

STA410 Assignment #3

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Question 1

(a)

Since we have,

$$\text{tr}(\hat{A}) = \frac{1}{m} \sum_{i=1}^m V_i^T A V_i$$

Then:

$$\text{Var}(\text{tr}(\hat{A})) = \frac{1}{m^2} \text{Var}(V^T A V) = \frac{1}{m^2} \{E[(V^T A V)^2] - \text{tr}(A)^2\}$$

, where $\frac{1}{m^2}$ and $\text{tr}(A)^2$ are constant.

So it is sufficient to minimize $E[(V^T A V)^2]$.

Since we are assuming that V_1, \dots, V_n are independent with mean 0 and variance 1, it follows that $E(V_i V_j V_k V_l) = 0$ or 1 unless $i = j = k = l$,

so we have:

$$\begin{aligned} E[(V^T A V)^2] &= \sum_{i=1}^n \sum_{j=1}^n \sum_{k=1}^n \sum_{l=1}^n a_{ij} a_{kl} E(V_i V_j V_k V_l) \\ &= a_{11}^2 E(V_1^4) + a_{22}^2 E(V_2^4) + \dots + a_{nn}^2 E(V_n^4) + \text{constant} \\ &= \sum_{i=1}^n a_{ii}^2 E(V_i^4) + \text{constant} \end{aligned}$$

So it is sufficient to minimize $E(V_i^4)$.

Since

$$\text{Var}(V_i) = E(V_i^2) - [E(V_i)]^2$$

so

$$E(V_i^2) = \text{Var}(V_i) + [E(V_i)]^2 = 1 + 0 = 1$$

Then

$$E(V_i^4) = \text{Var}(V_i^2) + [E(V_i^2)]^2 = \text{Var}(V_i^2) + 1$$

So it is sufficient to minimize $\text{Var}(V_i^2)$.

$\text{Var}(V_i^2)$ is minimized if V_i^2 is constant (with probability 1).

Since $E(V_i)$ and $E(V_i^2)$ must equal to 0 and 1 respectively, so that $\text{Var}(\text{tr}(\hat{A}))$ is minimized by taking the elements of V_i to be ± 1 , and each with probability $\frac{1}{2}$.

(b)

Since

$$H = X(X^T X)^{-1} X^T = \begin{pmatrix} H_{11} & H_{12} \\ H_{21} & H_{22} \end{pmatrix}$$

So

$$H \begin{pmatrix} V \\ 0 \end{pmatrix} = \begin{pmatrix} H_{11} & H_{12} \\ H_{21} & H_{22} \end{pmatrix} \begin{pmatrix} V \\ 0 \end{pmatrix} = \begin{pmatrix} H_{11}V \\ H_{21}V \end{pmatrix}$$

$$H \begin{pmatrix} H_{11}^{k-1}V \\ 0 \end{pmatrix} = \begin{pmatrix} H_{11} & H_{12} \\ H_{21} & H_{22} \end{pmatrix} \begin{pmatrix} H_{11}^{k-1}V \\ 0 \end{pmatrix} = \begin{pmatrix} H_{11}^k V \\ H_{21} H_{11}^{k-1} V \end{pmatrix}$$

(c)

Firstly, modify the given leverage function as follows:

```
#modify the given leverage function
leverage <- function(x1,x2,w,r=10,m=100) {
  qrx1 <- qr(x1)
  qrx2 <- qr(x2)
  n <- nrow(x1) #since X1 and X2 have the same #row=1000
  lev1 <- NULL
  lev2 <- NULL
  for (i in 1:m) {
    v <- ifelse(runif(n)>0.5,1,-1)
    v[-w] <- 0 #set other points = 0
    v01 <- qr.fitted(qrx1,v)
    v02 <- qr.fitted(qrx2,v)
    f1 <- v01
    f2 <- v02
    for (j in 2:r) {
      v01[-w] <- 0
      v02[-w] <- 0
      v01 <- qr.fitted(qrx1,v01)
      v02 <- qr.fitted(qrx2,v02)
      f1 <- f1 + v01/j
      f2 <- f2 + v02/j
    }
    lev1 <- c(lev1,sum(v*f1)) #estimated leverage of X1
    lev2 <- c(lev2,sum(v*f2)) #estimated leverage of X2
  }
  std.err1 <- exp(-mean(lev1))*sd(lev1)/sqrt(m) #standard error of leverage of X1
  std.err2 <- exp(-mean(lev2))*sd(lev2)/sqrt(m) #standard error of leverage of X2
  std.err.diff <- 0.5*(exp(-mean(lev1))+exp(-mean(lev2)))*sd(lev1-lev2)/sqrt(m)
  lev1 <- 1 - exp(-mean(lev1))
  lev2 <- 1 - exp(-mean(lev2))
  r <- list(lev = c(lev1,lev2), lev.diff = lev1-lev2, std.err=c(std.err1,std.err2,std.err.diff))
  r #($std.err[3]: the third element of $std.err is an estimate of the standard error of the difference.
}
```

