LINKÖPINGS UNIVERSITET

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Exam

Advanced R programming (732G50)

Time: 8-12, 2015-11-25

Material: The extra material is included in the zip-file exam material.zip.

Grades: A = 19-20 points.

B = 17-18 points. C = 12-16 points. D = 10-11 points. E = 8-9 points. F = 0-7 points.

Instructions

Write your code in an R script file named **Main.R**. The R code should be complete and readable code, possible to run by copying directly into a script. Comment directly in the code whenever something needs to be explained or discussed. Follow the instructions carefully.

Problem 1 (5 p)

a) Create a function you call rdices() with three arguments, n, eyes and dices. The functions should simulate throwing dices. The eyes argument should specify the number of eyes of the dice (six should be the default value), dices should specify the number of dices that is beeing throwned (two should be the default) and n is the number of throws that has been done. The function should return a vector of length n with the sum of the eyes in the thrown dices.

```
rdices(5)
[1] 11 8 4 6 10

mean(rdices(100000, dices = 1))
[1] 3.49532
```

- b) What is the complexity of this algorithm ith regard to n. Assume that drawing a random draw is a constant operation.
- c) Visualize 1000 draws from your function (with the default values) as histogram using ggplot2.

Problem 2 (5 p)

a) Create a function called inverse_triangular_block_matrix() that takes matrices A, B and C and return their inverse as follows:

$$\begin{bmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{0} & \mathbf{C} \end{bmatrix}^{-1} = \begin{bmatrix} \mathbf{A}^{-1} & -\mathbf{A}^{-1}\mathbf{B}\mathbf{C}^{-1} \\ \mathbf{0} & \mathbf{C}^{-1} \end{bmatrix}$$

```
inverse_triangular_block_matrix(diag(2), 2*diag(2), 3*diag(2))
     [,1] [,2]
                    [,3]
                               [,4]
[1,]
             0 -0.666667
                          0.000000
[2,]
             1
                0.000000 -0.666667
        0
[3,]
        0
             0
                0.333333
                          0.000000
[4,]
             0 0.000000 0.333333
inverse_triangular_block_matrix(diag(1), -1*diag(1), 5*diag(1))
     [,1] [,2]
[1,]
        1 0.2
[2,]
     0 0.2
```

b) Implement a test suite withs unit tests that check that the result is of the correct class, of the right size/dimensions and that the function correctly return one of the examples above.

Problem 3 (5 p)

a) Create a function to simulate draws from a multivariate normal distribution. The function should be called rmvn() and take the arguments n (number of draws), mu (a vector of means of length m) and Sigma (a square matrix of size $m \times m$). Below is the description of how to do a multivariate draw taken from Wikipedia:

A widely used method for drawing (sampling) a random vector \mathbf{x} from the N-dimensional multivariate normal distribution with mean vector μ and covariance matrix Σ works as follows:

- 1. Find any real matrix \mathbf{A} such that $\mathbf{A}\mathbf{A}^T = \Sigma$. When Σ is positive-definite, the Cholesky decomposition is typically used, and the extended form of this decomposition can always be used (as the covariance matrix may be only positive semi-definite) in both cases a suitable matrix \mathbf{A} is obtained. [...]
- 2. Let $\mathbf{z} = (z_1, \dots, z_N)^T$ be a vector whose components are N independent standard normal variates.
- 3. Let \mathbf{x} be $\mu + \mathbf{Az}$. This has the desired distribution [...].

```
Sigma <- matrix(c(1,0.5,0.5,1), ncol=2)
mu <- c(2,5)
rmvn(3, mu, Sigma)
```

```
[,1] [,2]
[1,] 2.34632 6.20888
[2,] 1.45695 3.96545
[3,] 2.27750 3.99149

var(rmvn(100000, mu, Sigma))

[,1] [,2]
[1,] 1.001480 0.505597
[2,] 0.505597 1.010933
```

b) Document your function using roxygen2. The documentation should contain the title, description, the arguments and the resulting value of the function.

Problem 4 (5 p)

a) Implement the binary search algorithm with two arguments A that is a sorted vector and key that is an element we want to search for. Below you can find the pseudocode for binary search from Wikipedia. Note that in your implementation you need to set imin and imax yourself in your function.

```
int binary_search(int A[], int key, int imin, int imax) {
  // continue searching while [imin,imax] is not empty
  while (imin <= imax)</pre>
    // calculate the midpoint for roughly equal partition
    int imid = midpoint(imin, imax);
    if(A[imid] == key)
      // key found at index imid
      return imid;
    // determine which subarray to search
    else if (A[imid] < key)</pre>
      // change min index to search upper subarray
      imin = imid + 1;
    else // change max index to search lower subarray
      imax = imid - 1;
  // key was not found
  return KEY_NOT_FOUND;
```

```
A <- 10:20
binary_search(A, 19)

[1] 10

binary_search(A, 11)
```

```
[1] 2
binary_search(A, 5)
[1] NA
```

b) Implement a linear search based on the following pseudocode (taken from Wikipedia).

```
for each item in the list:
   if that item has the desired value,
     stop the search and return the item's location.
return KEY_NOT_FOUND;
}
```

```
A <- 10:20
linear_search(A, 19)

[1] 10

linear_search(A, 11)

[1] 2

linear_search(A, 5)</pre>
```

c) Compare the speed of the two search algorithms for the key 91281 and 0 in the vector (1, 2, 3, ..., 1 000 000). Good luck!