof. First note that if the variance of a random variable is finite, this is equivalent to saying that the second moment of that variable is finite. So, consider the second moment of $\frac{h(Z)f(Z)}{g(Z)}$ where $Z \sim G$.

$$E_G \left[\left(\frac{h(Z)f(Z)}{g(Z)} \right)^2 \right] = \int_{\mathcal{X}} \frac{h(z)^2 f(z)^2}{g(z)^2} g(z) dz$$

$$= \int_{\mathcal{X}} h(z)^2 \frac{f(z)}{g(z)} f(z) dz$$

$$\leq M \int_{\mathcal{X}} h(z)^2 f(z) dz$$

$$= M E_F(h(X)^2) < \infty \quad \text{by assumption }.$$

Thus, if an accept-reject is possible for the propogal G, then a simple importance sampling estimator of θ , with a finite variance, is also possible. Now, we have a central limit theorem that can hold. Recall

$$\sigma_g^2 = \operatorname{Var}_G\left(\frac{h(Z)f(Z)}{g(Z)}\right)$$
 (1)

By the CLT, if $\sigma_g^2 < \infty$, then as $N \to \infty$,

$$\sqrt{N}(\hat{\theta}_g - \theta) \stackrel{d}{\to} N(0, \sigma_g^2)$$
. (2)

Further, an estimator of σ_g^2 is easily available since we have N samples of h(Z)f(Z)/g(Z) available. Thus, an estimator of σ_g^2 is the sample variance from all the samples:

$$\hat{\sigma}_g^2 := \frac{1}{N-1} \sum_{t=1}^N \left(\frac{h(Z_t) f(Z_t)}{g(Z_t)} - \hat{\theta}_g \right)^2.$$

Example 21 (Gamma continued). Recall from the accept-reject example for Gamma(α, β) with $\alpha \geq 1$ and Exponential(λ) proposal for an accept-reject sampler will work only if $\lambda < \beta$. That means, when $\lambda < \beta$, there exists a finite M, and the importance sampling estimator will have a finite variance.

Questions to think about

- 1. Can we construct G so that its support, \mathcal{Y} is larger than \mathcal{X} ?
- 2. Check what happens with $\beta = \lambda$ in this simulation.
- 3. Why would a CLT be useful here?
- 4. How would we check whether this importance sampler is better than IID Monte Carlo?

5.2.1 Optimal proposals

How do we choose the importance distribution g? The proposal g should be chosen so that:

- Sampling from G is relatively easy
- $\operatorname{Var}_g(\hat{\theta}_g) = \sigma_g^2/N$ is smaller than regular Monte Carlo variance estimator.

Note that, one reason to use importance sampling would be to obtain smaller variance estimators than the original. So, if we can choose g such that σ_g^2 is minimized that would be ideal!

Let's see this term:

$$\sigma_g^2 = \operatorname{Var}_G\left(\frac{h(Z)f(Z)}{g(Z)}\right) = \operatorname{E}_G\left[\frac{h(Z)^2f(Z)^2}{g(Z)^2}\right] - \theta^2 = \underbrace{\int_{\mathcal{X}} \frac{h(z)^2f(z)^2}{g(z)} dz}_{A} - \theta^2$$

For the above to be small, term A should be close to θ^2 . This logic leads to the following theorem.

Theorem 8. If $\int_{\mathcal{X}} |h(x)| f(x) dx \neq 0$, the importance density g^* that minimizes σ_g^2 is

$$g^*(z) = \frac{|h(z)|f(z)}{E_F[|h(x)|]}.$$

Proof. Consider the above importance density. The second moment of the importance sampling estimator with this density is:

$$\begin{split} \theta^2 + \sigma_{g^*}^2 \\ &= \operatorname{E}_{G^*} \left[\left(\frac{h(Z)f(Z)}{g^*(Z)} \right)^2 \right] \\ &= \int_{\mathcal{X}} \frac{h(z)^2 f(z)^2}{g^*(z)^2} g^*(z) dz \\ &= \int_{\mathcal{X}} \frac{h(z)^2 f(z)^2}{|h(z)|f(z)} \cdot \operatorname{E}_F \left[|h(x)| \right] dz \\ &= \operatorname{E}_F \left[|h(x)| \right] \int_{\mathcal{X}} |h(z)| f(z) dz \\ &= \left[\int_{\mathcal{X}} |h(z)| f(z) dz \right]^2 \\ &= \left[\int_{\mathcal{X}} \frac{|h(z)| f(z)}{g(z)} g(z) dz \right]^2 \quad \text{for any other } g \text{ defined on } \mathcal{X} \\ &= \left(\operatorname{E}_G \left[\frac{|h(z)| f(z)}{g(z)} \right] \right)^2 \\ &\leq \operatorname{E}_G \left[\frac{h(z)^2 f(z)^2}{g^2(z)} \right] \quad \text{By Jensen's inequality: for a convex function } \phi, \, \phi(E[x]) \leq E(\phi(x)) \\ &= \theta^2 + \sigma_g^2 \, . \end{split}$$

Thus, for any generic proposal q defined on \mathcal{X} , we have

$$\sigma_{g^*}^2 \le \sigma_g^2 \, .$$

Since this is true for all g, this implies that g^* produces the smallest $\sigma_{g^*}^2$.

Note that, with this choice of proposal,

$$\sigma_{g^*}^2 = \operatorname{Var}_{g^*} \left(\frac{h(Z)f(Z)}{g^*(Z)} \right)$$
$$= \operatorname{E}_F \left[|h(x)| \right]^2 \operatorname{Var}_{G^*} \left(\frac{h(Z)f(Z)}{|h(Z)|f(Z)} \right)$$

$$= \operatorname{E}_{F} \left[|h(z)| \right]^{2} \operatorname{Var}_{G^{*}} \left(\frac{h(Z)f(Z)}{|h(Z)|f(Z)} \right) .$$

If on the support \mathcal{X} , h(Z) = |h(Z)|, then the variance of the importance sampling estimator is zero!

Example 22 (Gamma distribution). Consider estimating moments of a Gamma(α, β) distribution. We actually know the optimal importance distribution here! For estimating the kth moment

$$g^*(z) \propto |h(z)|f(z)$$

$$= |x^k|x^{\alpha-1} \exp\{-\beta x\}$$

$$= x^{\alpha+k-1} \exp\{-\beta x\}.$$

So the optimum importance distribution is $Gamma(\alpha + k, \beta)$. The variance in this case of the estimator will be 0.

Example 23 (Mean of standard normal). Let h(x) = x and let f(x) be the density of a standard normal distribution. So we are interested in estimating the mean of the standard normal distribution. The universally optimal proposal in this case is

$$g^*(x) = \frac{|x|e^{-x^2/2}}{\int |x|e^{-x^2/2}dx}$$

But it may be quite challenging to draw samples from the above distribution! In order for importance sampling to be useful, we need not find the optimal proposal, as long as we can find a *more* efficient proposal than sampling from the target.

Consider an importance distribution of $N(0, \sigma^2)$ for some $\sigma^2 > 0$. The variance of the importance estimator is

$$\sigma_g^2 = \int_{-\infty}^{\infty} \frac{h(x)^2 f(x)^2}{g(x)} dx$$

$$= \int_{-\infty}^{\infty} x^2 \frac{\sigma}{\sqrt{2\pi}} \exp\left\{\frac{x^2}{2\sigma^2} - x^2\right\} dx$$

$$= \sigma \int_{-\infty}^{\infty} x^2 \frac{1}{\sqrt{2\pi}} \exp\left\{-\frac{x^2}{2} \left(2 - \frac{1}{\sigma^2}\right)\right\} dx$$

$$= \frac{\sigma}{\sqrt{2 - \sigma^{-2}}} \int_{-\infty}^{\infty} x^2 \cdot \sqrt{\frac{2 - \sigma^{-2}}{2\pi}} \exp\left\{-\frac{x^2}{2} \left(2 - \frac{1}{\sigma^2}\right)\right\} dx$$

$$= \frac{\sigma}{\sqrt{2 - \sigma^{-2}}} \frac{1}{2 - \sigma^{-2}}$$

$$= \frac{\sigma}{(2 - \sigma^{-2})^{3/2}} \quad \text{if } \sigma^2 > 1/2$$

else if $\sigma^2 < 1/2$, the integral diverges and the variance is infinite. Also, minimizing the variance:

$$\arg \min_{\sigma > \sqrt{1/2}} \frac{\sigma}{(2 - \sigma^{-2})^{3/2}} = \sqrt{2}.$$

Thus the optimal Normal proposal has standard deviation $\sigma = \sqrt{2}$, not 1! Also, at $\sigma^2 = 2$, the variance is .7698 which is less than 1.

5.2.2 Questions to think about

- Does this mean that N(0,2) is the optimal proposal for estimating the mean of a standard normal?
- What is the optimal proposal within the class of *Beta* proposals for estimating the mean of a Beta distribution?

5.3 Weighted Importance Sampling

Often for many distributions, we do not know the target distribution fully, but only know it up to a normalizing constant. That is, for some unknown a, the target density is

$$f(x) = a\tilde{f}(x)$$

and for some known or unknown b, the proposal density is

$$g(x) = b\tilde{g}(x)$$

For simplicity (or rather uniformity in complexity), we will assume that b is unknown. Even though f is not fully known, we are interested in expectations under f. Suppose for some function h, the following integral is of interest:

$$\theta := \int_{\mathcal{X}} h(x) f(x) dx.$$

We still want to use q as the importance distribution, so that

$$\theta = \int_{\mathcal{X}} \frac{h(x)f(x)}{g(x)} g(x) dx.$$

Since a and b are unknown, we can't evaluate f(x) and g(x). So our original estimator does not work anymore! We can evaluate $\tilde{f}(x)$ and $\tilde{g}(x)$. So if we can estimate a and b as well, that will allow us estimate θ . Instead, we will estimate b/a, which also works!