We define the two design matrices X1 and X2 as follows:

```
#construct two design matrices X1 and X2
x <- c(1:1000)/1000
X1 <- 1
for (k in 1:5) X1 <- cbind(X1,cos(2*k*pi*x),sin(2*k*pi*x))
library(splines) # loads the library of functions to compute B-splines
X2 <- cbind(1,bs(x,df=10))
```

Then we can use the above modified leverage function to compute the 20 leverages for X1 and X2 and their differences.

In this example, we chose $r = 20$ and $m = 100$.

```
lev1 <- NULL #empty list of 20 estimated leverages for X1
lev2 <- NULL #empty list of 20 estimated leverages for X2
lev.diff <- NULL #empty list of 20 differences of two estimated leverages
std.err.diff <- NULL #empty list of estimates of the standard errors of the differences.
for (k in (1:20)) {
  w_k <- c((50*k-49):(50*k))  #(k-1)/20 < xi <= k/20 eg. when k=1, 0<xi<=1/20, w_k = c(1:50)
  r <- leverage(X1,X2,w_k,r=20,m=100)
  lev1 <- c(lev1, r$lev[1])
  lev2 <- c(lev2, r$lev[2])
  lev.diff <- c(lev.diff, r$lev.diff)
  std.err.diff <- c(std.err.diff, r$std.err[3])
}
```

Here is a list of 20 estimated leverages for X1:

```
lev1 #a list of 20 estimated leverages for X1

## [1] 0.5305118 0.5599752 0.4236836 0.5238033 0.5026695 0.5138854 0.4797832
## [8] 0.4662939 0.5213244 0.5392356 0.4990046 0.6080412 0.6029045 0.4995600
## [15] 0.5896916 0.5199767 0.5997865 0.4923020 0.5622733 0.4899432
```

Here is a list of 20 estimated leverages for X2:

```
lev2 #a list of 20 estimated leverages for X2

## [1] 0.9785562 0.6697689 0.4146736 0.4563229 0.4009268 0.4432598 0.3156640
## [8] 0.4048661 0.3479995 0.4293976 0.3880660 0.4149868 0.5384891 0.3320690
## [15] 0.5144641 0.4090392 0.5238404 0.4823393 0.6710516 0.9619090
```

Here is a list of 20 differences of two estimated leverages:

```
lev.diff #a list of 20 differences of two estimated leverages

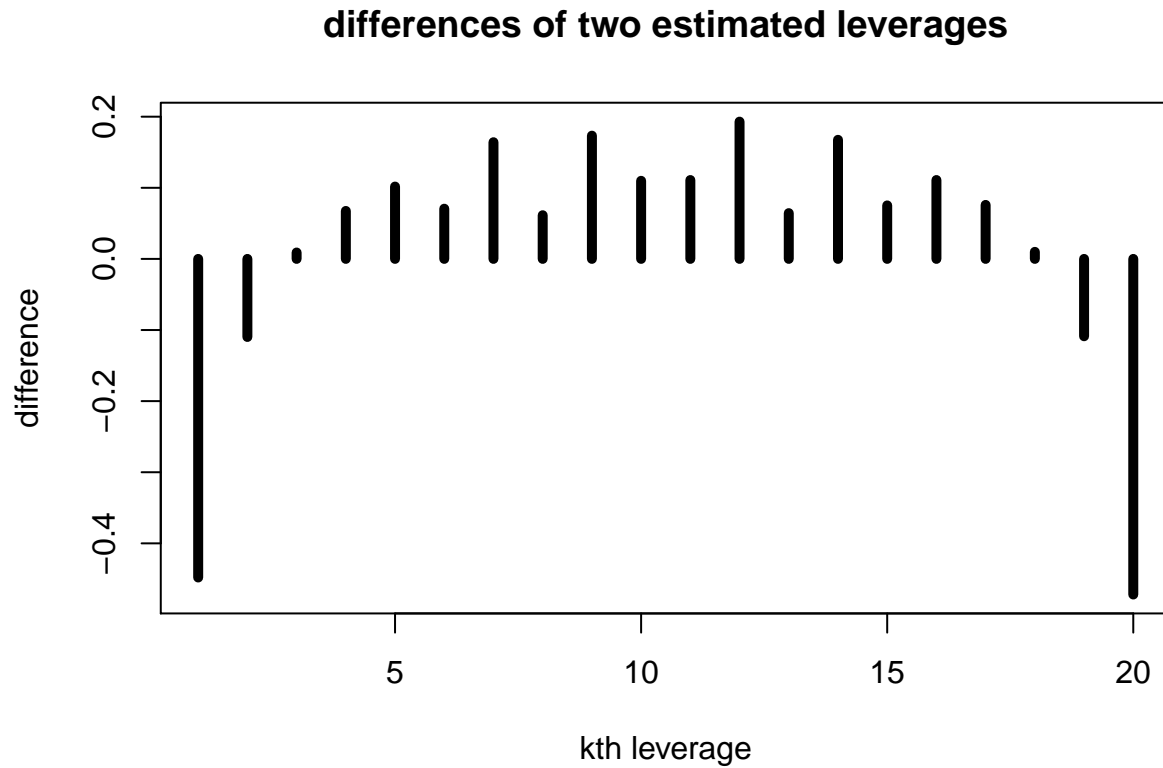
## [1] -0.448044371 -0.109793724 0.009009995 0.067480367 0.101742641
## [6] 0.070625601 0.164119261 0.061427770 0.173324886 0.109838041
## [11] 0.110938634 0.193054436 0.064415398 0.167490948 0.075227521
## [16] 0.110937470 0.075946109 0.009962668 -0.108778308 -0.471965779
```

