Exercise 1

TMA4300 Computer Intensive Statistical Models

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08 februar, 2021

Problem A: Stochastic simulation by the probability integral transform and bivariate techniques

1.

2.

(a)

(b)

3.

(a)

(b)

(c)

4.

5

Problem B: The gamma distribution

1.

(a)

(b)

2.

(a)

(b)

3.

(a)

(b)

4.

5.

(a)

(b)

Problem C: Monte Carlo integration and variance reduction

1.

2.

3.

(a)

(b)

Problem D: Rejection sampling and importance sampling

Subproblem 1.

We consider a vector of multinomially distributed counts $\mathbf{y} = \begin{bmatrix} y_1 & y_2 & y_3 & y_4 \end{bmatrix}^\top$ and the observed data is $\mathbf{y} = \begin{bmatrix} 125 & 18 & 20 & 34 \end{bmatrix}^\top$. The multinomial mass function is given as

and assuming a prior that is Uniform$(0,1)$ the posterior will be

$$f(\theta \mid \mathbf{y}) \propto f^*(\theta) := (2+\theta)^{y_1}(1-\theta)^{y_2+y_3}\theta^{y_3},$$

for $\theta \in (0,1)$. We wish to sample from this using a Uniform$(0,1)$ proposal density, that is, $g(\theta \mid \mathbf{y}) = 1$, for $\theta \in (0,1)$. To do a rejection sampling (not weighted rejection sampling), we need to know the normalizing constant of $f(\theta \mid \mathbf{y})$. That is, the constant k such that $f(\theta \mid \mathbf{y}) = kf^*(\theta \mid \mathbf{y})$. This can be found as

$$\frac{1}{K} = \int_{\mathbb{R}} f^*(\theta \mid \mathbf{y})\,\mathrm{d}\theta = \int_0^1 f^*(\theta \mid \mathbf{y})\,\mathrm{d}\theta \approx 2.3577 \cdot 10^{28},$$

and we find it using the `integrate()`-function in `R` below. To use the rejection sampling we also need that

$$\frac{f(\theta \mid \mathbf{y})}{g(\theta \mid \mathbf{y})} = f(\theta \mid \mathbf{y}) \leq k,$$

and a value for k is found in the code block below. We then simulate $\Theta \sim$ Uniform$(0,1)$ and $U \sim$ Uniform$(0,1)$ and calculate $\alpha = f(\theta \mid \mathbf{y})/k$. Then, if $U \leq \alpha$, Θ is returned, and if not, the procedure is run again. We then sample from the posterior distribution in the code block below.

```
y <- c(125, 18, 20, 34)    # Observed data

# Define the un-normalized posterior distribution f*(theta | y)
posterior_star <- function(theta, y) {
  return((2 + theta)^(y[1]) * (1 - theta)^(y[2] + y[3]) * theta^(y[4]))
}

# Find the normalizing constant 1 / K
norm_const <- integrate(function(theta)(posterior_star(theta, y)),
                        lower = 0,
                        upper = 1)$value

# Defining the normalized posterior distribution f(theta | y)
posterior <- function(theta, y) {
  return(posterior_star(theta, y) / norm_const)
}

# Finding the maximum
posterior_star_max <- optimize(function(theta)(posterior_star(theta, y)),
                               interval = c(0, 1),
                               maximum = TRUE)$objective

# k such that f(theta | y) <= k
k <- posterior_star_max / norm_const

# Rejection sampling algorithm
rejection_sampling <- function(M, y) {
  count <- 0       # Count how many times the algorithm runs
  accept <- c()    # List of the accepted samples

  while(length(accept) < M) {
    U <- runif(1)
    Theta <- runif(1)
    alpha <- posterior(Theta, y) / k
    if(U <= alpha) {
      accept <- rbind(accept, Theta)
```

```
    }
    count <- count + 1
  }
  return(list("accept" = accept, "co" = count))
}
```

Subproblem 2.