Consider $Z_1, \ldots, Z_t \stackrel{\text{iid}}{\sim} G$. The weighted importance sampling estimator of θ is

$$\hat{\theta}_w := \frac{\sum_{t=1}^N \frac{h(Z_t)\tilde{f}(Z_t)}{\tilde{g}(Z_t)}}{\sum_{t=1}^N \frac{\tilde{f}(Z_t)}{\tilde{g}(Z_t)}}.$$

Theorem 9. The weighted importance sampling estimator is consistent. So as $N \to \infty$, $\hat{\theta}_w \stackrel{p}{\to} \theta$.

Proof. Intuitively, both the numerator and the denominator are average, so the LLN applies to both individually. Then we may use Slutsky's theorem. This is the plan.

As a first step, note that as $N \to \infty$, by the law of large numbers

$$\frac{1}{N} \sum_{t=1}^{N} \frac{h(Z_t)\tilde{f}(Z_t)}{\tilde{g}(Z_t)} \stackrel{p}{\to} E_G \left[\frac{h(Z)\tilde{f}(Z)}{\tilde{g}(Z)} \right] \quad \text{and} \quad \frac{1}{N} \sum_{t=1}^{N} \frac{\tilde{f}(Z_t)}{\tilde{g}(Z_t)} \stackrel{p}{\to} E_G \left[\frac{\tilde{f}(Z)}{\tilde{g}(Z)} \right]$$

By an application of Slutsky's theorem, as $N \to \infty$

$$\hat{\theta}_w \stackrel{p}{\to} \frac{\mathrm{E}_G \left[\frac{h(Z)\tilde{f}(Z)}{\tilde{g}(Z)} \right]}{\mathrm{E}_G \left[\frac{\tilde{f}(Z)}{\tilde{g}(Z)} \right]}.$$

We need to show that

$$\theta = \frac{E_G \left[\frac{h(Z)\tilde{f}(Z)}{\tilde{g}(Z)} \right]}{E_G \left[\frac{\tilde{f}(Z)}{\tilde{g}(Z)} \right]}.$$

So we will first find both expectations. First

$$E_G \left[\frac{h(Z)\tilde{f}(Z)}{\tilde{g}(Z)} \right] = \int_{\mathcal{X}} \frac{h(z)\tilde{f}(z)}{\tilde{g}(z)} g(z)dz$$
$$= \frac{b}{a} \int_{\mathcal{X}} \frac{h(z)f(z)}{g(z)} g(z)dz$$
$$= \frac{b}{a}\theta.$$

Second,

$$E_G \left[\frac{\tilde{f}(Z)}{\tilde{g}(Z)} \right] = \int_{\mathcal{X}} \frac{\tilde{f}(z)}{\tilde{g}(z)} g(z) dz$$
$$= \frac{b}{a} \int_{\mathcal{X}} f(z) dz = \frac{b}{a}.$$

So,

$$\frac{E_G\left[\frac{h(Z)\tilde{f}(Z)}{\tilde{g}(Z)}\right]}{E_G\left[\frac{\tilde{f}(Z)}{\tilde{g}(Z)}\right]} = \frac{\frac{b}{a}\theta}{\frac{b}{a}} = \theta.$$

We will denote

$$w(Z) = \frac{\tilde{f}(Z)}{\tilde{g}(Z)}.$$

Then w(Z) is called the un-normalized importance sampling weight.

Thus, even though we do not know a and b, the weighted importance sampling estimator converges to the right quantity and is consistent.

However, not knowing b and a comes at a cost.

- Unlike the simple importance sampling estimator, the weighted importance sampling estimator $\hat{\theta}_w$ is not unbiased.
- Although it is challenging to show, the following asymptotic normality holds:

$$\sqrt{n}(\hat{\theta}_w - \theta) \stackrel{d}{\to} N(0, \sigma_w^2)$$

where

$$\sigma_w^2 = \frac{E_G [w(Z)(g(Z) - \theta)^2]}{(E_G [w(Z)])^2}$$

In general, we usually do not know how σ_w^2 compares with σ_g^2 or σ^2 even.

• Often, weighted importance sampling can work even better than simple importance sampling!

Example 24. Consider estimating

$$\theta = \int_0^{\pi} \int_0^{\pi} xy \pi(x, y) dx dy,$$

where

$$\pi(x,y) \propto e^{(\sin(xy))} \quad 0 \le x, y \le \pi$$

Notice that here, the target distribution is bivariate, but the function h(x,y) = xy is a univariate mapping. Further, the target distribution is not a product of two marginals, so we have to implement multivariate importance sampling. Also, we do not know the normalizing constants. So for some unknown a > 0

$$\pi(x,y) = a e^{(\sin(xy))}$$
 $0 \le x, y \le \pi$

We will use a weighted importance sampling distribution. Consider the importance distribution that is uniform on the box: $U[0, \pi] \times U[0, \pi]$. So that

$$g(z,t) = \frac{1}{\pi^2} I(0 < z < \pi) I(0 < t < \pi).$$

Since, we will assume b is unknown, we realize that

$$\tilde{g}(z,t) = 1 \cdot I(0 < z < \pi) I(0 < t < \pi) \,.$$

Sample $(Z_1, T_1), \ldots, (Z_N, T_N) \sim U[0, \pi] \times U[0, \pi]$. The weights are

$$w(Z_t, T_t) = \frac{\tilde{f}(Z_t, T_t)}{\tilde{g}(Z_t, T_t)} = e^{\sin(Z_t T_t)}.$$

The final estimator is

$$\hat{\theta}_w = \frac{\sum_{t=1}^{N} Z_t T_t w(Z_t, T_t)}{\sum_{t=1}^{N} w(Z_t, T_t)} = \frac{\sum_{t=1}^{N} Z_t T_t e^{\sin(Z_t T_t)}}{\sum_{t=1}^{N} e^{\sin(Z_t T_t)}}.$$

5.4 Questions to think about

- How would you choose a good proposal for weighted importance sampling? Would finding a proposal that yields a small variance suffice?
- Do you have intuition as to why often the variance of the weight importance estimator is larger than the variance of the simple importance sampler for the same importance proposal?

5.5 Exercises

- 1. Estimate $\int_0^1 e^x dx$ using importance sampling.
- 2. Estimate $\int_{-\infty}^{\infty} e^{-x^2/2}$ using importance sampling.
- 3. The inverse Gaussian distribution has density

$$f(x) = \sqrt{\frac{\lambda}{2\pi x^3}} \exp\left\{-\frac{\lambda(x-\mu)^2}{2\mu^2 x}\right\}$$
 $x > 0$ and $\mu, \lambda > 0$.

We are interested in estimating the moment generating function of this distribution, $E_F[e^{tX}]$ for $t \in \mathbb{R}$. Sampling from this Inverse Gaussian distribution can be quite challenging, so we will use importance sampling instead.

For $\mu = 1$ and $\lambda = 3$, using importance sampling with importance distribution Gamma(10,3) (rate = 3), make a plot with t on the x-axis and the importance sampling estimate of $E_F[e^{tX}]$ on the y-axis.

NOTE: First, use the Z_1, Z_2, \ldots, Z_N points for all chosen values of t, and then use different importance samples for different values of t.

4. Consider the problem of estimating the k-th moment of a Beta(α, β) distribution.

For which values of α and β are we sure to obtain importance estimators of the kth moment with a finite variance for a uniform proposal distribution.

- 5. In the previous problem, give other examples of importance proposal distributions that will give finite variance of the importance estimator? For what values of α and β is the finite variance of the estimator not guaranteed?
- 6. Consider estimating the mean of a standard Cauchy distribution using importance sampling with a normal proposal distribution. Does the estimator have finite variance?
- 7. For estimating kth moment of a Gamma(α, β) with $\alpha > 1$ with the importance distribution $\text{Exp}(\lambda)$, show that the importance sampling estimator has infinite variance when $\lambda > \beta$.
- 8. Considering a density f(x) and an importance proposal g(x). Suppose

$$\sup_{x} \frac{f(x)}{g(x)} < \infty.$$

In order to estimate the mean of the target density, is there any benefit to using importance sampling over accept-reject sampling?

9. For a target distribution f and a proposal g, if

$$\sup_{x} \frac{f(x)}{g(x)} < \infty$$

then we know that the simple importance estimator has finite variance. Does the weighted importance estimator also have finite variance?

10. For some known $y_i \in \mathbb{R}$, i = 1, ..., n and some $\nu > 2$, suppose the target density is

$$f(x) \propto e^{-x^2/2} \prod_{i=1}^{n} \left(1 + \frac{(y_i - x)^2}{\nu} \right)^{-(\nu+1)/2}$$
.

Generate ys using the following code for $\nu = 5$

Implement an importance sampling estimator with a N(0,1) proposal to estimate

the first moment of this distribution. Does the weighted importance sampling estimator seem to have finite variance? What happens if $\nu = 1$ and $\nu = 2$?

11. Suppose interest is in estimating

$$\theta = \int_0^{10} \exp\left\{-2|x-5|\right\} dx.$$

- What is the optimal simple importance proposal distribution? (*Hint:* look up Laplace distribution) and what is the corresponding simple importance sampling estimator?
- Implement a weighted importance sampling procedure with the same proposal distribution from above. How do the final estimators compare?

The quantity θ can be written as

$$\theta = \int_0^{10} 10 \exp\left\{-2|x-5|\right\} f(x) dx,$$

where $\pi(x)$ is the density of a U[0, 10]. We know we can do IID sampling that is, sample X_1, X_2, \ldots, X_N from Unif[0, 10] and estimate θ . But this will be simple Monte Carlo and not importance sampling. Using importance sampling, we can reduce the variance. First, note that the optimal importance distribution here is

$$g^*(z) = \frac{10 \exp\{-2|z-5|\} \frac{1}{10}}{\theta}$$
 $z \in (0,10)$.

