

## Homework 3

### Problem 1

#### 1a

For a proportionality constant  $k$  we can write:

$$\pi(x) = k \cdot x^{\alpha-1} (1-x)^{\beta-1}$$

$$\log \pi(x) = \log k + (\alpha - 1) \log x + (\beta - 1) \log(1 - x)$$

To use adaptive rejection sampling, we need the first derivative of  $\log \pi(x)$  to exist and the second derivative to be non positive.

$$\log \pi'(x) = \frac{\alpha-1}{x} - \frac{\beta-1}{1-x}$$

$$\log \pi''(x) = -\frac{\alpha-1}{x^2} - \frac{\beta-1}{(1-x)^2}$$

As we can see,  $\log \pi'(x)$  exists for  $0 < x < 1$ . Let's see which values of  $\alpha$  and  $\beta$  make  $\log \pi''(x)$  non positive.

$$\log \pi''(x) \leq 0 \quad \text{for all } 0 < x < 1$$

$$-\log \pi''(x) = \frac{\alpha-1}{x^2} + \frac{\beta-1}{(1-x)^2} \geq 0 \quad \text{for all } 0 < x < 1$$

Let's study two border cases first.

$$\lim_{x \rightarrow 0} -\log \pi''(x) = \lim_{x \rightarrow 0} \frac{\alpha-1}{x^2} + \frac{\beta-1}{(1-x)^2} = \lim_{x \rightarrow 0} \frac{\alpha-1}{x^2} + \frac{\beta-1}{1^2} = \lim_{x \rightarrow 0} \frac{\alpha-1}{x^2} \geq 0$$

Resulting in  $\alpha \geq 1$ .

$$\lim_{x \rightarrow 1} -\log \pi''(x) = \lim_{x \rightarrow 1} \frac{\alpha-1}{x^2} + \frac{\beta-1}{(1-x)^2} = \lim_{x \rightarrow 1} \frac{\alpha-1}{1^2} + \frac{\beta-1}{(1-x)^2} = \lim_{x \rightarrow 1} \frac{\beta-1}{(1-x)^2} \geq 0$$

Resulting in  $\beta \geq 1$ .

From these border cases we see, that  $\alpha \geq 1$  and  $\beta \geq 1$ . It is also easy to see, that these constraints on  $\alpha$  and  $\beta$  make term  $-\log \pi''(x) = \frac{\alpha-1}{x^2} + \frac{\beta-1}{(1-x)^2}$  non-negative for all values of  $0 < x < 1$ .

Therefore,  $\log \pi''(x)$  is non positive for  $\alpha \geq 1$  and  $\beta \geq 1$ .

Answer: adaptive rejection sampling can be used for  $\alpha \geq 1$  and  $\beta \geq 1$ .

#### 1b

```
library(ars)
set.seed(42)
```

```

# log pi(x)
f <- function(x, a, b){
  return((a-1) * log(x) + (b-1) * log(1-x))
}

# log pi'(x)
fprima <- function(x, a, b){
  return((a-1) / (x) - (b-1) / (1-x))
}

```

```

R = 10^5
a = 2
b = 6

gen_beta <- ars(R,f,fprima, x=c(0.3,0.6),
               m=2, lb=TRUE, xlb=0, ub=TRUE, xub=1,
               a=a ,b=b)

