

Statistical Programming

Worksheet 2

By the end of the practical you should feel confident writing and calling functions, and using `if()`, `for()` and `while()` constructions.

You should complete and understand questions 1–6 for next time.

1. Review

- (a) Let $t = 2$: create a vector with $(i + 1)$ th entry $\frac{e^{-t}t^i}{i!}$ for $i = 0, \dots, 10$ (you might want to use the function `factorial()` for this).

```
t <- 2
n <- 10
sq <- 0:n
exp(-t) * t^sq/factorial(sq)
```

- (b) Write a function with arguments `t` and `n` that evaluates $\sum_{i=0}^n \frac{e^{-t}t^i}{i!}$.

```
parsum = function(t, n) {
  sq <- seq(from = 0, to = n)
  exp(-t) * sum(t^sq/factorial(sq))
}
```

Note this is the same as $P(N \leq n)$ where $N \sim \text{Poisson}(t)$, which is coded as `ppois()` in R:

```
ppois(15, lambda = 10)
parsum(t = 10, 15)
```

But notice that our version doesn't work well for large n : try `parsum(t=200, 200)`.

- (c) Write your function again using a `for()` loop. Do not use vectors, or the `sum()` function. Check it gives the same answers as (b).

```
parsum2 = function(t, n) {
  out <- 0
  for (i in 0:n) {
    out <- out + exp(-t) * t^i/factorial(i)
  }
  out
}
```

- 2. Solving a Quadratic.** Write a function with three arguments `a`, `b` and `c`, that returns the **real** roots of the equation $ax^2 + bx + c = 0$, if any. Your function should behave well if $a = 0$ and return an empty vector when there are no real roots.

```
roots <- function(a, b, c) {
  dis <- b^2 - 4 * a * c
  ## if discriminant is negative, no solutions
```

```

if (dis < 0)
  return(numeric(0))
if (a == 0) {
  ## if a=0 reduce to linear case
  if (b != 0)
    return(-c/b) else return(NaN) # if a=b=0 this is degenerate
}
## quadratic formula
(-b + c(1, -1) * sqrt(dis))/(2 * a)
}

```

3. Sieve of Eratosthenes. The Sieve of Eratosthenes is a method for finding all the prime numbers less than some specified n . Here is an outline of the algorithm:

- Create a vector x of integers from 2 to n , and an empty vector p .
- Given x , append the first element (say z) to p ; then remove any multiples of z (including z itself) from x .
- Stop when x is empty, and return p .

Write a function to implement this of Eratosthenes. It should take one argument n , and return all the primes up to n .

```

sieve <- function(n) {
  x <- 2:n
  p <- numeric(0)
  while (length(x) > 0) {
    p = c(p, x[1])
    x = x[x%%x[1] != 0]
  }
  p
}
sieve(1000)

```

Try it for $n = 10^3, 10^4, 10^5$. [Optional: can you see any way to speed this procedure up?]

Once $x[1]$ is greater than \sqrt{n} it's clear that we won't find more prime factors, so we could just stop and return $c(p, x)$ at this point.

4. Random Walks. Write a function `rndwlk`, with an argument k , that simulates a symmetric random walk (see lecture), stopping when the walk reaches k (or $-k$). After stopping it should return the entire walk.

```

rndwlk = function(k) {
  curr = 0 # current position
  out = 0 # vector of all positions
  while (abs(curr) < k) {
    curr = curr + sample(c(1, -1), 1) # new position
    out = c(out, curr) # add to vector
  }
  out
}

```

```

    }
    out
}

```

Try calling `plot(rndw1k(10))` a few times to see how it looks.

5. Simulating Discrete Distributions. In lectures you've seen that we can sample X from a discrete distribution on $\{1, \dots, k\}$ as follows: let $p_i = P(X = i)$. Then:

- generate $U \sim \text{Unif}[0, 1]$;
- set $X = \min\{i \mid \sum_{j=1}^i p_j \geq U\}$.