Drawing $\Theta_1, \ldots, \Theta_M \sim f(\theta \mid \mathbf{y})$, the Monte Carlo estimate of $\mu = \mathrm{E}(\theta \mid \mathbf{y})$ is

$$\hat{\mu} = \frac{1}{M} \sum_{i=1}^{M} \Theta_i.$$

We do this for $M = 10000$ in the code block below. Figure 1 shows the result of this. We see the estimation of the posterior mean $\mathrm{E}(\theta \mid \mathbf{y})$ using Monte Carlo integration and numerical integration together with the theoretical posterior density distribution and a generated histogram of the samples. In the figure the posterior density is plotted using a normalizing constant we find by numerical integration in `R` below, giving the normalizing constant `norm_const`.

```
M <- 10000    # Number of samples from f(theta | y)

Theta_samp <- rejection_sampling(M, y)   # M samples from f(theta | y)
mu_est <- mean(Theta_samp$accept)    # = 1/M * sum(Theta_samp)

mu_num <- integrate(function(theta)(theta * posterior(theta, y)),
                    lower = 0,
                    upper = 1)$value    # Value of mu by numerical integration

# Plot
ggplot() +
  geom_histogram(
    data = as.data.frame(Theta_samp$accept),
    mapping = aes(x = Theta_samp$accept, y = ..density..),
    binwidth = 0.01,
    boundary = 0
  ) +
  stat_function(
    fun = posterior,
    args = list(y = y),
    aes(col = "Posterior density")
  ) +
  geom_vline(
    aes(xintercept = c(mu_est, mu_num),
        col = c("Estimated posterior mean", "Numerical posterior mean"),
        linetype = c("dashed", "dotted"))
  ) +
  guides(linetype = FALSE) +    # Remove linetype from label
  ggtitle("Estimation of the posterior mean") +
  xlab("theta") +
  ylab("Density") +
  theme_minimal() +
  theme(plot.title = element_text(hjust = 0.5)) +
  theme(legend.title = element_blank())
```

In the following code block we find the values of `mu_est` and `mu_num`.

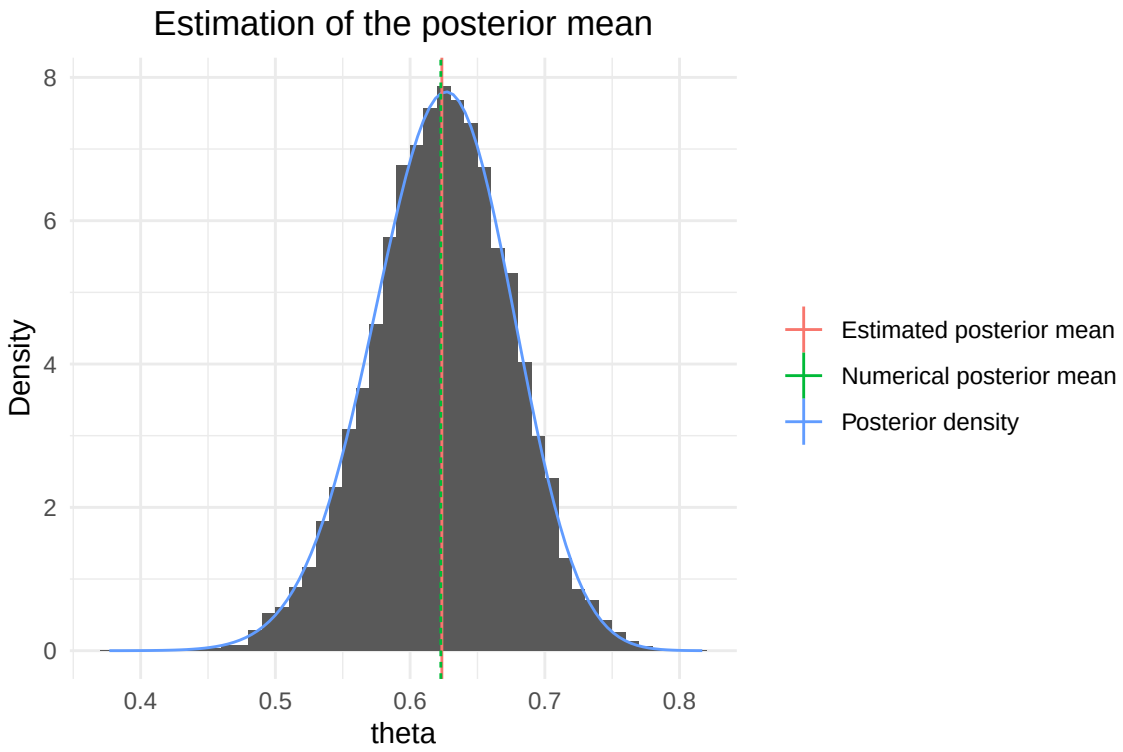

Figure 1: Estimation of the posterior mean $\mathrm{E}(\theta \mid \mathbf{y})$ using Monte Carlo integration and numerical integration. A histogram of the samples is also shown together with the theoretical posterior density distribution.

```
mu_est
```

```
## [1] 0.6235803
```

```
mu_num
```

```
## [1] 0.6228061
```

