732A90: Computational Statistics

Computer lab6 - Group11

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Question 1: Genetic algorithm

In this exercise we are going to perform one-dimensional maximization by using a genetic algorithm.

1.

Firstly, we define the function f() as

$$f(x) := \frac{x^2}{e^x} - 2 \exp(-(9\sin x)/(x^2 + x + 1)).$$

2.

Secondly, we define the function crossover(), that takes two scalars x and y as inputs, and returns a child as $\frac{x+y}{2}$.

3.

Thirdly, we define the function mutate(), that performs the integer division $x^2 \mod 30$, for a scalar input x.

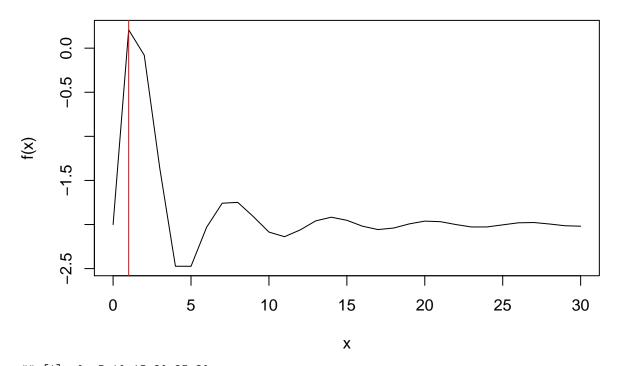
4.

Further, we will create a function called genetic(), with the parameters maxiter and mutprob. The settings of this genetic() function, as well as its output results, are presented in (a)-(e). The code can be found in the Appendix.

- (a). The function f() is plotted in the range from 0 to 30 in Figure X, and we can observe that there is a maximum value located around x = 1.
- (b). An initial population for the genetic algorithm is defined as X = (0, 5, 10, 15, ..., 30).
- (c). A vector called Values are computed, containing the function values for each population point.
- (d). The genetic() function performs maxiter iterations. For each iteration...
- (e).

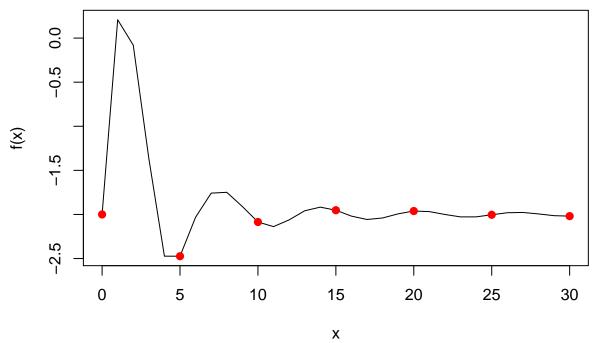
5.

By using the defined functions from previous tasks (1.1-1.4), we are going to observe the initial population and final population. This is done by running the code with different combinations of maxiter= 10,100 and mutprob= 0.1,0.5,0.9.



[1] 0 5 10 15 20 25 30

[1] -2.000000 -2.473573 -2.085654 -1.951947 -1.961344 -2.003663 -2.019194



Question 2: EM algorithm

The purpose with this exercise is to implement the EM algorithm. For this, we are given the data file physicall.csv, containing a behavior of two related physical processes Y = Y(X) and Z = Z(X).

1.

The first step is to examine the data set physical1.csv, to see if the two processes are related to each other.

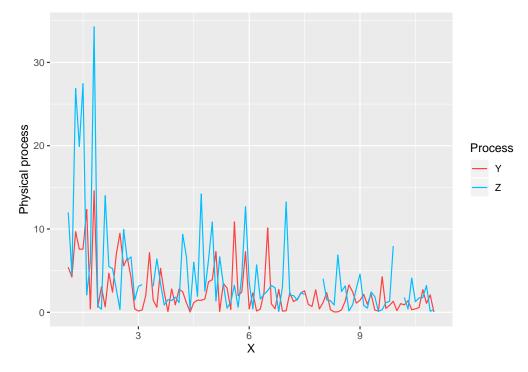


Figure 1: Time series plot of the dependence of Z and Y versus X.

In Figure 1 it seems that the two processes are related to each other, with respect to X, since the graphs follows similar patterns. We can also observe that the physical process Z has a greater variation, especially at the beginning of the series, but also in general, compared to the process Y.

2.

Using the following model,

$$Y_i \sim exp\left(\frac{X_i}{\lambda}\right)$$

$$Z_i \sim exp\left(\frac{X_i}{2\lambda}\right)$$

where λ is an unknown parameter, we derive the EM algorithm to estimate λ .

Y and Z are defined by the following density functions,

$$Z(X) = \frac{X_i}{2\lambda} exp\left(-\frac{X_i}{2\lambda}Z_i\right)$$
$$Y(X) = \frac{X_i}{\lambda} exp\left(-\frac{X_i}{\lambda}Y_i\right)$$

Assuming that Y(X) and Z(X) are i.i.d. we can obtain the joint density function,

$$f(Y,Z) = \frac{X_i}{2\lambda} exp\left(-\frac{X_i}{2\lambda}Z_i\right) * \frac{X_i}{\lambda} exp\left(-\frac{X_i}{\lambda}Y_i\right)$$
$$= \frac{X_i^2}{2\lambda^2} exp\left[-\frac{X_i(Z_i - 2Y_i)}{2\lambda}\right]$$

From this point, we compute the likelihood of the distribution,

$$\mathcal{L}(Y, Z|X, \lambda) = \prod_{i=1}^{n} \frac{X_i^2}{2\lambda^2} exp\left[-\frac{X_i(Z_i - 2Y_i)}{2\lambda}\right]$$

The code can be found in the Appendix.

