

# Homework 4

*Xiaokang Liu*

*28 September 2018*

## Contents

<b>1</b>	<b>Many local maxima</b>	<b>1</b>
1.1	Part 1 . . . . .	1
1.2	Part 2 . . . . .	2
1.3	Part 3 . . . . .	2
1.4	Part 4 . . . . .	3
1.5	Part 5 . . . . .	4
<b>2</b>	<b>Modeling beetle data</b>	<b>5</b>
2.1	Fit the population growth model . . . . .	5
2.2	Show the contour plot . . . . .	6
2.3	Log-Normality . . . . .	7

## 1 Many local maxima

Consider the probability density function with parameter  $\theta$ :

$$f(x; \theta) = \frac{1 - \cos(x - \theta)}{2\pi}, \quad 0 \leq x \leq 2\pi, \quad \theta \in (-\pi, \pi).$$

A random sample from the distribution is

```
samp <- c(3.91, 4.85, 2.28, 4.06, 3.70, 4.04, 5.46, 3.53, 2.28, 1.96,  
          2.53, 3.88, 2.22, 3.47, 4.82, 2.46, 2.99, 2.54, 0.52)
```

### 1.1 Part 1

The log-likelihood function of  $\theta$  based on the sample is

$$l(\theta) = \sum_{i=1}^n \ln(1 - \cos(x_i - \theta)) - n \ln(2\pi).$$

And the plot of it between  $-\pi$  and  $\pi$  is:

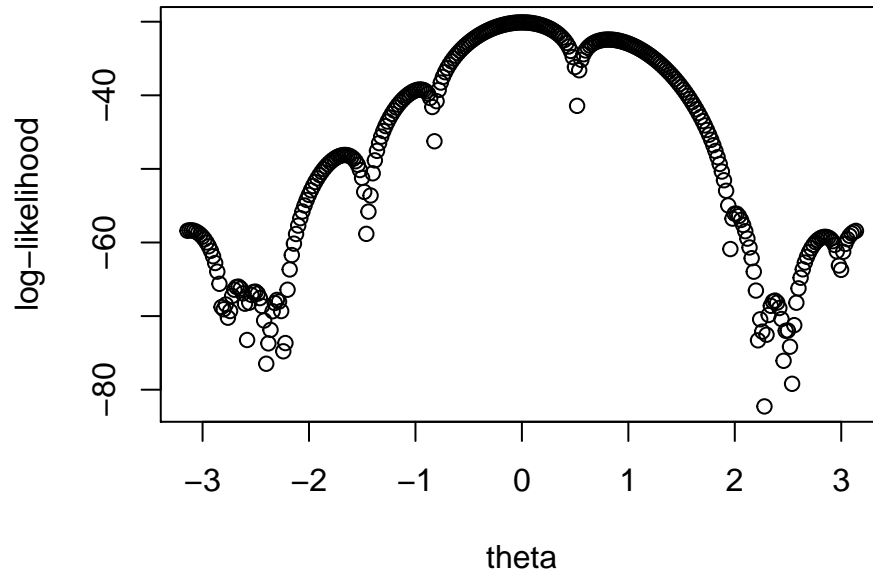
```
logf <- function(dat,para){  
  f <- sum(log(1-cos(dat-para))-log(2*pi))  
  return(f)  
}  
  
int <- seq(-pi,pi,0.02)
```

```

val <- vector()
for (i in 1:length(int)){
  val <- c(val,logf(samp,int[i]))
}

plot(int,val,xlab = "theta",ylab = "log-likelihood")

```



## 1.2 Part 2

The expectation of  $X$  can be derived by

$$\begin{aligned}
 E(x|\theta) &= \int_0^{2\pi} x \frac{1 - \cos(x - \theta)}{2\pi} dx \\
 &= \pi - \frac{1}{2\pi} \int_{-\theta}^{2\pi - \theta} x \cos(x) dx \\
 &= \pi + \sin(\theta).
 \end{aligned}$$

Thus, the moment estimator of  $\theta$  is  $\tilde{\theta}_n = \arcsin(\bar{X}_n - \pi)$ . Since we have defined an interval for  $\theta$ , thus the only possible moment estimator from the sample is 0.095394.