```

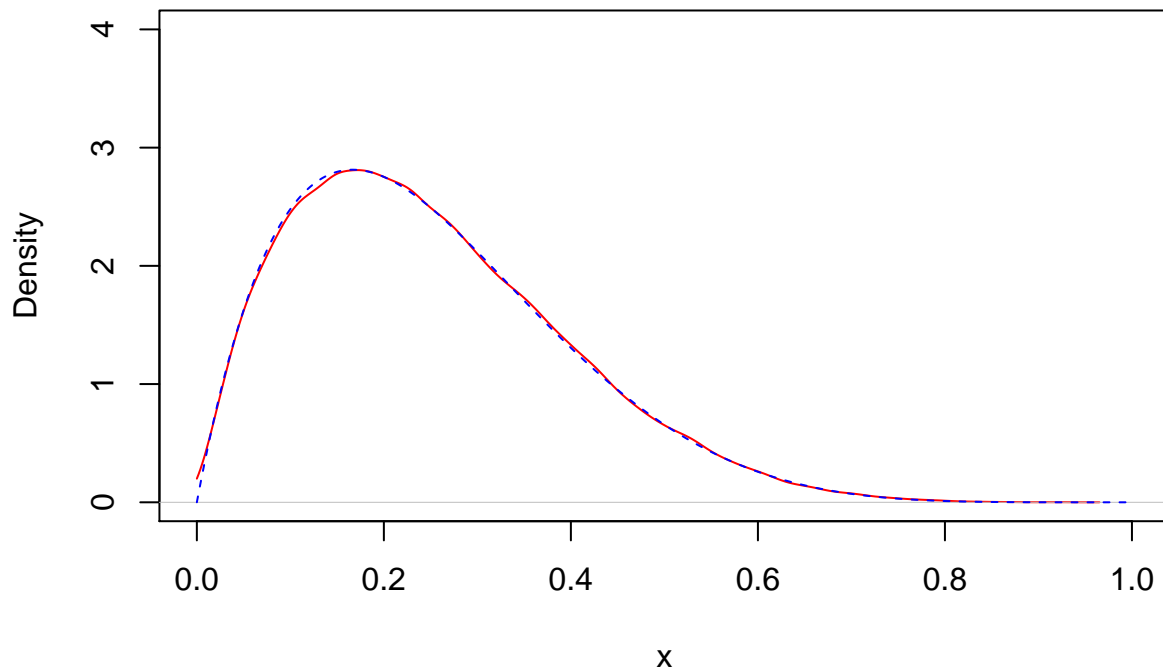
```

x <- seq(0, 1, length=1000)

plot(density(gen_beta, from=0), xlim=c(0,1), ylim=c(0,4), col='red', xlab="x",
     main="Estimated kernel density of our implementation G(0.5,1) - red,
          and true density of G(0.5,1) - blue")
lines(x, dbeta(x, a, b),col='blue', lty=2)

```

**Estimated kernel density of our implementation G(0.5,1) – red,  
and true density of G(0.5,1) – blue**



## Problem 2

Let  $I$  denote if individual has HIV and  $T$  denoting if the test is positive. Then:

$$\begin{aligned} P(T = 1|I = 1) &= 0.99 \\ P(T = 0|I = 0) &= 0.98 \\ P(I = 1) &= 0.001 \end{aligned} \tag{1}$$

### 2.a

$P(I = 1|T = 1)$ —?

$$\begin{aligned} P(I = 1|T = 1) &\stackrel{\text{Bayes Theorem}}{=} \frac{P(T=1|I=1) \cdot P(I=1)}{P(T=1)} \\ P(I = 0) &= 1 - P(I = 1) = 0.999 \\ P(T = 1) &= P(T = 1|I = 1) \cdot P(I = 1) + P(T = 1|I = 0) \cdot P(I = 0) = \\ &= 0.99 \cdot 0.001 + 0.02 \cdot 0.999 = 0.02097 \\ P(I = 1|T = 1) &= \frac{0.99 \cdot 0.001}{0.02097} \approx 0.04721 = 4.721\% \end{aligned} \tag{2}$$

### 2.b

$\text{percentage}(I = 0|T = 1)$ —?

$$\text{percentage}(I = 0|T = 1) = P(I = 0|T = 1) = 1 - P(I = 1|T = 1) \approx 0.95279 = 95.279\% \tag{3}$$

## Problem 3

### 3.a

For i.i.d. data we can write likelihood as follows:

$$\begin{aligned} \ell(\lambda) &= f(\mathbf{y}|\lambda) \stackrel{i.i.d.}{=} \prod_{i=1}^n f(y_i|\lambda) \\ &= \prod_{i=1}^n \frac{\lambda^{y_i} e^{-\lambda}}{y_i!} \\ &\propto \prod_{i=1}^n \lambda^{y_i} e^{-\lambda} \\ &= e^{-\lambda n} \cdot \lambda^{\sum_{i=1}^n y_i} \end{aligned} \tag{4}$$