3.

4.

As a final task, we are going to see if the optimal value $\lambda=10.69566$ is reasonable for the physical processes Y=Y(X) and Z=Z(X). We examine the results by computing the expected values of Y and Z, which is given by $E[Y_i]=\frac{\lambda}{X_i}$ and $E[Z_i]=\frac{2\lambda}{X_i}$, respectively.

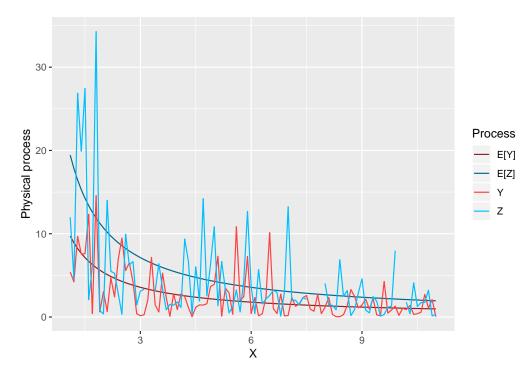


Figure 2: Time series plot of E[Y] and E[Z] versus X, and the dependence of Z and Y versus X.

In Figure 2, we can see that E[Y] and E[Z] versus X are in the same plot as Y and Z versus X. From this plot it is clear that the optimal λ is reasonable, since the exponential curves E[Y] and E[Z], the follow their corresponding values of Y and Z.

Appendix

```
knitr::opts_chunk$set(echo = FALSE)
# R version
RNGversion('3.5.1')
library("ggplot2")
#1.1
f <- function(x){</pre>
  return(x^2/\exp(x) - 2*\exp(-1*(9*\sin(x)) / (x^2 + x + 1)))
}
crossover <- function(x,y){</pre>
  return((x+y) / 2)
}
mutate <- function(x){</pre>
  return(x^2 %% 30)
}
#4
genetic <- function(maxiter, mutprob){</pre>
  plot(x = seq(0,30), y = f(seq(0,30)), type = "l", xlab = "x", ylab = "f(x)")
  abline(v=seq(0,30)[which.max(f(seq(0,30)))], col="red")
  #b
  X = seq(0,30,5)
  Values = f(X)
  \#d
  #set seed
  set.seed(1234567890)
  for (i in 1:maxiter) {
    parents = match(sample(X, 2),X)
    \#ii
    victim = order(Values)[1]
    \#iii
    kid = round(crossover(parents[1],parents[2]))
    p = runif(1)
```

```
if (p < mutprob) {</pre>
      kid = mutate(kid)
    \#iv
    X[victim] = kid
    Values = f(X)
    #υ
    max = max(Values)
  }
  #e
  print(X)
  print(Values)
  plot(x = seq(0,30), y = f(seq(0,30)), type = "l", xlab = "x", ylab = "f(x)")
  points(x = X, y = Values, col = "red", pch = 19)
# Just testing no change, i.e. initial population
genetic(1,0)
# 2.1
physical <- read.csv2("physical1.csv", sep = ",")</pre>
X <- as.numeric(as.character(physical$X))</pre>
Y <- as.numeric(as.character(physical$Y))
Z <- as.numeric(as.character(physical$Z))</pre>
data \leftarrow data.frame(X = c(X,X), value = c(Y,Z), Process= rep(c("Y","Z"), each= 100))
# var(Z, na.rm = TRUE)
# var(Y, na.rm = TRUE)
# Time series plot
ggplot(data = data, aes(x = X, y = value, col = Process)) +
  geom_line() +
  ylab("Physical process")+
  scale_color_manual(values=c( "brown1", "deepskyblue1"))
## 2.2
EM_algorithm <- function(X, Y, Z, lambda, eps, k_max) {</pre>
    Z_{obs} \leftarrow Z[!is.na(Z)]
    Z_miss <- Z[is.na(Z)]</pre>
    X_obs <- X[!is.na(Z)]</pre>
    X_miss <- X[is.na(Z)]</pre>
    Y_obs <- Y[!is.na(Z)]
    Y_miss <- Y[is.na(Z)]</pre>
```

```
n <- length(c(Z_obs, Z_miss))</pre>
   r <- length(Z_obs)
   k<-1
   lambda_prev <- lambda+10+100*eps #random number to initialize the algorithm
   lambda_curr <- lambda</pre>
   while (k<k_max+1 && abs(lambda_prev-lambda_curr)>=eps) {
       lambda_prev<-lambda_curr
       ## E-step
       lambda_curr <- (sum(X_obs*(Z_obs+2*Y_obs)) + 2*sum(lambda_curr+X_miss*Y_miss))/(4*n)</pre>
       k<-k+1
   }
   return(c(number_of_iterations = k-1, optimal_lambda = lambda_curr))
}
##2.3
EM_algorithm(X,Y,Z, 100, 0.001, 100)
optimal_lambda <- EM_algorithm(X,Y,Z, 100, 0.001, 100)[2]
# Expected values
EY <- optimal_lambda/X</pre>
EZ <- 2*optimal_lambda/X
 data2 \leftarrow data.frame(X = c(X,X), value = c(Y,Z,EY,EZ), Process = rep(c("Y","Z","E[Y]","E[Z]"), each = 100) 
ggplot(data = data2, aes(x = X, y = value, col = Process)) +
 geom_line() +
 ylab("Physical process") +
 scale_color_manual(values=c("brown4", "deepskyblue4", "brown1", "deepskyblue1"))
```