## 1.3 Part 3

```

init <- asin(mean(samp)-pi) # it's okay to +/- 2*k*pi
l1 <- function(para,dat){
  f <- -sum(sin(samp-para)/(1-cos(samp-para)))
  return(f)
}
l2 <- function(para,dat){
  f <- sum((cos(samp-para)-1)/(1-cos(samp-para))^2)
}

```

```

}

NR <- function(ini,x,tol,max_ite){
  err <- 100
  iter <- 0
  conver <- 0
  while ((err > tol) & (iter < max_ite)) {
    ini1 <- ini-l1(ini,x)/l2(ini,x)
    err <- abs(ini1-ini)
    ini <- ini1
    iter <- iter+1
  }
  if (iter >= max_ite) conver <- 1
  return(list(ini1=ini1,iter=iter,err=err,conver=conver))
}

NR(init,samp,tol = .Machine$double.eps^0.5,max_ite = 200)

```

```

## $ini1
## [1] 0.003118157
##
## $iter
## [1] 4
##
## $err
## [1] 1.525437e-10
##
## $conver
## [1] 0

```

## 1.4 Part 4

```

NR(-2.7,samp,tol = .Machine$double.eps^0.5,max_ite = 200)

## $ini1
## [1] -2.668857
##
## $iter
## [1] 4
##
## $err
## [1] 8.772369e-09
##
## $conver
## [1] 0

```

```
NR(2.7,samp,tol = .Machine$double.eps^0.5,max_ite = 200)
```

```
## $ini1
## [1] 2.848415
##
## $iter
## [1] 5
##
## $err
## [1] 1.682339e-10
##
## $conver
## [1] 0
```

## 1.5 Part 5

```
init <- seq(-pi, pi, length.out = 200)
n <- length(init)
eps0 <- .Machine$double.eps^0.5
max_ite0 <- 200
```

```
val_NR <- rep(0,n)
conv_NR <- rep(0,n)
iter_NR <- rep(0,n)
```

```
for (i in 1:n){
  res <- NR(init[i],samp,eps0,max_ite0)
  val_NR[i] <- res$ini1
  iter_NR[i] <- res$iter
  conv_NR[i] <- res$conver
}
```

```
val_NR1 <- round(val_NR,9)
uni <- unique(val_NR1)
gr <- matrix(nrow = 18, ncol = 200)
for (j in 1:18){
  gr[j,] <- init*(val_NR1==uni[j])
}
# group 1 to group 18
table(val_NR1)
```

```
## val_NR1
## -3.112470507 -2.786556852 -2.668857459 -2.509356033 -2.388266628
##           11           2           5           6           1
## -2.297925969 -2.232191899 -1.662712395 -1.447502553 -0.954405837
```

```
##           4           1           24           1           19
## 0.003118157 0.812637417 2.007223238 2.237012923 2.374711666
##           42           46           8           2           6
## 2.488449651 2.848415325 3.1707148
##           2           15           5
```

```
for (i in 1:18){
  x <- gr[i,]
  z <- x[ min( which ( x != 0 )) : max( which( x != 0 )) ]
  cat("At group", i, ",values are:",z,"\n")
}
```

```
## At group 1 ,values are: -3.141593 -3.110019 -3.078445 -3.046871 -3.015297 -2.983724 -2.95211
## At group 2 ,values are: -2.794281 -2.762707
## At group 3 ,values are: -2.731133 -2.69956 -2.667986 -2.636412 -2.604838
## At group 4 ,values are: -2.573264 -2.541691 -2.510117 -2.478543 -2.446969 -2.415395
## At group 5 ,values are: -2.383822
## At group 6 ,values are: -2.352248 -2.320674 -2.2891 -2.257526
## At group 7 ,values are: -2.225953
## At group 8 ,values are: -2.194379 -2.162805 -2.131231 -2.099657 -2.068084 -2.03651 -2.00493
## At group 9 ,values are: -1.436608
## At group 10 ,values are: -1.405034 -1.37346 -1.341886 -1.310313 -1.278739 -1.247165 -1.2155
## At group 11 ,values are: -0.8051318 -0.773558 -0.7419842 -0.7104104 -0.6788366 -0.6472628 -0
## At group 12 ,values are: 0.5209676 0.5525414 0.5841152 0.615689 0.6472628 0.6788366 0.71041
## At group 13 ,values are: 1.973362 2.004936 2.03651 2.068084 2.099657 2.131231 2.162805 2.19
## At group 14 ,values are: 2.225953 2.257526
## At group 15 ,values are: 2.2891 2.320674 2.352248 2.383822 2.415395 2.446969
## At group 16 ,values are: 2.478543 2.510117
## At group 17 ,values are: 2.541691 2.573264 2.604838 2.636412 2.667986 2.69956 2.731133 2.76
## At group 18 ,values are: 3.015297 3.046871 3.078445 3.110019 3.141593
```

