

## 1 Example 1: computing the posterior for a discrete distribution

Suppose we are interested in estimating the number of different fish in a particular lake. We extract 50 fish, and mark them. A week later we extract another 50 fish, and count how many of them have marks. This approach is called "Capture-Mark-Recapture". Suppose  $X$  of them are marked, and we want to compute the posterior distribution on the number of fish.

I am going to use a prior  $p(N) \propto 1/N$  for  $N = 50, \dots, 10^6$ . Note that this is approximately uniform on the log scale, so represents being uncertain about  $N$  over orders of magnitude.

The likelihood is the number of ways of drawing 50 with  $X$  marked and  $50 - X$  unmarked, divided by the total number of ways of drawing 50 fish.

$$P(X|N) = \frac{\binom{50}{X} \binom{N-50}{50-X}}{\binom{N}{50}} \quad (1)$$

Here's example R code for  $X = 1$ :

```
N=50:10^6
X=1 # for example!
prior = 1/N
prior = prior/sum(prior)
lik = choose(N-50,50-X)/choose(N,50) #I dropped the term not depending on N
post = prior*lik
post=post/sum(post) #normalize to sum to 1
```

Now we can compute the probability that  $N$  lies in any given interval. For example

```
sum(post[N<10^4])
[1] 0.7798534
```

so our posterior probability that there are fewer than 10,000 fish is about 0.78.

## 2 Example 2: Computation by naive Monte Carlo

This example is modeled on an example from Jim Berger.

Suppose we have an imperfect test for whether someone is affected with a condition (eg a virus). Let  $f$  be the overall prevalence of the condition,  $p_0$  be the probability of a positive test result when the individual is unaffected,  $p_1$  be the probability of a positive test result if the individual is affected.

Given a positive test result the probability of being affected is

$$p_A := fp_1 / (fp_1 + (1 - f)p_0).$$

Now suppose  $f$ ,  $p_0$  and  $p_1$  are not known with certainty, but estimated based on the following pilot samples.

- For  $f$ : 1 out of 100 individuals were affected.
- For  $p_0$ : 1 out of 10 unaffected individuals gave a positive result when tested.
- For  $p_1$ : 8 out of 10 affected individuals gave a positive result when tested.

Assuming independent  $\text{Beta}(0.5, 0.5)$  priors on  $f, p_0, p_1$  the posteriors given these pilot data are also independent and  $f \sim \text{Beta}(1.5, 99.5)$ ,  $p_0 \sim \text{Beta}(1.5, 9.5)$  and  $p_1 \sim \text{Beta}(8.5, 2.5)$ .

Because it is straightforward to simulated from these posterior distributions, we can easily estimate a posterior median and 95% Credible Interval for  $p_A$  by naive Monte Carlo simulation from the posterior:

Nsim=10000



Figure 1: Histogram of posterior samples of  $p_A$  from example 2

```
f = rbeta(Nsim,1.5,99.5)
p0= rbeta(Nsim,1.5,9.5)
p1= rbeta(Nsim,8.5,2.5)
pA = f*p1/(f*p1+(1-f)*p0)
quantile(pA,c(0.025,0.5,0.975))

          2.5%          50%          97.5%
0.005374967 0.074966382 0.542589567
```

Note: the posterior of  $p_A$  (Figure ??) is sufficiently skew that the symmetric interval is not ideal here, but it is simple.

### 3 Example 3: Naive Monte Carlo vs Importance Sampling

This example is entirely artificial, to illustrate importance sampling. Throughout we assume that  $X \sim \text{Beta}(2, 2)$ . Suppose we want to estimate the probability that  $X < 0.25$ . We can do it easily by naive Monte Carlo simulation:

```
mean(rbeta(10000,2,2)<0.25)
[1] 0.1522
> pbeta(.25,2,2)
[1] 0.15625
```

The correct answer is 0.15625, so we're getting reasonable accuracy from the Monte Carlo estimate with only 10,000 simulations.

But if we want to estimate the probability that  $X < 0.001$  we have more trouble:

```
> mean(rbeta(10000,2,2)<0.0001)
[1] 0
> pbeta(0.0001,2,2)
[1] 2.9998e-08
```

Well you might argue that 0 is not a bad estimate in absolute terms, but the *relative* error is large, and sometimes we might care about this.

Here importance sampling can help. The intuition is that we need to sample from a distribution that has more mass  $< 0.0001$ . How about  $\text{Beta}(0.5, 0.5)$ ?

```
y = rbeta(10000,0.5,0.5)
w = dbeta(y,2,2)/dbeta(y,0.5,0.5)
mean(w*(y<0.0001))
```

```
[1] 2.727679e-08
sd(w*(y<0.0001))
[1] 5.434473e-07
```

which is at least the right order of magnitude. But the standard deviation of the estimator is rather large, and we can do better. Note in particular that the  $\text{Beta}(0.5, 0.5)$  spends a lot of its time sampling parts of the space where the summands ( $wI(y < 0.001)$ ) are 0. Recall, also, that the optimal IS distribution is proportional to  $\text{Beta}(2, 2)I(X < 0.0001)$ . Based on this, let's try uniform on  $(0, 0.0001)$ .

```
y = runif(10000, 0, 0.0001)
w = dbeta(y, 2, 2)/dunif(y, 0, 0.0001)
> mean(w*(y<0.0001))
[1] 2.959391e-08
> sd(w*(y<0.0001))
[1] 1.732338e-08
```

which we can see gives better accuracy and lower standard error.