The function h is identical to the density of a Laplace (Double exponential) distribution. For a Laplace random variable with parameters μ and b

$$l(x) = \frac{1}{2b} \exp\left\{-\frac{|z-\mu|}{b}\right\} - \infty < z < \infty$$

So h(x) is the density of Laplace(5, 1/2), but truncated between 0 and 10. So the optimum proposal is a truncated on (0, 10) Laplace(5, 1/2).

You can simulate $Z_1, Z_2, ... Z_N$ from this g^* using accept-reject algorithm (recall previous exercises) and then the optimal importance sampling estimator of θ is

$$\hat{\theta}_{g^*} = \frac{1}{N} \sum_{t=1}^{N} \frac{h(Z_t) f(Z_t)}{g^*(Z_t)} = \frac{1}{N} \sum_{t=1}^{N} \frac{10 \exp\{-2|z-5|\} \frac{1}{10}}{\frac{10 \exp\{-2|z-5|\} \frac{1}{10}}{\theta}} = \theta!$$

So notice that for the optimal proposal, the IS estimator is the quantity we want itself! Thus it is impossible to implement the simple importance sampler here.

However, we can do a weighted importance sampling estimator since the normalizing constant for g^* (which is θ) is unknown. So we have

$$g^*(z) = \underbrace{\frac{1}{\theta}}_{b} \underbrace{\exp\{-2|z-5|\}}_{\tilde{g}^*} \qquad z \in (0,10)$$

We will assume $\tilde{f} = f$, that is a = 1.

$$\hat{\theta}_w = \frac{\sum_{t=1}^{N} \frac{h(Z_t)\tilde{f}(Z_t)}{\tilde{g}^*(Z_t)}}{\sum_{t=1}^{N} \frac{\tilde{f}(Z_t)}{\tilde{g}^*(Z_t)}}$$

$$= \frac{\sum_{t=1}^{N} \frac{\exp\{-2|Z_t-5|\}}{\exp\{-2|Z_t-5|\}}}{\sum_{t=1}^{N} \frac{10^{-1}}{\exp\{-2|Z_t-5|\}}}$$

$$= 10 \frac{N}{\sum_{t=1}^{N} \exp\{2|Z_t-5|\}}$$

6 Likelihood Based Estimation

We have learned a fair amount about sampling from various distributions and estimating integrals. For the next few weeks we will focus our attention to optimization methods for certain statistical procedures.

Before we study optimization, we want to motivate why exactly optimization is useful to statisticians. One common use of optimization in statistics is when obtaining a maximum likelihood estimator (MLE) for a parameter. Thus, we first introduce MLE below briefly, before going into optimization methods.

6.1 Likelihood Function

Suppose $X_1, X_2, ..., X_n$ is a random sample from a distribution with density $f(x|\theta)$ for $\theta \in \Theta$, where Θ is the parameter space. The "x given θ " implies that given a particular value of θ , $f(\cdot|\theta)$ defines a density. $f(\cdot|\theta)$ is also written sometimes as $f_{\theta}(x)$.

The parameter θ can be a vector of parameters. After having obtained *real data*, from F, we want to

- 1. estimate θ
- 2. and assess the quality of the estimator of θ .

A useful method of estimating θ is the method of maximum likelihood estimation. Let $\mathbf{X} = (X_1, \dots, X_n)$. The idea is that we define a function $L(\theta|\mathbf{X})$ which measures "how likely is a particular value of θ given the data observed". In general $L(\theta|\mathbf{X} = \mathbf{x})$ is the joint distribution of all the Xs

$$L(\theta|\mathbf{X}=\mathbf{x}) = f(\mathbf{x}|\theta) = f(x_1, x_2, \dots, x_n|\theta).$$

When the sample is independent as well, this likelihood becomes

$$L(\theta|\mathbf{X}) = \prod_{i=1}^{n} f(x_i|\theta).$$

It is important to note that $L(\theta|\mathbf{x})$ is not a distribution over θ , it is just a function of θ . It is a function that quantifies how likely a value of θ is.

Example 25. Suppose we obtain $X_1, X_2 \stackrel{\text{iid}}{\sim} N(\theta, 1)$, where we don't know θ . Suppose

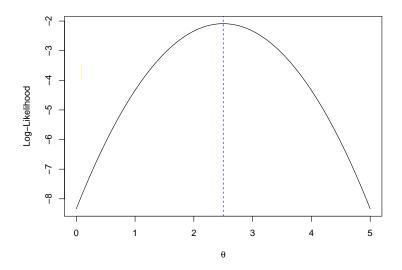
we obtain $X_1 = 2 =: x_1$ and $X_2 = 3 =: x_2$. Then the likelihood function is:

$$L(\theta|x_1, x_2) = f(x_1|\theta) \cdot f(x_2|\theta)$$

$$= f(2|\theta)f(5|\theta)$$

$$= \frac{1}{2\pi} \exp\left\{-\frac{(2-\theta)^2}{2} - \frac{(3-\theta)^2}{2}\right\}.$$

Below is the plot of the above function of θ for different values of θ to understand what the likelihood of every value of θ is.



6.2 Maximum Likelihood Estimation

The "most likely" value of θ having observed the data is the value that maximizes the likelihood

$$\hat{\theta}_{\text{MLE}} = \arg \max_{\theta \in \Theta} L(\theta | \mathbf{x}).$$

 $\hat{\theta}_{\text{MLE}}$ is called the maximum likelihood estimator of θ . As you can understand, maximizing the function $L(\theta|\mathbf{x})$ may be complex for some problems and not possible to do analytically. This is where we require numerical optimization methods to help us obtain these MLEs. MLEs have nice theoretical properties and you will learn them in MTH211a or MTH418a.

Before we continue with some examples, we recall a few definitions:

Definition 1. Concave function (one-dimension): a function h(x) is concave if $h''(x) \le 0$ for all x. If the equality never holds, it is strictly concave.

Definition 2. Concave function: a function $h(\mathbf{x})$ is concave if the Hessian matrix, $\nabla^2 h(\mathbf{x})$, is negative semi-definite for all \mathbf{x} . That is, if all eigenvalues of the Hessian are non-positive.

Example 26 (Bernoulli). Let $X_1, \ldots, X_n \stackrel{iid}{\sim} \mathrm{Bern}(p), \ 0 \le p \le 1$. Then the likelihood is

$$L(p|\mathbf{x}) = \prod_{i=1}^{n} \Pr(X_i = x_i|p)$$

$$= \prod_{i=1}^{n} \left[p^{x_i} (1-p)^{1-x_i} \right]$$

$$= p^{\sum x_i} (1-p)^{n-\sum x_i}.$$

To obtain the MLE of θ , we will maximize the likelihood. Note that maximizing the likelihood is the same as maximizing the log of the likelihood, but the calculations are easier after taking a log. So we take a log:

$$\Rightarrow l(p) := \log L(p|\mathbf{x}) = \left(\sum_{i=1}^{n} x_{i}\right) \log p + \left(n - \sum_{i=1}^{n} x_{i}\right) \log(1 - p)$$

$$\Rightarrow \frac{dl(p)}{dp} = \frac{\sum x_{i}}{p} - \frac{n - \sum x_{i}}{1 - p} \stackrel{\text{set}}{=} 0$$

$$\Rightarrow \hat{p} = \frac{1}{n} \sum_{i=1}^{n} x_{i}.$$

Taking the second derivative, we obtain

$$\frac{d^2l(p)}{dp^2} = -\frac{\sum x_i}{p^2} - \frac{n - \sum x_i}{(1 - p)^2} < 0 \qquad \text{for all } p.$$

Thus, the likelihood is concave, and \hat{p} is a global maxima.

In any example, if I do not check the second derivative, you HAVE to check it for yourself.

Thus,

$$\hat{p}_{\text{MLE}} = \frac{1}{n} \sum_{i=1}^{n} x_i.$$

6.3 Regression

We will focus a lot on variants of linear regression and thus it is important to setup the premise of linear regression.

Let Y_1, Y_2, \ldots, Y_n be observations known as the *response*. Let $x_i = (x_{i1}, \ldots, x_{ip})^T \in \mathbb{R}^p$ be the *i*th corresponding vector of covariates for the *i*th observation. Let $\beta \in \mathbb{R}^p$ be the *regression coefficient* so that for $\sigma^2 > 0$,

$$Y_i = x_i^T \beta + \epsilon_i$$
 where $\epsilon_i \sim N(0, \sigma^2)$.

Let $X = (x_1^T, x_2^T, \dots, x_n^T)^T$. In vector form we have,

$$Y = X\beta + \epsilon \sim N_n(X\beta, \sigma^2 I_n)$$
.

Note: I use capital Y to denote the population random variable and will use the small y to denote realized observations.

The linear regression model is built to estimate β , which measures the linear effect of X on Y. There is much more to linear regression and multiple courses are required to study all aspects of it. However, here we will just focus on the mathematical properties and optimization tools required to study them.