Write a function that, given `p` containing (p_1, \dots, p_k) can simulate X from this distribution. You may find the functions `cumsum()` and `which()` useful.

```

simDisc <- function(p) {
  U <- runif(1)
  cp <- cumsum(p) # get cumulative distribution
  which(cp > U)[1] # return which entry is first > U
}

```

Modify your function so that it takes an argument `n`, and produces a vector of `n` i.i.d. values from the distribution p . Comment on how you could check that your function worked as expected.

```

simDisc2 <- function(n, p) {
  U <- runif(n)
  out <- numeric(n) # numeric vector of length n
  cp <- cumsum(p)
  for (i in 1:n) {
    out[i] <- which(cp > U[i])[1]
  }
  out
}

```

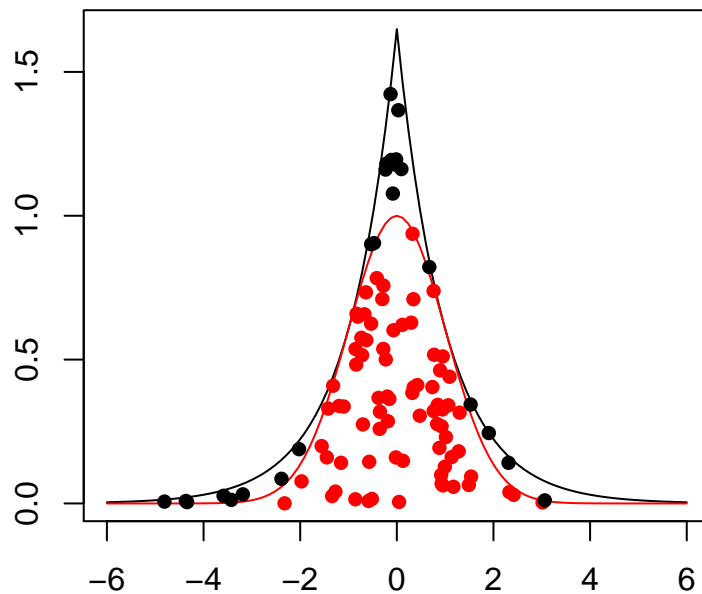
We can generate a large number of values and compare empirical (i.e. observed) frequencies with the probabilities. The CLT tells us how close the numbers should be.

6. Rejection Sampling. (You wont have seen this in simulation, so consider this an intro - the algorithm below is well defined and you can certainly implement it) We will write an R function to simulate $X \sim N(0,1)$ using rejection sampling with the double exponential proposal. That is, from a random variable Y with density

$$f_Y(y) = \exp(-|y|), \quad y \in \mathbb{R}.$$

```
#a plot for some intuition
#Throwing darts
x=seq(-6,6,length.out = 101)
plot(x,exp(-abs(x)+1/2),type='l',col=1,
     main="",
     xlab="Throwing darts: red points x-val ~ N(0,1)",
     ylab="",cex.lab=1)

lines(x,exp(-x^2/2),col=2)
for (k in 1:100) {
  y=sample(c(-1,1),1)*rexp(1);
  u=runif(1); #lines(c(y,y),c(0,exp(-abs(y)+1/2)),lwd=0.1);
  points(y,u*exp(-abs(y)+1/2),
        pch=16,cex=1,
        col=1+(u*exp(-abs(y)+1/2)<exp(-y^2/2)))
}
```



Throwing darts: red points $x\text{-val} \sim N(0,1)$

- (i) Write a function to simulate n i.i.d. values of Y . [Hint: you might want to start thinking about how to simulate an exponential random variable.] Here are a couple of possible solutions. You could also use a for-loop, though that is less efficient and harder to read.

```
rdbexp <- function(n) {
  mag <- log(runif(n)) # gets an exponential by inversion
  sign <- sample(c(-1, 1), n, replace = TRUE)
  out <- mag * sign
  return(out)
}
```

An alternative to `sample()` would be `Y <- 2*round(runif(n)) - 1`.

- (ii) Write a function implementing rejection for X . The algorithm is:
1. simulate $Y \sim \exp(-|x|)$ and $U \sim U(0, 1)$;
 2. if $U < \exp(-Y^2/2 + |Y| - 1/2)$ accept $X = Y$ and stop. Otherwise repeat 1.

