

Note that you must use “MockPractical.Rmd” for your solutions so you can produce a pdf of your solutions (and associated R output).

This test contains **four** questions. The maximum mark is 50.

## Bayesian computation

1. Suppose that  $X$  follows a Cauchy distribution with density

$$f(x) = \frac{1}{\pi(1+x^2)}, \quad -\infty < x < \infty.$$

We wish to estimate the probability  $p = \mathbb{P}(X > 2)$ . We can write this as the integral

$$p = \mathbb{P}(X > 2) = \int_{-\infty}^{\infty} I(x > 2)f(x)dx$$

where  $I(x > 2)$  is the indicator function returning the value 1 if  $x > 2$  and 0 otherwise.

- 1.1 A Monte Carlo estimator of  $p$  based on  $N$  independent realisations  $X_1, \dots, X_N$  of  $X$  is

$$\hat{p}_N = \frac{1}{N} \sum_{i=1}^N I(X_i > 2).$$

Write an R function `mc` that takes a single argument `N` and returns a single realisation of  $\hat{p}_N$ . You may use the intrinsic function `rcauchy` for generating draws of  $X$ .

- 1.2 Run your `mc` function with  $N = 1000$  and compare the resulting estimate to the true value of  $p$ . Hint: `pcauchy` might be helpful here.
- 1.3 Write a function `mcMany` with arguments `N` and `n` that returns a vector of length `n` whose elements are independent realisations of  $\hat{p}_N$ . Hint: use either a `for` loop or `replicate`.
- 1.4 Using your `mcMany` function and `n=1000`, estimate the mean and variance of  $\hat{p}_N$  for  $N = 500$ . How do these compare (in terms of relative error) to the true mean and variance of  $\hat{p}_N$  for this choice of  $N$ ?
- 1.5 For  $x > 2$  the behaviour of  $f(x)$  is similar to

$$g(x) = \frac{2}{x^2}, \quad x > 2.$$

Hence, given independent draws  $X_1, \dots, X_n$  from  $g$ , the **importance sampling** estimator of  $p$  is

$$\tilde{p}_N = \frac{1}{N} \sum_{i=1}^N \frac{X_i^2}{2\pi(1+X_i^2)}.$$

- (a) Briefly explain why the indicator function is not needed in the expression for  $\tilde{p}_N$ .
- (b) The following R function takes a single argument `N` and returns a single realisation of  $\tilde{p}_N$ .

```
imp=function(N)
{
  x <- 2/runif(N)
  return((1/N)*sum(x^2/(2*pi*(1+x^2))))
}
```

Write a function `mcMany2` with arguments `N` and `n` that returns a vector of length `n` whose elements are independent realisations of  $\tilde{p}_N$ .

(c) Using your `mcMany2` function and `n=1000`, estimate the mean and variance of  $\tilde{p}_N$  for  $N = 500$ . How do these compare to the mean and variance of  $\hat{p}_N$ ?

2. A doctor is interested in the survival time of patients (in years) following a major surgery. It is proposed that the survival time,  $X$ , follows a Gamma distribution with shape parameter  $\alpha = 2$  and an unknown rate parameter  $\beta$ . Hence,  $X$  has probability density function

$$f(x|\beta) = \beta^2 \exp\{-\beta x\}, \quad x > 0.$$

Our prior distribution is  $\beta \sim \text{Gamma}(a, b)$  with  $a = b = 1$ .

Suppose that the survival times of  $m$  patients are recorded precisely as  $\mathbf{x}^o = (x_1^o, \dots, x_m^o)'$ . A further  $n - m$  patients were only monitored for  $s^* = 5$  years, i.e. their unknown survival times that exceeded  $s^* = 5$  were recorded as  $s^*$ . Denote the censored observations by  $\mathbf{x}^c = (x_1^c, \dots, x_{n-m}^c)' = (s^*, \dots, s^*)'$ . Associated with each censored observation  $x_i^c$  is a latent variable  $z_i$  representing the true survival time before censoring. Denote the latent variables by  $\mathbf{z} = (z_1, \dots, z_{n-m})'$ .

The joint posterior for  $\beta$  and  $\mathbf{z}$  is then

$$\pi(\beta, \mathbf{z}|\mathbf{x}) \propto \beta^{a-1} \exp(-b\beta) \prod_{i=1}^m \beta^2 \exp(-\beta x_i^o) \prod_{j=1}^{n-m} f(x_j^c, z_j|\beta).$$

The term  $f(x_j^c, z_j|\beta)$  is the probability density of obtaining  $z_j$  from a  $\text{Gamma}(2, \beta)$  distribution *and* that it exceeds  $s^*$ ; that is,

$$\beta^2 z_j \exp(-\beta z_j) I(z_j > s^*)$$

where the indicator  $I(z_j > s^*)$  takes the value 1 if  $z_j > s^*$  and 0 otherwise.

- 2.1 Using the *inverse sampling* method, write a function called `gammaT` that takes arguments `N`, `sstar` and `beta` and returns `N` draws from a left truncated (at `sstar`)  $\text{Gamma}(2, \beta)$  distribution. Hint: the functions `pGamma(x, a, b)` and `qGamma(x, a, b)` evaluate the cdf and inverse cdf (resp.) of a Gamma random variable with shape and rate parameters given by `a` and `b`, at the value `x`.