Before we continue, we will need to review some matrix-vector differentiation. For vectors $\mathbf{x}, \mathbf{a} \in \mathbb{R}^p$, and $p \times p$ matrix A, taking gradient wrt \mathbf{x}

- if A is symmetric as well, then $\nabla \mathbf{x}^T A \mathbf{x} = 2A\mathbf{x}$

Example 27 (MLE for Linear Regression). In order to understand the linear relationship between X and β , we will need to estimate β . Since we assume that the errors are normally distributed, we have a distribution available for Ys and we may use the

method of MLE. We have

$$L(\beta, \sigma^2 | y) = \prod_{i=1}^n f(y_i | X, \beta, \sigma^2)$$

$$= \left(\frac{1}{\sqrt{2\pi\sigma^2}}\right)^n \exp\left\{-\frac{1}{2} \frac{(y - X\beta)^T (y - X\beta)}{\sigma^2}\right\}$$

$$\Rightarrow l(\beta, \sigma^2) := \log L(\beta, \sigma^2 | y) = -\frac{n}{2} \log(2\pi) - \frac{n}{2} \log(\sigma^2) - \frac{1}{2} \frac{(y - X\beta)^T (y - X\beta)}{\sigma^2}$$

Note that

$$(y - X\beta)^{T}(y - X\beta) = (y^{T} - \beta^{T}X^{T})(y - X\beta)$$
$$= y^{T}y - y^{T}X\beta - \beta^{T}X^{T}y + \beta^{T}X^{T}X\beta$$
$$= y^{T}y - 2\beta^{T}X^{T}y + \beta^{T}X^{T}X\beta.$$

Using this we have (recall your multivariable calculus courses)

$$\begin{split} \frac{dl}{d\beta} &= -\frac{1}{2\sigma^2} \left[-2X^T y + 2X^T X \beta \right] = \frac{X^T y - X^T X \beta}{2\sigma^2} \stackrel{set}{=} 0 \\ \frac{dl}{d\sigma^2} &= -\frac{n}{2\sigma^2} + \frac{(y - X\beta)^T (y - X\beta)}{2\sigma^4} \stackrel{set}{=} 0 \,. \end{split}$$

The first equation leads to $\hat{\beta}_{\text{MLE}}$ satisfying

$$X^T y - X^T X \hat{\beta}_{\text{MLE}} = 0 \Rightarrow \hat{\beta}_{\text{MLE}} = (X^T X)^{-1} X^T y$$
,

if $(X^TX)^{-1}$ exists. And $\hat{\sigma}^2_{MLE}$ is

$$\hat{\sigma}_{MLE}^2 = \frac{(y - X\hat{\beta}_{MLE})^T (y - X\hat{\beta}_{MLE})}{n}$$
.

Verify: that the Hessian matrix is negative definite, and thus the objective function is concave.

Note: What if $(X^TX)^{-1}$ does not exist?

For example, if p > n, then the number of observations is less than the number of parameters, and since X is $n \times p$, (X^TX) is $p \times p$ of rank n < p. So X^TX is not full rank and cannot be inverted. In this case, the MLE does not exist and other estimators need to be constructed. This is one of the motivations of *penalized regression*, which we will discuss in detail.

6.4 Penalized Regression

Note that in the Linear regression setup, the MLE for β satisfied:

$$\hat{\beta}_{\text{MLE}} = \arg\min_{\beta} (y - X\beta)^T (y - X\beta)$$

Suppose X is such that (X^TX) is not invertible, then we don't know how to estimate β . In such cases, we may used *penalized likelihood*, that penalizes the coefficients β so that some of the β s are "pushed towards zero". The corresponding Xs to those small β s are essentially not important, removing singularity from X^TX .

Instead of looking at the likelihood, we consider a penalized likelihood. Since the optimization of $L(\beta|y)$ only depends on $(y-X\beta)^T(y-X\beta)$ term, a penalized (negative) log-likelihood is used and the final penalized (negative) log-likelihood is

$$Q(\beta) = -\log L(\beta|y) + P(\beta)$$

Here $P(\beta)$ is a penalization function. Note that since we are now looking at the *negative* log-likelihood, we now want to minimize $Q(\beta)$. The penalization function assigns large values for large β , so that the optimization problem favors small values of β .

There are many ways of penalizing β and each method yields a different estimator. A popular one is the ridge penalty.

Example 28 (Ridge Regression). The ridge penalization term is $P(\beta) = \lambda \beta^T \beta/2$ for some $\lambda > 0$ for

$$Q(\beta) = \frac{(y - X\beta)^{T}(y - X\beta)}{2} + \frac{\lambda}{2}\beta^{T}\beta.$$

We will minimize $Q(\beta)$ over the space of β and since we are adding an arbitrary term that depends on the size of β , smaller sizes of β will be preferred. Small sizes of β means X are less important, and this will eventually nullify the singularity in X^TX . The larger λ is, the more "penalization" there is for large values of β ; λ is typically user-chosen. We will study choosing λ when we cover "cross-validation" later.

We are now interested in finding:

$$\hat{\beta} = \arg\min_{\beta} \left\{ \frac{(y - X\beta)^T (y - X\beta)}{2} + \frac{\lambda}{2} \beta^T \beta \right\}.$$

To carry out the minimization, we take the derivative:

$$\frac{dQ(\beta)}{d\beta} = \frac{1}{2}(-2X^Ty + 2X^TX\beta) + \lambda\beta \stackrel{set}{=} 0$$

$$\Rightarrow (X^TX + \lambda I_p)\hat{\beta} - X^Ty = 0$$

$$\Rightarrow \hat{\beta}_{\text{ridge}} = (X^TX + \lambda I_p)^{-1}X^Ty.$$

Note: Verify that the Hessian matrix is positive definite for yourself.

Note that $(X^TX + \lambda I_p)$ is always positive definite for $\lambda > 0$ since for any $a \in \mathbb{R}^p \neq 0$

$$a^T(X^TX + \lambda I_n)a = a^TX^TXa + \lambda a^Ta > 0$$

Thus, the final ridge solution always exists even if X^TX is not invertible.

Pros:

We have an estimate of $\beta!$

In terms of a certain criterion (we will learn later), we actually do better than nonpenalized estimation even when (X^TX) is invertible.

Cons:

The estimator is not an MLE, so we cannot use distributional properties to construct confidence intervals. This is a big problem, and is addressed by bootstrapping, which we will get to.

We will study one more penalization method later.

Questions to think about

- 1. Under the normal likelihood, what is the distribution of $\hat{\beta}_{MLE}$ (when it exists) and $\hat{\beta}_{ridge}$? Are they unbiased? Which one has a smaller variance (covariance)?
- 2. What other penalization functions can you think of? Recall that $\beta^T \beta = \|\beta\|_2^2$.

6.5 No closed-form MLEs

Obtaining MLE estimates for a problem requires maximizing the likelihood. However, it is possible that no analytical form of the maxima is possible!

This is a common challenge in many models and estimation problems, and requires sophisticated optimization tools. In the next few weeks, we will go over some of these

optimization methods.

Example 29 (Gamma Distribution). Let $X_1, \ldots, X_n \stackrel{iid}{\sim} \text{Gamma}(\alpha, 1)$. The likelihood function is

$$L(\alpha|x) = \prod_{i=1}^{n} \frac{1}{\Gamma(\alpha)} x_i^{\alpha-1} e^{-x_i}$$

$$= \frac{1}{\Gamma(\alpha)^n} e^{-\sum x_i} \prod_{i=1}^{n} x_i^{\alpha-1}$$

$$\Rightarrow l(\alpha) := \log L(\alpha|x) = -n \log(\Gamma(\alpha)) + (\alpha - 1) \sum_{i=1}^{n} \log x_i - \sum_{i=1}^{n} x_i$$

Taking first derivative

$$\frac{dl(\alpha)}{d\alpha} = -n\frac{\Gamma'(\alpha)}{\Gamma(\alpha)} + \sum_{i=1}^{n} \log x_i \stackrel{set}{=} 0$$

Solving the above analytically is not possible. In fact, the form of $\frac{\Gamma'(\alpha)}{\Gamma(\alpha)}$ is challenging to write analytically.

However, taking second derivative

$$\frac{d^2 l(\alpha)}{d\alpha^2} = -n \frac{d^2}{d\alpha^2} \log(\Gamma(\alpha)) < 0 \quad \text{(polygamma function of order 1 is } > 0\text{)}$$

In the above, we use that

$$\frac{d^2}{d\alpha^2}\log(\Gamma(\alpha))$$

is the *polygamma function of order 1*, which is always positive (look it up). So we know that the function is concave and a unique maximum exists, but not available in closed form.

We cannot get an analytical form of the MLE for α . In such cases, we will use optimization methods.

6.6 Exercises

1. Two parameter exponential: The density of a two parameter exponential distribution is

$$f(x|\mu,\lambda) = \lambda e^{-\lambda(x-\mu)}$$
 $x \ge \mu$, $\mu \in \mathbb{R}, \lambda > 0$.

Compute the MLEs of both λ and μ .

The likelihood is

$$L(\lambda, \mu | \mathbf{x}) = \prod_{i=1}^{n} f(x_i | \mu, \lambda)$$

$$= \prod_{i=1}^{n} \lambda e^{-\lambda(x_i - \mu)} I(x_i \ge \mu)$$

$$= \lambda^n \exp \left\{ -\lambda \left(\sum_{i=1}^{n} x_i - n\mu \right) \right\} I(x_1, \dots, x_n \ge \mu) \quad \forall \mu \text{ and } \lambda > 0.$$

But if $X_1, \ldots, X_n \ge \mu \Rightarrow \min\{X_i\} \ge \mu$. So

$$L(\lambda, \mu | \mathbf{x}) = \lambda^n \exp \left\{ -\lambda \left(\sum_i x_i - n\mu \right) \right\} I\left(\min_i \{x_i\} \ge \mu \right) \quad \forall \mu \text{ and } \lambda > 0.$$

We will first try to maximize with respect to μ and then with respect to λ . Note that $L(\lambda, \mu | \mathbf{x})$ is an increasing function of μ within the restriction. So that the MLE of μ is the largest value in the support of μ where $\mu \leq \min\{X_i\}$. So

$$\hat{\mu}_{\text{MLE}} = \min_{1 \le i \le n} \{X_i\} =: X_{(1)}.$$

Next, note that

$$L(X_{(1)}, \lambda | \mathbf{x}) = \lambda^n \exp\left\{-\lambda \left(\sum_i X_i - nX_{(1)}\right)\right\}$$

$$\Rightarrow l(X_{(1)}, \lambda) := \log L(X_{(1)}, \lambda | \mathbf{x}) = n \log \lambda - \lambda \left(\sum_i X_i - nX_{(1)}\right)$$

$$\Rightarrow \frac{dl}{d\lambda} = \frac{n}{\lambda} - \left(\sum_i X_i - nX_{(1)}\right) \stackrel{\text{set}}{=} 0 \quad \text{and}$$

$$\frac{d^2l}{d\lambda^2} = -\frac{n}{\lambda^2} < 0.$$

So, the log-likelihood function is concave, thus there is a unique maximum. Set

$$\frac{dl}{d\lambda} = 0$$

$$\Rightarrow \frac{n}{\lambda} = \sum_{t=1}^{n} X_i - nX_{(1)}$$

$$\Rightarrow \hat{\lambda}_{\text{MLE}} = \frac{n}{\sum X_i - nX_{(1)}}.$$

2. Simple linear regression: Load the cars dataset in R:

data(cars)

Fit a linear regression model using maximum likelihood with response y being the distance and x being speed. Remember to include an intercept term in X by making the first column as a column of 1s. Do not use inbuilt functions in R to fit the model.