From this it is clear that the estimated posterior mean is $\hat{\mu} \approx 0.623$ using Monte Carlo integration, and $\mu \approx 0.623$ using numerical integration with `integrate()`. Figure 1 also shows that these means corresponds well to the real posterior mean.

Subproblem 3.

We are now interested in the number of random numbers the sampling algorithm needs to obtain one sample from $f(\theta \mid \mathbf{y})$. The expected number of trials up to the first sample from $f(\theta \mid \mathbf{y})$ is c given by the condition

$$\frac{f(\theta \mid \mathbf{y})}{g(\theta \mid \mathbf{y})} = f(\theta \mid \mathbf{y}) \leq c.$$

We may then choose

$$c \geq \max_{\theta \in [0,1]} f(\theta \mid \mathbf{y}),$$

and we choose the equality. Thus we may find c in `R` using the `optimize()`-function, and call this `const_num` to symbolize that this is the numerically calculated value, as in the following code block.

```
const_num <- optimize(function(theta)(posterior_f(theta, y)),
                      interval = c(0, 1),
```

```
                        maximum = TRUE)$objective
const_num
```

```
## [1] 7.799308
```

Using the sampler, the expected number of random numbers that has to be generated in order to obtain one sample of $f(\theta \mid \mathbf{y})$ is given in the following code block.

```
Theta_samp$co / M
```

```
## [1] 7.8237
```

Subsection 4.

We now assume a Beta$(1, 5)$ prior

$$\tilde{f}(\theta) = \frac{1}{\mathrm{B}(1,5)}(1-\theta)^4 \propto (1-\theta)^4,$$

where $\mathrm{B}(\cdot, \cdot)$ is the beta function. This gives the posterior

$$\tilde{f}(\theta \mid \mathbf{y}) \propto f(\mathbf{y} \mid \theta)\tilde{f}(\theta) \propto \tilde{f}^*(\theta \mid \mathbf{y}) := (2+\theta)^{y_1}(1-\theta)^{y_2+y_3+4}\theta^{y_4}.$$

We wish to estimate μ by importance sampling, and to avoid needing to know the normalizing constant, we use the self-normalizing importance sampling estimator

$$\tilde{\mu}_{\mathrm{IS}} = \frac{\sum_{i=1}^n \Theta_i w(\Theta_i)}{\sum_{i=1}^n w(\Theta_i)},$$

where

$$w(\Theta_i) = \frac{\tilde{f}^*(\Theta_i \mid \mathbf{y})}{f^*(\Theta_i)} = (1-\Theta_i)^4.$$

In the limit $n \to \infty$, the estimator $\tilde{\mu}_{\mathrm{IS}}$ is unbiased. Here $\Theta_1, \ldots, \Theta_n \sim f(\theta \mid \mathbf{y})$.

```
set.seed(69)

posterior_tilde_star <- function(theta, y) {
  return((2 + theta)^(y[1]) * (1 - theta)^(y[2] + y[3] + 4) * theta^(y[4]))
}

importance_sampling <- function(n, y) {
  Theta <- rejection_sampling(n, y)$accept
  w <- (1 - Theta)^4
  importance_mean <- sum(Theta * w) / sum(w)
  return(importance_mean)
}

# Test
n <- 10000
y <- c(125, 18, 20, 34)   # Observed data

mu_is <- importance_sampling(n, y)
mu_is
```

```
## [1] 0.5947878
```

```r
# True mean
const <- integrate(function(theta)(posterior_tilde_star(theta, y)),
                   lower = 0,
                   upper = 1)$value
true_mu <- integrate(function(theta)(theta * posterior_tilde_star(theta, y) / const),
                     lower = 0,
                     upper = 1)$value
true_mu
```

```
## [1] 0.5959316
```