[Hint: you can do this using a while statement. You should call the function you wrote in (a) to simulate Y . Your function should have no inputs, and return the simulated value of X .]

```
my_rnorm <- function(n) {
  out <- numeric(n)

  for (i in 1:n) {
    finished <- FALSE

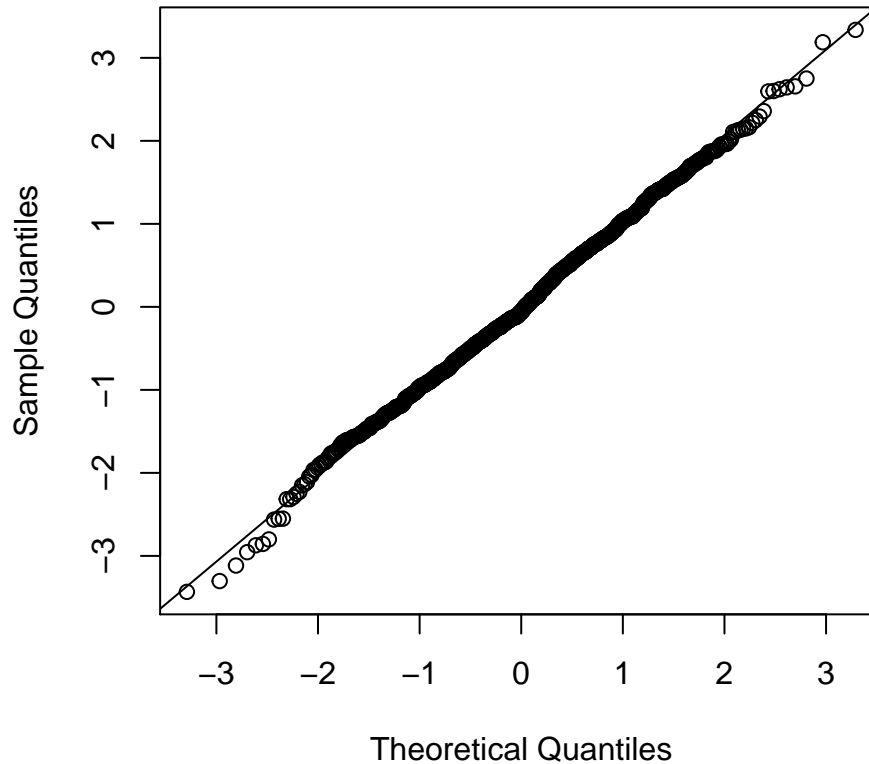
    while (!finished) {
      ## try a Y, see if
      Y <- rdbexp(1)
      U <- runif(1)
      finished <- (U < exp(-Y^2/2 + abs(Y) - 0.5))
    }
    out[i] <- Y
  }

  return(out)
}
```

- (iii) Test your rejection sampler by simulating 1000 samples and checking they are normal using the `qqnorm()` function.

```
X <- my_rnorm(1000)
qqnorm(X) # should be roughly straight
qqline(X) # to make the comparison easier
```

Normal Q-Q Plot



7. **Double for() Loop.** Using two `for()` loops, write a function with an argument `n`, which constructs the $n \times n$ matrix with entries $a_{ij} = i - j$.

```
addMat <- function(n) {  
  out <- matrix(0, n, n)  
  for (i in 1:n) {  
    for (j in 1:n) {  
      out[i, j] = i - j  
    }  
  }  
  out  
}  
addMat(4)
```

8. Moving Averages

- (a) Write a function to calculate the moving averages of length 3 of a vector $(x_1, \dots, x_n)^T$. That is, it should return a vector $(z_1, \dots, z_{n-2})^T$, where

$$z_i = \frac{1}{3}(x_i + x_{i+1} + x_{i+2}), \quad i = 1, \dots, n-2.$$

Call this function `ma3()`.

```

ma3 = function(x) {
  n = length(x)
  x1 = x[-(1:2)]
  x2 = x[-c(1, n)]
  x3 = x[-c(n - 1, n)]
  (x1 + x2 + x3)/3
}

x = rnorm(100)
plot(ma3(x), type = "l")

```

- (b) Write a function which takes two arguments, x and k , and calculates the moving average of x of length k . [Use a `for()` loop.]

```

ma = function(x, k) {
  n = length(x)
  out = x[-(1:(k - 1))]/k
  for (i in 2:k) {
    out = out + x[seq(from = k + 1 - i, to = n + 1 -
      i)]/k
  }

  out
}

max(abs(ma(x, 3) - ma3(x)))

```

- (c) How does your function behave if k is larger than (or equal to) the length of x ? You can tell it to return an error in this case by using the `stop()` function. Do so.
- (d) How does your function behave if $k = 1$? What should it do? Fix it if necessary. *It should just return x , but it may cause the `for()` loop to misbehave if you used $1:(k-1)$ in it.*

```

ma = function(x, k) {
  if (k == 1)
    return(x)
  n = length(x)
  out = x[-(1:(k - 1))]/k
  for (i in 2:k) {
    out = out + x[seq(from = k + 1 - i, to = n + 1 -
      i)]/k
  }

  out
}

max(abs(ma(x, 3) - ma3(x)))

```