- 2.2 Generate a (synthetic) data set by running the following R code:

```
set.seed(3421)
data <- rgamma(100, 2, 0.5)
#Truncate anything bigger than 5
data[data > 5] <- 5
data <- sort(data) #x=(x^o, x^c)
```

How many precise observations are there?

- 2.3 You are given the full conditional distributions

$$\beta|\mathbf{x}, \mathbf{z} \sim \text{Gamma}\left(a + 2n, b + \sum_{i=1}^m x_i^o + \sum_{j=1}^{n-m} z_j\right)$$

$$Z_j | \mathbf{x}, \beta, \mathbf{z}_{-j} \sim \text{Gamma}_{(z_j > s^*)}(2, \beta)$$

where  $\text{Gamma}_{(z_j > s^*)}(2, \beta)$  denotes a gamma density truncated at the left at  $s^*$ .

Complete the following R function which takes as arguments the number of iterations  $N$ , a data set  $\mathbf{x}$  and the left truncation limit  $sstar$ , and runs a Gibbs sampler to generate draws from the posterior  $\pi(\beta, \mathbf{z} | \mathbf{x})$ .

```
gibbs = function(N,x,sstar)
{
  a <- 1; b <- 1 #prior hyper-parameters
  n <- length(x)
  m <- #WRITE YOUR SOLUTION
  x0 <- x[1:m] #observed survival times
  betaVec <- rep(0,N) #store beta samples here
  zMat <- matrix(0,nrow=N,ncol=(n-m)) #store z samples here
  #Initialise
  beta <- #WRITE YOUR SOLUTION
  z <- rep(sstar,n-m)
  betaVec[1] <- beta; zMat[1,] <- z
  for(i in 2:N)
  {
    #update beta
    beta <- #WRITE YOUR SOLUTION
    #update z
    z <- #WRITE YOUR SOLUTION
    #store
    betaVec[i] <- beta; zMat[i,] <- z
  }
  return(list(betaVec,zMat))
}
```

- 2.4 By running the sampler for 5,000 iterations, briefly investigate the mixing of the  $\beta$  chain and the  $\mathbf{z}$  component chains. Remove any necessary burn-in, produce a kernel density estimate of the marginal  $\beta$  posterior and compare it to the ground truth value of  $\beta$  that generated the data.

## Bayesian modelling

3. We have been given two competing models:

$$\begin{aligned}
 \mathcal{M}_1 : X | \mu, \phi &\sim \text{NegBin}(\mu, \phi), \\
 \mu &\sim \text{N}(5, 4) \text{ truncated to have support } [0, \infty], \\
 \phi &\sim \text{Gamma}(2, 2); \\
 \mathcal{M}_2 : X | \mu, \phi &\sim \text{NegBin}(\mu, \phi), \\
 \mu &\sim \text{N}(5, 4) \text{ truncated to have support } [2, 10], \\
 \phi &\sim \text{Gamma}(2, 2) \text{ truncated to have support } [10, \infty].
 \end{aligned}$$

Note that, with this parameterisation of the negative binomial distribution, we have

$$E(X | \mu, \phi) = \mu \quad \text{and} \quad \text{Var}(X | \mu, \phi) = \mu + \mu^2 / \phi.$$

We receive the following data:

$$\underline{x} = \{1, 3, 5, 6, 5, 2, 8, 2, 9, 7, 3, 5, 5, 1, 8, 1, 8, 4, 1, 0\}.$$

- 3.1 Write some model code for Stan that covers each of the models (with two separate Stan files).
- 3.2 Write out R code to sample from the posterior for each model, and estimate a 95% credible interval for each of the model parameters.
- 3.3 Use the `loo` package to find pseudo-BMA weights for the models.

4. We have data about average student-to-teacher ratios for seven- and eight-year-olds for a number of towns, across different districts, for two different regions. If we let  $R_{ijk}$  be the average ratio for the  $i$ th town from the  $j$ th district for the  $k$ th region, we have the following model:

$$\begin{aligned}
 R_{ijk} | \mu_{jk}, \sigma_k^2 &\sim \text{log-N}(\mu_{jk}, \sigma_k^2), \\
 \sigma_k^2 &\sim \text{Exp}(\lambda), \\
 \lambda &\sim \text{Gamma}(5, 1), \\
 \mu_{jk} &\sim \text{N}(\beta_k, 1/\nu), \\
 \beta_k &\sim \text{N}(\alpha, 1/\tau), \\
 \nu &\sim \text{Exp}(3), \\
 \tau &\sim \text{Exp}(1) \\
 \alpha &\sim \text{???}.
 \end{aligned}$$

where  $i = 1, \dots, n_{jk}$ ,  $j = 1, \dots, n_k$ ,  $k = 1, 2$ , and the prior distribution for  $\alpha$  has not yet been determined.

- 4.1 Using the bisection method for elicitation of expert knowledge, we elicit judgements of 2.3, 3 and 3.85 for the 25th, 50th and 75th percentiles for  $\alpha$  respectively. Use <https://jeremy-oakley.shinyapps.io/SHELF-single/> to determine a suitable prior distribution for  $\alpha$  using the elicited judgements.
- 4.2 The collected data have been recorded in "Mock\_practical.Rmd". How many distinct districts do we have data for?
- 4.3 Write model code for Stan that covers the given model and generates a sample from the predictive distribution for the ratio for an unsampled town in the second district in the first region.
- 4.4 Find the posterior means and standard deviations for all the model parameters.
- 4.5 Which town is best and which region is best overall in terms of having the lowest student-to-teacher ratio? Justify your answers.