Then the maximum likelihood estimate is:

$$\begin{aligned} \hat{\lambda}_{MLE} &= \arg \max_{\lambda} \ell(\lambda) = \arg \max_{\lambda} \log \ell(\lambda) \\ &= \arg \max_{\lambda} \left[ -n \cdot \lambda + \log \lambda \cdot \sum_{i=1}^n y_i \right] \end{aligned} \tag{5}$$

$$\begin{aligned}
\frac{d}{d\lambda} \log \ell(\lambda) &= -n + \frac{1}{\lambda} \sum_{i=1}^n y_i \stackrel{!}{=} 0 \\
\lambda &= \frac{\sum_{i=1}^n y_i}{n} \\
\hat{\lambda}_{MLE} &= \frac{\sum_{i=1}^n y_i}{n}
\end{aligned} \tag{6}$$

### 3.b

$$\begin{aligned}
p(\lambda) &= G(\alpha, \beta) = \frac{\beta^\alpha \cdot \lambda^{\alpha-1} \cdot e^{-\beta\lambda}}{\Gamma(\alpha)} \\
&\propto \lambda^{\alpha-1} e^{-\beta\lambda}
\end{aligned} \tag{7}$$

$$\begin{aligned}
p(\lambda|\mathbf{y}) &= \frac{\ell(\lambda) \cdot p(\lambda)}{f(\mathbf{y})} = \\
&\propto \ell(\lambda) \cdot p(\lambda) \\
&\propto \lambda^{\alpha-1} \cdot e^{-\beta\lambda} \cdot e^{-n\lambda} \cdot \lambda^{\sum_{i=1}^n y_i} \\
&= \lambda^{(\alpha-1+\sum_{i=1}^n y_i)} \cdot e^{-\lambda(\beta+n)} \\
&\propto G\left(\alpha + \sum_{i=1}^n y_i, \beta + n\right)
\end{aligned} \tag{8}$$

As we can see, posterior is Gamma-distributed with  $\alpha_{post} = \alpha + \sum_{i=1}^n y_i$  and  $\beta_{post} = \beta + n$ .

### 3.c

```
lambda <- 5

n <- 10^3

a_prior <- 70
b_prior <- 10

n_1 <- 10
n_2 <- 100
n_3 <- 1000
```

```
set.seed(42)
samples <- rpois(n, lambda)
```

```
ab_posterior <- function(a_prior, b_prior, y){
  a_post = a_prior + sum(y)
  b_post = b_prior + length(y)
  c(a_post, b_post)
}

mle_lambda <- function(y){
  return(mean(y))
}
```

```

res_1 <- ab_posterior(a_prior, b_prior, samples[1:n_1])
res_2 <- ab_posterior(a_prior, b_prior, samples[1:n_2])
res_3 <- ab_posterior(a_prior, b_prior, samples[1:n_3])
a_post_1 <- res_1[1]
b_post_1 <- res_1[2]
a_post_2 <- res_2[1]
b_post_2 <- res_2[2]
a_post_3 <- res_3[1]
b_post_3 <- res_3[2]
mle_1 <- mle_lambda(samples[1:n_1])
mle_2 <- mle_lambda(samples[1:n_2])
mle_3 <- mle_lambda(samples[1:n_3])

```

```

x <- seq(4, 10, length=1000)

plot(x, dgamma(x, a_prior, b_prior), ylim=c(0,4), ylab="density",
      lwd=1, col='red', type="l")
title("
True lambda - black
Prior - red
Posterior and MLE after 10 obs - blue
Posterior and MLE after 100 obs - orange
Posterior and MLE after 1000 obs - green
", line = -5, cex.main = 0.7,
)
lines(x, dgamma(x, a_post_1, b_post_1), lwd=1, col='blue')
lines(x, dgamma(x, a_post_2, b_post_2), lwd=1, col='darkorange2')
lines(x, dgamma(x, a_post_3, b_post_3), lwd=1, col='darkgreen')
abline(v=lambda, lty=2, lwd=2, col=c("black"))

abline(v=mle_1, lty=2, lwd=1, col=c("blue"))
abline(v=mle_2, lty=2, lwd=1, col=c("darkorange2"))
abline(v=mle_3, lty=2, lwd=1, col=c("darkgreen"))

```