## 2 Modeling beetle data

The counts of a floor beetle at various time points (in days) are given in a dataset as below:

```
beetles1 <- data.frame(
  days = c(0, 8, 28, 41, 63, 69, 97, 117, 135, 154),
  beetles = c(2, 47, 192, 256, 768, 896, 1120, 896, 1184, 1024))
```

### 2.1 Fit the population growth model

```
lk <- function(t,a,b){
  f <- (4-4*exp(-b*t))/(2+(a-2)*exp(-b*t))^2
  return(f)
}
```

```

lr <- function(t,a,b){
  f <- (2*a*(a-2)*t*exp(-b*t))/(2+(a-2)*exp(-b*t))^2
  return(f)
}

GN <- function(t,y,ini,tol,max_ite){
  err <- 100000000
  iter <- 0
  conver <- 0
  while ((err > tol)&(iter < max_ite)) {
    f1 <- lk(t,ini[1],ini[2])
    f2 <- lr(t,ini[1],ini[2])
    A <- cbind(f1,f2)
    ini1 <- ini+solve(t(A)%*%A+0.00001*diag(nrow = 2))%*%t(A)%*%(y-2*ini[1]/(2+(ini[1]-2)*exp(-ini[2]*t)))^2)
    err <- sum((ini-ini1)^2)
    ini <- ini1
    iter <- iter+1
  }
  ss <- sum((y-2*ini[1]/(2+(ini[1]-2)*exp(-ini[2]*t)))^2)
  if (iter >= max_ite) conver <- 1
  return(list(est=ini,ss=ss,iter=iter,err=err,conver=conver))
}
GN(beetles1$days,beetles1$beetles,c(1100,0.1),.Machine$double.eps^0.5,200)

```

```

## $est
##           [,1]
## f1 1049.4072555
## f2    0.1182684
##
## $ss
## [1] 73419.7
##
## $iter
## [1] 9
##
## $err
## [1] 3.052186e-09
##
## $conver
## [1] 0

```

## 2.2 Show the contour plot

Show the contour plot of the sum of squared errors.

```

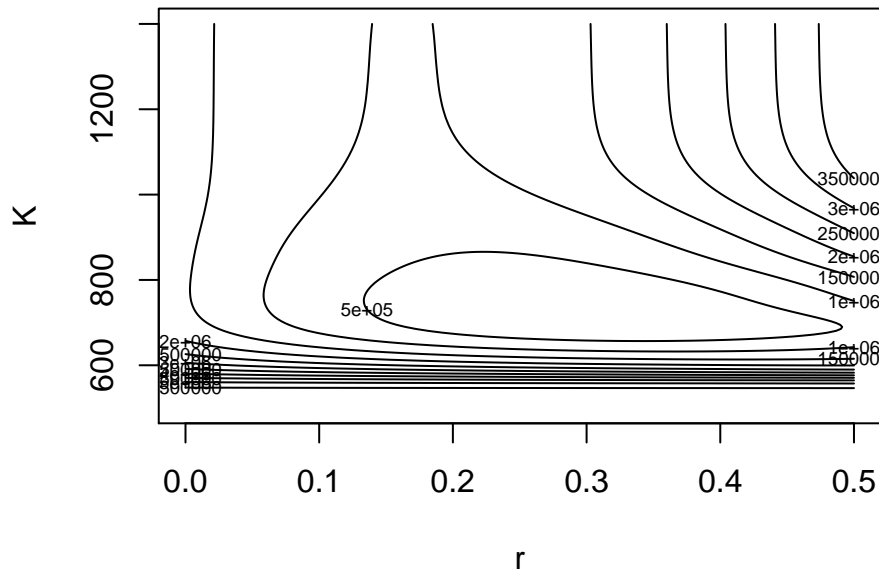
kseq <- seq(500, 1400, length.out = 200)
rseq <- seq(0, 0.5, length.out = 200)