3. Multiple linear regression: Load the fuel2001 dataset in R:

fuel2001 <- read.csv("https://dvats.github.io/assets/fuel2001.csv",
row.names = 1)</pre>

Fit the linear regression model using maximum likelihood with response FuelC. Remember to include an intercept in X.

4. Simulating data in R:

Let $X \in \mathbb{R}^{n \times p}$ be the design matrix, where all entries in its first column equal one (to form an intercept). Let $x_{i,j}$ be the (i,j)th element of X. For the *ith* case, $x_{i1} = 1$ and x_{i2}, \ldots, x_{ip} are the values of the p-1 predictors. Let y_i be the response for the *i*th case and define $y = (y_1, \ldots, y_n)^T$. The model assumes that y is a realization of the random vector

$$Y \sim N_n(X\beta_*, \sigma_*^2 I_n)$$
,

where $\beta_* \in \mathbb{R}^p$ are unknown regression coefficients and $\sigma_*^2 > 0$ is the unknown variance.

For our simulation, let's pick $n = 50, p = 5, \sigma^2 = 1/2$ and generate the entries of β_* as p independent draws from N(0, 1):

```
set.seed(1)
n <- 50
p <- 5
sigma2.star <- 1/2
beta.star <- rnorm(p)
beta.star # to output
[1] -0.6264538  0.1836433 -0.8356286  1.5952808  0.3295078</pre>
```

We will create the design matrix $X \in \mathbb{R}^{n \times p}$, so that $x_{i1} = 1$ and the other entries are from N(0, 1).

$$X \leftarrow cbind(1, matrix(rnorm(n*(p-1)), nrow = n, ncol = (p-1)))$$

Now we will generate a realization of $Y \sim N_n(X\beta_*, \sigma_*^2 I_n)$:

In this way we have generate *simulated* data to be used in regression.

- 5. Find the MLE estimator of β and σ^2 from the previous dataset. Is it close to β_* and σ_*^2 ? Find the ridge regression solution with $\lambda = 0.01, 1, 10, 100$.
- 6. Regression: an equivalent optimization

In our original setup $X \in \mathbb{R}^{n \times p}$, all entries in its first column equal to one to form an intercept. The MLE estimate (when it exists) is

$$\hat{\beta} = \arg\min_{\beta \in \mathbb{R}^p} (y - X\beta)^T (y - X\beta).$$

Define X_{-1} be the matrix X with its first column removed. Let $\bar{y} = n^{-1} \sum_{i=1}^{n} y_i$ and $\bar{x}^T = n^{-1} 1_n^T X_{-1} = (n^{-1} \sum_{i=1}^{n} x_{i2}, \dots, n^{-1} \sum_{i=1}^{n} x_{ip})$. Let $\tilde{y} = (y_1 - \bar{y}, \dots, y_n - \bar{y})^T$ and $\tilde{X} = X_{-1} - 1_n \bar{x}^T$. Then \tilde{y} is the centered response and \tilde{X} is the centered design matrix.

Suppose that

$$\hat{\beta}_{-1} = \arg\min_{\tilde{\beta} \in \mathbb{R}^{p-1}} (\tilde{y} - \tilde{X}\tilde{\beta})^T (\tilde{y} - \tilde{X}\tilde{\beta})$$
$$\hat{\beta}_1 = \bar{y} - \bar{x}^T \hat{\beta}_{-1}.$$

Then $(\hat{\beta}_1, \hat{\beta}_{-1}^T)^T$ is equivalent to $\hat{\beta}$ above. Verify this for the dataset generated in Exercise 3.

7. Logistic Regression: Often in regression, the response may be a 0 or 1. That is, the response is a Bernoulli random variable. Let the covariate vector for the *i*th observation be $x_i = (1, x_{i2}, \dots, x_{ip})^T$. Suppose y_i is a realization of Y_i where

$$Y_i \sim \text{Bern}\left(\frac{\exp(x_i^T \beta)}{1 + \exp(x_i^T \beta)}\right).$$

Find the MLE of β_* . Does a closed form solution exist?

7 Numerical optimization methods

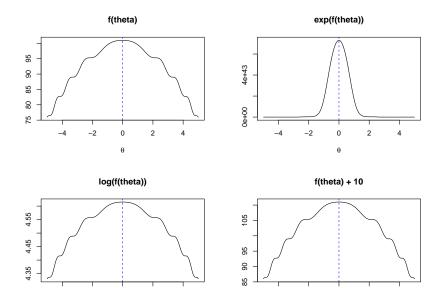
A general numerical optimization problem is framed in the following way. Let $f(\theta)$ be an *objective function* that is the main function of interest and needs to be either maximized or minimized. Then, we want to solve the following maximization

$$\theta_* = \arg\max_{\theta} f(\theta)$$

That is, we want to find that value of θ that maximizes the function f. We are often not going to be interested in the value of the function at that optimum.

This also means that we can apply transformations to the function $f(\theta)$ that retain θ_* as the argument that maximizes it.

$$\theta_* = \arg\max_{\theta} f(\theta) = \arg\max_{\theta} e^{f(\theta)} = \arg\max_{\theta} \log\{f(\theta)\} = \arg\max_{\theta} [f(\theta) + 100]$$



-2

We will cover three optimization methods:

-2

2

- Newton-Raphson method
- Gradient ascent (descent)
- The MM algorithm

All the above three algorithms are such that they generate a sequence of $\{\theta_{(k)}\}$ such that the goal is for $\theta_{(k)} \to \theta_*$ in a deterministic manner (non-random convergence).

All methods that we will learn will find a local optima. Some will guarantee a local maxima, but not guarantee a global maxima, some will guarantee a local optima (so max or min), but not a global maxima. If the objective function is concave, then all methods will guarantee a global maxima!

Recall the following:

- a (univariate) function f is concave if f'' < 0
- a (multivariate) function f is concave if its Hessian is negative definite: for all $a \neq 0 \in \mathbb{R}^{p}$,

$$a^T \left[\nabla^2 f \right] a < 0.$$

7.1 Taylor Series Approximation

For a univariate function $f(\theta)$, it's Taylor series representation around a point θ_0 is

$$f(\theta) = f(\theta_0) + f'(\theta_0)(\theta - \theta_0) + f''(\theta_0)\frac{(\theta - \theta_0)^2}{2} + f'''(\theta_0)\frac{(\theta - \theta_0)^3}{3!} + \dots$$

We will often employ the above Taylor series often. However, the infinite series representation will not be very useful. So the following approximations will be used.

Linear approximation

Ignoring all terms except the first two terms in the Taylor-series approximation gives us:

$$f(\theta) \approx f(\theta_0) + f'(\theta_0)(\theta - \theta_0)$$

Since θ_0 is just a constant, the RHS can be written as:

$$\underbrace{f(\theta_0) - f'(\theta_0)\theta_0}_{b} + \underbrace{f'(\theta_0)}_{m} \theta$$

which describes a line with intercept b and slope m. Note that when $\theta = \theta_0$, the above takes the value $f(\theta_0)$. Thus $(\theta_0, f(\theta_0))$ is on the line.

In summary, the <u>first-order</u> Taylor series approximates the function $f(\theta)$ with a line of slope $f'(\theta_0)$ passing through $(\theta_0, f(\theta_0))$.

Quadratic approximation

Ignoring all terms except the first three terms in the Taylor-series approximation gives us:

$$f(\theta) \approx f(\theta_0) + f'(\theta_0)(\theta - \theta_0) + f''(\theta_0) \frac{(\theta - \theta_0)^2}{2}$$

Since θ_0 is just a constant again, the RHS can be written as:

$$\theta^{2} \frac{f''(\theta_{0})}{2} - \theta \left[f''(\theta_{0})\theta_{0} - f'(\theta_{0}) \right] + \left[f(\theta_{0}) - f'(\theta_{0})\theta_{0} + f''(\theta_{0}) \frac{\theta_{0}^{2}}{2} \right]$$

which describes a quadratic curve. This curve also passes through the point $(\theta_0, f(\theta_0))$. In summary, the <u>second-order</u> Taylor series approximates the function $f(\theta)$ with a quadratic curve passing through $(\theta_0, f(\theta_0))$.

If $f''(\theta_0) < 0$, the quadratic is concave and if $f''(\theta_0) > 0$ the quadratic is convex.

Tayor Series' Approximation in Higher Dimensions

Suppose $f: \mathbb{R}^p \to \mathbb{R}$, that is, $\theta \in \mathbb{R}^p$ (or even some subset of \mathbb{R}^p). Recall that taking the derivative of $f(\theta)$ with respect to θ is a vector (denoted by $\nabla f(\theta)$ and the second derivative is the Hessian, which is a $p \times p$ matrix denoted by $\nabla^2 f(\theta)$.