We then see that the estimated $\tilde{\mu}_{\text{IS}} \approx 0.5948$, while the true posterior mean is $\mu \approx 0.5959$. We notice that $\tilde{\mu}_{\text{IS}} < \hat{\mu}$, as found in Subproblem 2. This can be explained by looking at the priors. The previous prior $\text{Beta}(1, 1)$ and the new prior $\text{Beta}(1, 5)$ are shown in Figure 2. From this it is clear that the $\text{Beta}(1, 5)$ prior favors lower values for θ compared to the uniform prior, which do not favor any particular θ. It is therefore expected that $\tilde{\mu}_{\text{IS}} < \hat{\mu}$, which is shown to be true.

```r
ggplot() +
  stat_function(
    fun = dbeta,
    args = list(shape1 = 1, shape2 = 1),
    aes(col = "Beta(1, 1)")
  ) +
  stat_function(
    fun = dbeta,
    args = list(shape1 = 1, shape2 = 5),
    aes(col = "Beta(1, 5)")
  ) +
  ggtitle("The different priors") +
  xlab("theta") +
  ylab("Density") +
  theme_minimal() +
  theme(plot.title = element_text(hjust = 0.5)) +
  theme(legend.title = element_blank())
```

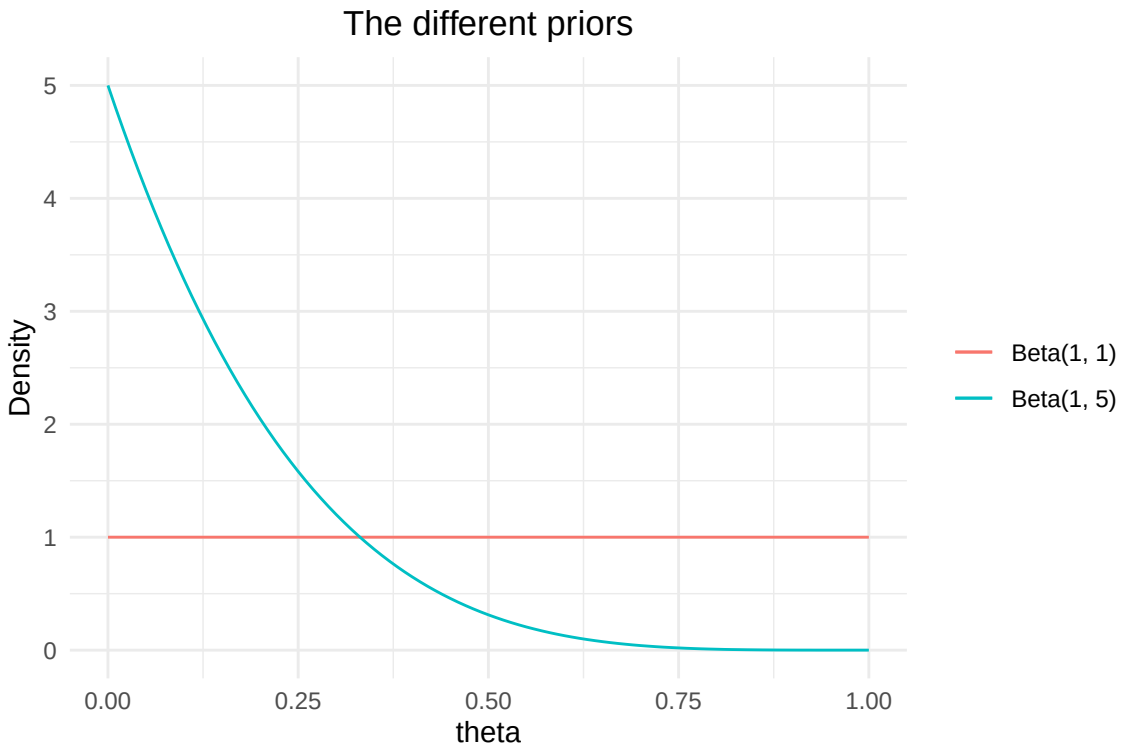

Figure 2: The two priors Beta$(1, 1)$ and Beta$(1, 5)$.