```

```

cont <- matrix(nrow = length(kseq), ncol = length(rseq))
y <- beetles1$beetles
t <- beetles1$days
for (i in 1:length(kseq)){
  for (j in 1:length(rseq)){
    cont[i,j] <- sum((y-2*kseq[i]/(2+(kseq[i]-2)*exp(-rseq[j]*t)))^2)
  }
}
# contour plot
contour(rseq,kseq,cont,xlab="r",ylab="K",method = "simple")

```



## 2.3 Log-Normality

If we assume  $\log N_t$  are independent and normally distributed with mean  $\log f(t)$  and variance  $\sigma^2$ . The the log-likelihood function is

$$l(x, r, K, \sigma^2) = -\frac{n}{2} \ln \sigma^2 - \frac{1}{2\sigma^2} \sum_{i=1}^n (\log(x_i) - \log \frac{2k}{2 + (k-2)e^{-rt_i}})^2 + C.$$

where  $C$  is a constant. Here we use BFGS with linear constraints to solve this problem. Since the commonly used optimization functions in `r` will do minimization in default, here we use the negative log-likelihood function as the objective function. We provide the function for the objective function and its first order derivative function.

```

y <- beetles1$beetles
t <- beetles1$days
n <- length(y)
fr <- function(x){
  x1 <- x[1] # sigma^2
  x2 <- x[2] # k
  x3 <- x[3] # r

```

```

log(x1)*n/2+sum((log(y)-log(2*x2/(2+(x2-2)*exp(-x3*t))))^2)/2/x1
}
grr <- function(x){
  x1 <- x[1] # sigma^2
  x2 <- x[2] # k
  x3 <- x[3] # r
  c(n/2/x1-sum((log(y)-log(2*x2/(2+(x2-2)*exp(-x3*t))))^2)/2/x1^2,
    sum((log(y)-log(2*x2/(2+(x2-2)*exp(-x3*t))))*(2*exp(-x3*t)-2)/(2+(x2-
      2)*exp(-x3*t)))/x2/x1,
    sum((log(y)-log(2*x2/(2+(x2-2)*exp(-x3*t))))*((2-x2)*t*exp(-x3*t))/(2+
      (x2-2)*exp(-x3*t)))/x1
  )
}

# add linear constraints on parameters, they are all positive
mod1 <- constrOptim(c(0.01,2200,0.1), fr, grr, ui=diag(nrow = 3),
  ci=c(0,0,0), hessian = TRUE)
mod1

## $par
## [1] 0.8559366 2151.1598581 0.1431047
##
## $value
## [1] 4.221507
##
## $counts
## function gradient
## 152 101
##
## $convergence
## [1] 0
##
## $message
## NULL
##
## $hessian
## [,1] [,2] [,3]
## [1,] 6.822987683 -2.915587e-03 -0.0111829
## [2,] -0.002915587 2.755498e-07 0.0175775
## [3,] -0.011182899 1.757750e-02 1186.2494911
##
## $outer.iterations
## [1] 3
##
## $barrier.value
## [1] -1.435492

```



```
var.est <- diag(solve(-mod1$hessian))  
var.est
```

```
## [1] 1.797321e-03 8.125787e+05 -6.646766e-04
```

From the result, we know that the estimated variances from Hessian matrix are  $1.797 \times 10^{-3}$ ,  $8.126 \times 10^5$  and 0. Since the last one is negative, thus we set it to be 0.