• Linear Approximation:

$$f(\theta) = f(\theta_0) + (\theta - \theta_0)^T \nabla f(\theta_0)$$

• Quadratic Approximation:

$$f(\theta) = f(\theta_0) + (\theta - \theta_0)^T \nabla f(\theta_0) + (\theta - \theta_0)^T \nabla^2 f(\theta_0) (\theta - \theta_0)$$

7.2 Newton-Raphson's method

Recall that for a function $f(\theta)$, our goal is to solve the optimization problem:

$$\theta_* = \arg\max_{\theta} f(\theta)$$

As mentioned before, we will generate a sequence of points $\{\theta_{(k)}\}$ that will (hopefully) converge to θ_* . This process will be done sequentially. Consider a given iteration $\theta_{(k)}$ – this is just a constant number that is our current guess for θ_* . Consider a quadratic

Taylor series approximation of $f(\theta)$ around this number $\theta_{(k)}$. That is:

$$f(\theta) \approx f(\theta_{(k)}) + f'(\theta_{(k)})(\theta - \theta_{(k)}) + f''(\theta_{(k)})\frac{(\theta - \theta_{(k)})^2}{2} =: \tilde{f}_Q(\theta).$$

Given this approximation, the Newton-Raphson algorithm finds the optima of the quadratic curve $\tilde{f}_Q(\theta)$. That is, take a derivative wrt θ of $\tilde{f}_Q(\theta)$, set it zero and solve for θ . This *optima* occurs at:

$$\theta_{(k)} - \frac{f'(\theta_{(k)})}{f''(\theta_{(k)})}.$$

The Newton-Raphson method updates using iterations:

$$\theta_{(k+1)} = \theta_{(k)} - \frac{f'(\theta_{(k)})}{f''(\theta_{(k)})}$$

Algorithm 16 Newton-Raphson Algorithm

1: Choose starting value $\theta_{(0)}$ and tolerance ϵ .

2: For any
$$k$$
 find $\theta_{(k+1)} = \theta_{(k)} - \frac{f'(\theta_{(k)})}{f''(\theta_{(k)})}$

3: if $|f'(\theta_{(k+1)})| < \epsilon$ then

4: Return $\theta_{(k+1)}$ and stop

5: **else** Continue step 2

If the objective function is concave, the N-R method will converge to the global maxima. Otherwise it converges to a local optima or diverges!

Example 30 (Gamma distribution continuted). Our objective function is the log-likelihood:

$$f(\alpha) = -n\log(\Gamma(\alpha)) + (\alpha - 1)\sum_{i=1}^{n}\log x_i - \sum x_i$$

First derivative

$$f'(\alpha) = -n \frac{\Gamma'(\alpha)}{\Gamma(\alpha)} + \sum_{i=1}^{n} \log X_i$$

Second derivative

$$f''(\alpha) = -n \frac{d^2}{d\alpha^2} \log(\Gamma(\alpha)) < 0.$$

Thus the log-likelihood is concave, which implies there is a global maxima! The Newton-Raphson algorithm will converge to this global maxima.

Start with a reasonable starting value: α_0 . Then iterate with

$$\alpha_{(k+1)} = \alpha_{(k)} - \frac{f'(\alpha_{(k)})}{f''(\alpha_{(k)})}$$

Polygamma functions are in psi function in the pracma R package.

What is a good starting value α_0 ? Well, we know that the mean of a Gamma(α , 1) is α , so a good starting value is $\alpha_0 = n^{-1} \sum_{i=1}^n X_i$.

Example 31 (Location Cauchy distribution). Consider the location Cauchy distribution with mode at $\mu \in \mathbb{R}$. The goal is to find the MLE for μ .

$$f(x|\mu) = \frac{1}{\pi} \frac{1}{(1 + (x - \mu)^2)}$$
.

First, we find the log-likelihood

$$L(\mu|X) = \prod_{i=1}^{n} f(X_i|\mu) = \pi^{-n} \prod_{i=1}^{n} \frac{1}{1 + (x_i - \mu)^2}$$

$$\Rightarrow l(\mu) := \log L(\mu|X) = -n \log \pi - \sum_{i=1}^{n} \log(1 + (X_i - \mu)^2).$$

It is evident that a closed form solution is difficult. So, we find the derivates.

$$l'(\mu) = 2\sum_{i=1}^{n} \frac{X_i - \mu}{1 + (X_i - \mu)^2}.$$

$$l''(\mu) = 2\sum_{i=1}^{n} \left[2\frac{(X_i - \mu)^2}{[1 + (X_i - \mu)^2]^2} - \frac{1}{1 + (X_i - \mu)^2} \right],$$

which may be positive or negative. So this is not a concave function. This implies that Newton-Raphson is not guaranteed to converge to the global maxima! It may not even converge to a local maxima, and may converge to a local minima. Thus, we will need to be careful in choosing starting values.

- 1. Set $\mu_0 = \text{Median}(X_i)$ since the mean of Cauchy does not exist and the Cauchy centered at μ is symmetric around μ .
- 2. Determine:

$$\mu_{(k+1)} = \mu_{(k)} - \frac{f'(\mu_{(k)})}{f''(\mu_{(k)})}$$

3. Stop when $|l'(\mu_{(k+1)})| < \epsilon$ for a chosen tolerance level ϵ .

Newton-Raphson in Higher Dimensions

The NR method can be found in the same way using the multivariate Taylor's expansion. Let $\theta = (\theta_1, \theta_2, \dots, \theta_p)$. Then first let ∇f denote the gradient of f and $\nabla^2 f$ denote the Hessian. So

$$\nabla f = \begin{bmatrix} \frac{\partial f}{\partial \theta_1} \\ \vdots \\ \frac{\partial f}{\partial \theta_p} \end{bmatrix} \quad \text{and} \quad \nabla^2 f = \begin{bmatrix} \frac{\partial^2 f}{\partial \theta_1^2} & \frac{\partial^2 f}{\partial \theta_1 \theta_2} & \dots \\ \vdots & \vdots & \vdots \\ \frac{\partial^2 f}{\partial \theta_p \theta_1} & \dots & \vdots \frac{\partial^2 f}{\partial \theta_p^2} \end{bmatrix}$$

Then, the function f is concave if $\nabla^2 f$ is negative definite. So always check that first to know if there is a unique maximum.

Using a similar multivariate Taylor series quadratic expansion, the Newton-Raphson updates are as follows. If $\nabla^2 f(\theta_{(k)})$ is invertible, then you have that

$$\theta_{(k+1)} = \theta_{(k)} - \left[\nabla^2 f(\theta_{(k)}) \right]^{-1} \nabla f(\theta_{(k)}).$$

Algorithm 17 Newton-Raphson Algorithm in Higher Dimensions

- 1: Choose starting value $\theta_{(0)}$ and tolerance ϵ .
- 2: For any k find $\theta_{(k+1)} = \theta_{(k)} \left[\nabla^2 f(\theta_{(k)}) \right]^{-1} \nabla f(\theta_{(k)})$.
- 3: if $||f'(\theta_{(k+1)})|| < \epsilon$ then
- 4: Return $\theta_{(k+1)}$ and stop
- 5: **else** Continue step 2

Example 32 (Ridge regression). In the ridge regression problem, we have an analytical solution available. But assume that we did not have the analytical solution. In that case we have our objective function to *minimize*:

$$Q(\beta) = \frac{(y - X\beta)^T (y - X\beta)}{2} + \frac{\lambda}{2} \beta^T \beta.$$

We are now interested in finding:

$$\hat{\beta} = \arg\min_{\beta} \left\{ \frac{(y - X\beta)^T (y - X\beta)}{2} + \frac{\lambda}{2} \beta^T \beta \right\}.$$

Recall that we obtained

$$\nabla Q(\beta) = (X^T X + \lambda I_p)\beta - X^T y$$

and the Hessian was

$$\nabla^2 Q(\beta) = (X^T X + \lambda I_p)$$

So the iterates of Newton-Raphson are then

$$\beta_{k+1} = \beta_{(k)} - (X^T X + \lambda I_p)^{-1} ((X^T X + \lambda I_p) \beta_{(k)} - X^T y)$$

$$= \beta_{(k)} - \beta_{(k)} + (X^T X + \lambda I_p)^{-1} X^T y$$

$$= (X^T X + \lambda I_p)^{-1} X^T y$$

This simplification gives us the actual right solution! So one NR step will lead us to the analytical solution in this case! This is because $Q(\beta)$ is a quadratic function to begin with; a quadratic approximation of a quadratic is the same quadratic.

Questions

- 1. Can you implement the Newton-Raphson procedure for linear regression and ridge regression?
- 2. What are some of the issues in implementing Newton-Raphson? Can we use it for any problem?
- 3. If the function is not concave and different starting values yield convergence to

different points (or divergence), then what do we do?

7.3 Gradient Ascent (Descent)

For concave objective functions, Newton-Raphson is essentially the best algorithm. However, there are a few flaws of the algorithm that make it challenging to use it more generally:

- when the objective function is not concave, NR is not guaranteed to even converge.
- when the objective function is complicated and high-dimensional, finding the Hessian, and inverting it repeatedly may be expensive.

In such a case, gradient ascent (or gradient descent if the problem is a minimizing problem) is a useful alternative as it does not require the Hessian.

