Tutorial 6 sample answer

Question 1

```
Command/code and output:
a)
        > dpois(3, lambda=3)
        [1] 0.2240418
     Command/code and output:
b)
        > ppois(3, lambda=3)
        [1] 0.6472319
     Command/code:
c)
        set.seed(100)
        Y <- rpois(100, lambda=4)
        mean(Y)
        var(Y)
     Output:
        > mean(Y)
        [1] 4.1
        > var(Y)
        [1] 3.060606
     Comment:
     Using different seed will give different result. Additionally, if we increase the
     number of sample, mean(Y) and var(Y) will get closer to 4.
     Command/code:
d)
        generate_Z <- function(n){</pre>
             X <- rpois(n, lambda=3)</pre>
             Y <- rpois(n, lambda=4)
             Z < - X + Y
             return(Z)
        }
        set.seed(100)
        Z <- generate Z(1000)</pre>
        mean(Z)
        var(Z)
        table(Z)/1000 # estimated probability
        dpois(7, lambda=7) # theoretical probability
```

```
Output:
```

```
> mean(Z)
[1] 7.148
> var(Z)
[1] 7.657754
> table(Z)/1000
                2
                      3
0.003 0.010 0.016 0.056 0.085 0.105 0.150 0.145 0.139
         10
               11
                     12
                            13
                                  14
                                        15
                                              16
                                                     17
0.111 0.069 0.049 0.031 0.013 0.005 0.008 0.001 0.002
   18
         19
0.001 0.001
> dpois(7, lambda=7)
[1] 0.1490028
```

Comment:

The mean and variance of Z is close to 7, as expected. And if we increase the number of samples, the values will get closer and closer to 7.

From the generated Z, we estimate P(Z=7) to be 0.145. And as shown, the true value for P(Z=7) is 0.149.

Question 2

our result.

```
Command/code:
a)
         f <- function(x) {
              return(x^x)
         }
b)
      Command/code:
         monte_carlo_int <- function(n) {</pre>
              X \leftarrow runif(n, min=1, max=2)
              est <-1/n*sum(f(X))
              return(est)
         }
c)
      Command/code:
         set.seed(100)
         monte carlo int(100000)
         integrate(f, 1, 2) # compare using built-in function
      Output:
         > set.seed(100)
         > monte_carlo int(100000)
         [1] 2.048201
         > integrate(f, 1, 2)
         2.050446 with absolute error < 2.3e-14
      Comment:
      In this question, we attempt to find the estimate of the integral using Monte Carlo
      method. The integral cannot be solved analytically, and we cannot compare our
      result with the true/analytical value.
      Using Monte Carlo integration, we estimate
                                 \int_{0}^{2} x^{x} dx \approx 2.048201
      using 100 000 samples. As usual, if we increase the number of samples, the
      estimation will be more accurate, but may become computationally costly. We can
      compare the result from our Monte Carlo method with the built-in function to
      integrate numerically in R which gives the value 2.050446, which is not far from
```

```
Command/function:
  generate x <- function(n){</pre>
       x <-_C()
       n_accept <- 0
       while(n accept < n){</pre>
            x1 <- runif(1, min=0, max=1)</pre>
            u <- runif(1, min=0, max=2.4)
            if(u \le dbeta(x1,3,5)){
                 x \leftarrow c(x, x1)
       }
       return(x)
  }
  set.seed(100)
  x <- generate_x(1000)</pre>
  length(x)
  hist(x, freq=FALSE)
  curve (dbeta(x, 3, 5), from=0, to=1, col="red",
  add=TRUE)
Example output:
  > length(x) #check the size of generated x
  [1] 1000
                             Histogram of x
            LO.
            0.0
            LO.
        Density
            0.
            0.5
            0.0
                         0.2
                 0.0
                                  0.4
                                          0.6
                                                  8.0
                                     Χ
```

Comment:

A common mistakes among student is by generating n samples of x_i , and keeping the values of x_i such that $u_i \le f(x)$. For example:

```
Tut6 <- function(n) {
    x <- runif(n, 0, 1)
    u <- runif(n, 0, 2.4)
    fx <- dbeta(x, 3, 5)
    samples <- x[u <= fx]
    return(samples)
}</pre>
```

This is not fully correct because the size of the generated sample will almost always be less than n. In this questions, to have n samples, you will have to use while loop.