Consider the objective function $f(\theta)$ that we want to maximize and suppose θ_* is the true maximum. Then, by the Taylor series approximation at a fixed θ_0

$$f(\theta) \approx f(\theta_0) + f'(\theta_0)(\theta - \theta_0) + \frac{f''(\theta_0)}{2}(\theta - \theta_0)^2$$

If $f''(\theta)$ is unavailable or we don't want to use it, consider assuming that the double derivative is a negative constant: $f''(\theta) = -1/t$ for some t > 0. That is, assume that f is quadratic and concave. Then,

$$f(\theta) \approx f(\theta_0) + f'(\theta_0)(\theta - \theta_0) - \frac{1}{2t}(\theta - \theta_0)^2$$

Maximizing $f(\theta)$ and using this crude approximation would imply maximizing the right hand side. Taking the derivative with respect to θ setting it to zero:

$$f'(\theta_0) - \frac{\theta - \theta_0}{t} \stackrel{set}{=} 0 \Rightarrow \theta = \theta_0 + tf'(\theta_0)$$
.

Using this intuition, given a $\theta_{(k)}$, the gradient ascent algorithm does the update

$$\theta_{(k+1)} = \theta_{(k)} + tf'(\theta_{(k)}),$$

for t > 0. The iteration can be stopped when $|\theta_{(k+1)} - \theta_{(k)}| < \epsilon$ for $\epsilon > 0$ or when $|f'(\theta_{k+1})| \approx 0$.

For concave functions, there exists a t such that gradient ascent converges to the global maxima. In general (when the function is not concave), there exists a t such that gradient ascent converges to a local maxima, as long as you don't start from a local minima.

The algorithm essentially does a local concave quadratic approximation at the current point $\theta_{(k)}$ and then maximizes that quadratic equation. The value of t indicates how far do we want to jump and is a tuning parameter. If t is large, we take big jumps, as opposed to t small, where the jumps are smaller.

Example 33 (Location Cauchy distribution). Recall the location Cauchy distribution with mode at $\mu \in \mathbb{R}$, where the log-likelihood is not guaranteed to be concave and thus Newton-Raphson is difficult to use. The log-likelihood was

$$L(\mu|X) = \prod_{i=1}^{n} f(X_i|\mu) = \pi^{-n} \prod_{i=1}^{n} \frac{1}{1 + (x_i - \mu)^2}$$

$$\Rightarrow l(\mu) := \log L(\mu|X) = -n \log \pi - \sum_{i=1}^{n} \log(1 + (X_i - \mu)^2).$$

We found the first derivative:

$$l'(\mu) = 2\sum_{i=1}^{n} \frac{X_i - \mu}{1 + (X_i - \mu)^2}.$$

I choose t = 0.3 in this case so that we obtain the following gradient ascent iterative scheme:

$$\mu_{(k+1)} = \mu_{(k)} + (0.3) \left(2 \sum_{i=1}^{n} \frac{X_i - \mu}{1 + (X_i - \mu)^2} \right).$$

- 1. Set $\mu_0 = \text{Median}(X_i)$ since the mean of Cauchy does not exist and the Cauchy centered at μ is symmetric around μ .
- 2. Determine:

$$\mu_{(k+1)} = \mu_{(k)} + (0.3) \left(2 \sum_{i=1}^{n} \frac{X_i - \mu}{1 + (X_i - \mu)^2} \right).$$

3. Stop when $|l'(\mu_{(k+1)})| < \epsilon$ for a chosen tolerance level ϵ .

Gradient Ascent in Higher Dimensions

By a multivariate Taylor series expansion, we can obtain a similar motivation and the iteration in the algorithm is:

$$\theta_{(k+1)} = \theta_{(k)} + t\nabla f(\theta_{(k)}).$$

Example 34 (Logistic regression). We have studied linear regression for modeling continuous responses. But when Y is a response of 1s and 0s (Bernoulli) then assuming the Ys are normally distributed is not appropriate. Instead, when the ith covariate is $(x_{i1}, \ldots, x_{ip})^T$, then for $\beta \in \mathbb{R}^p$ logistic regression assumes the model

$$Y_i \sim \mathrm{Bern}\left(rac{e^{x_i^Teta}}{1+e^{x_i^Teta}}
ight)$$
 .

In other words, the probability that any response takes the value 1 is

$$\Pr(Y_i = 1) = \frac{e^{x_i^T \beta}}{1 + e^{x_i^T \beta}} =: p_i.$$

Our goal is to obtain the MLE of β . As usual, first we write down the log-likelihood.

$$L(\beta|Y) = \prod_{i=1}^{n} (p_i)^{y_i} (1 - p_i)^{1 - y_i}$$

$$\Rightarrow l(\beta) = \sum_{i=1}^{n} y_i \log p_i + \sum_{i=1}^{n} (1 - y_i) \log(1 - p_i)$$

$$= \sum_{i=1}^{n} \log(1 - p_i) + \sum_{i=1}^{n} y_i (\log p_i - \log(1 - p_i))$$

$$= -\sum_{i=1}^{n} \log (1 + \exp(x_i^T \beta)) + \sum_{i=1}^{n} y_i x_i^T \beta$$

In order to understand taking the derivative, we can do is elementwise for each β_s or do it using matrix calculus. Let's first do it element-wise. In order to do it element-wise, we can first solve the above

$$l(\beta) = -\sum_{i=1}^{n} \log \left(1 + \exp\left(\sum_{j} x_{ij} \beta_{j}\right) \right) + \sum_{i=1}^{n} \sum_{j=1}^{p} y_{i} x_{ij} \beta_{j}$$

$$\frac{\partial l(\beta)}{\beta_s} = \sum_{i=1}^n y_i x_{is} - \sum_{i=1}^n \frac{x_{is} e^{\sum_j x_{ij} \beta_j}}{1 + e^{\sum_j x_{ij} \beta_j}}$$

Assembling the whole vector for all j

$$\Rightarrow \nabla l(\beta) = \sum_{i=1}^{n} x_i \left[y_i - \frac{1}{1 + e^{-x_i^T \beta}} \right]$$

The above can be done directly for the full vector as well in one go:

$$\nabla l(\beta) = \sum_{i=1}^{n} x_i y_i - \sum_{i=1}^{n} \frac{x_i e^{x_i^T \beta}}{1 + e^{x_i^T \beta}}$$

$$= \sum_{i=1}^{n} x_i \left[y_i - \frac{e^{x_i^T \beta}}{1 + e^{x_i^T \beta}} \right]$$

$$= \sum_{i=1}^{n} x_i \left[y_i - \frac{1}{1 + e^{-x_i^T \beta}} \right] \stackrel{set}{=} 0$$

An analytical solution here is not possible, thus a numerical optimization tool is required. In order to know what kind of function we have, we also obtain the Hessian. In order to calculate the Hessian, we can, once again do this element-wise or do it together. Let's do it elementwise: For $j \neq k$

$$\frac{\partial^2 l(\beta)}{\beta_k \beta_s} = \frac{\partial}{\partial \beta_k} \left[\sum_{i=1}^n y_i x_{is} - \sum_{i=1}^n \frac{x_{is} e^{\sum_j x_{ij} \beta_j}}{1 + e^{\sum_j x_{ij} \beta_j}} \right]$$

$$= -\frac{\partial}{\partial \beta_k} \left[\sum_{i=1}^n \frac{x_{is} e^{\sum_j x_{ij} \beta_j}}{1 + e^{\sum_j x_{ij} \beta_j}} \right]$$

$$= -\sum_{i=1}^n x_{is} e^{\sum_j x_{ij} \beta_j} \frac{x_{ik}}{(1 + e^{\sum_j x_{ij} \beta_j})^2}$$

$$= -\sum_{i=1}^n x_{ik} x_{is} \frac{e^{\sum_j x_{ij} \beta_j}}{(1 + e^{\sum_j x_{ij} \beta_j})^2}$$

$$= -X^T W X.$$

where W is the $n \times n$ diagonal matrix with diagonal elements $e^{x_i^T \beta}/(1+e^{x_i^T \beta})^2$.

Note: You can verify by studying the Hessian above that it is negative semi-dfinite, and thus that the likelihood function is indeed concave and thus we can use either Newton-Raphson or Gradient Ascent successfully. We will use gradient ascent, and

you should on your own, implement N-R.

Gradient Ascent for Logistic regression:

- 1. Set $\beta_{(0)} = \mathbf{0}_p$ (since there is no information available, and it is reasonable to assume that none of the covariates are important).
- 2. Set some appropriate t.
- 3. For iteration k + 1:

$$\beta_{(k+1)} = \beta_{(k)} + t \left[\sum_{i=1}^{n} x_i \left[y_i - \frac{1}{1 + e^{-x_i^T \beta_{(k)}}} \right] \right]$$

4. Stop when $|\nabla f'(\beta_{(k)})| < \epsilon$.

7.4 Exercises

1. Find the MLE of (α, μ) for a Pareto distribution with density

$$f(x) = \frac{\alpha \mu^{\alpha}}{x^{\alpha+1}}$$
 $x \ge \mu$, $\mu, \alpha > 0$.

Do you need numerical procedures to calculate the MLE here?

- 2. Consider objective function of the form $f(\theta) = a\theta^2 + b\theta + c$. Writes the iterates of the Newton-Raphson algorithm. In how many iterates do you expect NR to converge for any starting value $\theta_{(0)}$?
- 3. (Using R) Find the MLE of (μ, σ^2) for the $N(\mu, \sigma^2)$ distribution using Newton-Raphson's method. Compare with the closed form estimates.
- 4. (Using R) Using both Newton-Raphson and gradient ascent algorithm, maximize objective function

$$f(x) = \cos(x) \quad x \in [-\pi, 3\pi].$$

5. Consider estimating the first moment of $\text{Exp}(\lambda)$ using simple importance sampling with a $\text{Gamma}(\alpha, \beta)$ proposal distribution. The density of an exponential is

$$\pi(x) = \lambda e^{-\lambda x} \quad x > 0 \,,$$

and the density of $Gamma(\alpha, \beta)$ is

$$g(x) = \frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha-1} e^{\beta x} \quad x > 0.$$

- (a) Construct the simple importance sampling expression for general α , β , and λ ?
- (b) Denote the variance of the estimator in (a) as $\kappa_{\lambda}(\alpha, \beta)$. For what values of α and β is $\kappa_{\lambda}(\alpha, \beta)$ infinite? You will have have to solve the integral here.
- (c) Set $\beta = 1$. Describe an algorithm for obtaining the best proposal within this family of proposals. Give all details. Use the above constructed algorithm to obtain the best $Gamma(\alpha, 1)$ for $\lambda = 5$. Here you may require an optimization algorithm.
- (d) Is the proposal obtained in (d) the universally optimal proposal for this problem?
- 6. Generate data according to the logistic regression model above with n = 50, and use Newton-Raphson's and gradient ascent algorithm to find the MLE.
- 7. Implement Newton-Raphson and gradient ascent to find the MLE of (α, β) for a Beta (α, β) distribution. Generate your own data to implement this in R.
- 8. (Using R) Find the MLE of a Gamma(α , 1) distribution using Newton-Raphson's method. Set $\alpha = 4$ and n = 10 and generate your own data. Rerun the Newton-Raphson's algorithm with different starting values.
- 9. (Using R) Find the MLE of a location Cauchy distribution with density

$$f(x) = \frac{1}{\pi} \frac{1}{1 + (x - \mu)^2} \quad \mu \in \mathbb{R}, x \in \mathbb{R}.$$

Set $\mu = -2$ and n = 4,100, and run the Newton-Raphson's algorithm with different starting values. Are starting values more impactful here than compared to the Gamma problem? Why or why not? Now repeat the same for the gradient ascent algorithm.

10. (Modified Newton-Raphson): It is possible to "overshoot" when using Newton-Raphson's algorithm, when the objective function is not concave. In these scenarios, we use a modified Newton-Raphson's approach, with step factors.

Suppose the kth iteration is such that $f'(\theta_{(k)}) > 0$ (that is the function is increasing at $\theta_{(k)}$), and NR takes us to $\theta_{(k+1)}$ where $f'(\theta_{(k+1)}) < 0$, so that now the function is decreasing. This means we may have overshot! As a compromise, we may want to implement the following algorithm:

$$\theta_{(k+1)} = \theta_{(k)} - \lambda_{(k)} \frac{f'(\theta_{(k)})}{f''(\theta_{(k)})}$$

where $\lambda_{(k)}$ is a step-factor sequence chosen at every iteration so that

$$f(\theta_{(k+1)}) > f(\theta_{(k)}).$$

That is, the next value must be such that we achieve an increase in the objective function. You can choose $\lambda_{(k)}$ so that

- (a) If $f(\theta_{(k+1)}) > f(\theta_k)$, then $\lambda_{(k)} = 1$. Else $\lambda_{(k)} = 1/2$, and recalculate $f(\theta_{(k+1)})$
- (b) If $f(\theta_{(k+1)}) > f(\theta_k)$, then continue, else set $\lambda_{(k)} = 1/2^2$ and so on...

Implement this modified algorithm for the Gamma and Cauchy examples.

11. Consider the logistic regression model: for i = 1, ..., n, let $x_i = (1, x_{i2}, ..., x_{ip})^T$ be the vector of covariates for the *i*th observation and $\beta \in \mathbb{R}^p$ be the corresponding vector of regression coefficients. Suppose response y_i is a realization of Y_i with

$$Y_i \sim \text{Bern}(p_i)$$
 where $p_i = \frac{\exp(-x_i^T \beta)}{1 + \exp(-x_i^T \beta)}$.

- (a) Write the negative log-likelihood, $-l(\beta)$.
- (b) Consider the ridge logistic regression problem, which *minimizes* the following penalized negative log-likelihood:

$$Q(\beta) = -l(\beta) + \frac{\lambda}{2} \sum_{i=1}^{p} \beta_i^2.$$

Is $Q(\beta)$ a convex function in β ?

- (c) Write the Newton-Raphson algorithm for minimizing $Q(\beta)$. Write all steps clearly.
- 12. Consider a typical Poisson regression model. For i = 1, ..., n, let $x_i \in \mathbb{R}^p$ be the vector of covariates for the *i*th observation and $\beta \in \mathbb{R}^p$ are the corresponding

regression coefficients. Let y_i be the observed response, which is count data. That is,

$$Y_i \sim \text{Poisson}\left(e^{x_i^T \beta}\right)$$
.

- (a) Find the joint likelihood $L(\beta \mid y_1, \dots y_n)$ and obtain the maximum likelihood estimator of β . If not available in closed-form, present the complete optimization algorithm to obtain $\hat{\beta}_{\text{MLE}}$.
- 13. **Probit Regression:** Logistic regression is only one way to model the probabilities. For p_i , we may actually use any mapping that will transport $x_i^T \beta$ to a probability p_i . This can be done through a CDF function.

For i = 1, ..., n, let $x_i = (1, x_{i2}, ..., x_{ip})^T$ be the vector of covariates for the *i*th observation and $\beta \in \mathbb{R}^p$ be the corresponding vector of regression coefficients. Suppose response y_i is a realization of Y_i with

$$Y_i \sim \operatorname{Bern}\left(\Phi(x_i^T\beta)\right)$$
,

where $\Phi(\cdot)$ is the CDF of a standard Normal distribution. Implement Newton-Raphson/Gradient Ascent for the Titanic dataset to find the MLE of β .

The likelihood is given by

$$L(\beta) = \prod_{i=1}^{N} \Phi(x_i^T \beta)^{y_i} (1 - \Phi(x_i^T \beta))^{(1-y_i)}.$$

We know that $1 - \Phi(x) = \Phi(-x)$. Let $z_i = 2y_i - 1$, which implies $y_i = 1 \implies z_i = 1$ and $y_i = 0 \implies z_i = -1$. This transformation will help us.

So for $y_i = 1$, $\Phi(x_i^T \beta)^{y_i} = \Phi(z_i x_i^T \beta)^{y_i}$, and for $y_i = 0$, $\Phi(x_i^T \beta)^{y_i} = 1 = \Phi(z_i x_i^T \beta)^{y_i}$. Similarly, for $y_i = 0$, $\Phi(-x_i^T \beta)^{(1-y_i)} = \Phi(z_i x_i^T \beta)^{(1-y_i)}$, and for $y_i = 1$, $\Phi(-x_i^T \beta)^{(1-y_i)} = 1 = \Phi(z_i x_i^T \beta)^{(1-y_i)}$.

Therefore,

$$\Phi(x_i^T \beta)^{y_i} = \Phi(z_i x_i^T \beta)^{y_i}$$
$$(1 - \Phi(x_i^T \beta))^{(1-y_i)} = \Phi(z_i x_i^T \beta)^{(1-y_i)}$$

As a consequence, likelihood can be written concisely as

$$L(\beta) = \prod_{i=1}^{N} \Phi(z_i x_i^T \beta)^{y_i} \Phi(z_i x_i^T \beta)^{(1-y_i)}$$
$$= \prod_{i=1}^{N} \Phi(z_i x_i^T \beta)$$

Let the log likelihood be denoted by $l(\beta)$. We have

$$l(\beta) = \sum_{i=1}^{N} \log(\Phi(z_i x_i^T \beta)).$$

We first will find the gradient vector and the Hessian matrix, to also determine whether the objective function is concave or not. Let f denote the density of a standard Normal distribution

$$\nabla l(\beta) = \frac{d}{d\beta} \sum_{i=1}^{N} \log(\Phi(z_i x_i^T \beta))$$
$$= \sum_{i=1}^{N} \frac{f(z_i x_i^T \beta) z_i x_i}{\Phi(z_i x_i^T \beta)}.$$

Next, we find the Hessian:

$$\nabla^{2}l(\beta) = \frac{d}{d\beta} \sum_{i=1}^{N} \frac{f(z_{i}x_{i}^{T}\beta)z_{i}x_{i}}{\Phi(z_{i}x_{i}^{T}\beta)}$$

$$\Rightarrow \nabla^{2}l(\beta)_{pq} = \sum_{i=1}^{N} \left[\frac{d}{d\beta_{q}} \frac{f(z_{i}x_{i}^{T}\beta)z_{i}x_{ip}}{\Phi(z_{i}x_{i}^{T}\beta)} \right]$$

$$= \sum_{i=1}^{N} \left[-\frac{f(z_{i}x_{i}^{T}\beta)^{2}}{\Phi(z_{i}x_{i}^{T}\beta)^{2}} (z_{i}x_{iq})(z_{i}x_{ip}) + \frac{f'(z_{i}x_{i}^{T}\beta)}{\Phi(z_{i}x_{i}^{T}\beta)} (z_{i}x_{iq})(z_{i}x_{ip}) \right]$$

We can use that $f'(x) = \frac{-x}{\sqrt{2\pi}}e^{-x^2/2} = -xf(x)$. This gives us the following formulation for Hessian matrix

$$\nabla^2 l(\beta)_{pq} = -\sum_{i=1}^N \left[x_{ip} \left(\frac{f(z_i x_i^T \beta)}{\Phi(z_i x_i^T \beta)} \right)^2 x_{iq} + x_{ip} \left(\frac{f(z_i x_i^T \beta)(z_i x_i^T \beta)}{\Phi(z_i x_i^T \beta)} \right) x_{iq} \right]$$

Therefore,

$$\nabla^2 l(\beta) = -\sum_{i=1}^N x_i \left[\left(\frac{f(z_i x_i^T \beta)}{F(z_i x_i^T \beta)} \right)^2 + \left(\frac{f(z_i x_i^T \beta)(z_i x_i^T \beta)}{\Phi(z_i x_i^T \beta)} \right) \right] x_i^T$$

One can show that the above is negative definite, and thus the function is concave (this will be an exercise). Now we can apply Gradient Ascent algorithm or Newton-Raphson.