Here is a list of estimates of the standard errors of the differences.

```
std.err.diff #a list of estimates of the standard errors of the differences.

## [1] 0.097816078 0.018897767 0.003763207 0.010334948 0.013176147
## [6] 0.007289563 0.020473768 0.006070282 0.025450391 0.013948671
## [11] 0.014781839 0.022027997 0.007904531 0.023221664 0.007347576
## [16] 0.012058281 0.012519934 0.003739175 0.015854084 0.096327879
```

```
plot(c(1:20), lev.diff, type="h", lwd=5,
     main="differences of two estimated leverages",
     xlab="kth leverage", ylab="difference")
```



From the above plot, we find that the first($k=1$) and last($k=20$) differences are the two largest, and other differences are small, with absolute values less than 0.2. Additionally, the differences of the first 10 leverages($k = 1, \dots, 10$) are almost symmetric with differences of the other 10 leverages($k = 11, \dots, 20$).

Question 2

(a)

Let \bar{x} be the sample mean and s^2 be the sample variance of the data x_1, \dots, x_n . Since for Gamma Distribution, we have

$$E(x) = \frac{\alpha}{\lambda} \quad \text{and} \quad \text{Var}(x) = \frac{\alpha}{\lambda^2}$$

Then

$$\alpha = \frac{[E(x)]^2}{\text{Var}(x)} \Rightarrow \hat{\alpha} = \frac{\bar{x}^2}{s^2}$$

$$\lambda = \frac{E(x)}{\text{Var}(x)} \Rightarrow \hat{\lambda} = \frac{\bar{x}}{s^2}$$

(b)

Since we have:

$$f(x; \alpha, \lambda) = \frac{\lambda^\alpha x^{\alpha-1} \exp(-\lambda x)}{\Gamma(\alpha)} \quad \text{for } x > 0$$

Then, likelihood function for α and λ is:

$$L(\alpha, \lambda) = \prod_{i=1}^n \frac{\lambda^\alpha x_i^{\alpha-1} \exp(-\lambda x_i)}{\Gamma(\alpha)}$$

log-likelihood function for α and λ is:

$$\begin{aligned} l(\alpha, \lambda) &= \sum_{i=1}^n \ln \left\{ \frac{\lambda^\alpha x_i^{\alpha-1} \exp(-\lambda x_i)}{\Gamma(\alpha)} \right\} \\ &= \sum_{i=1}^n \alpha \ln(\lambda) + (\alpha - 1) \ln(x_i) - \lambda x_i - \ln(\Gamma(\alpha)) \\ &= n\alpha \ln(\lambda) + (\alpha - 1) \sum_{i=1}^n \ln(x_i) - \lambda \sum_{i=1}^n x_i - n \ln(\Gamma(\alpha)) \end{aligned}$$

Then take the first partial derivatives:

$$\begin{aligned} \frac{\partial l(\alpha, \lambda)}{\partial \alpha} &= n \ln(\lambda) + \sum_{i=1}^n \ln(x_i) - n \frac{\Gamma'(\alpha)}{\Gamma(\alpha)} \\ \frac{\partial l(\alpha, \lambda)}{\partial \lambda} &= \frac{n\alpha}{\lambda} - \sum_{i=1}^n x_i \end{aligned}$$

If we let $\psi(\alpha)$ and $\psi'(\alpha)$ be the first and second derivatives of $\ln(\Gamma(\alpha))$, then we can write the score vector as follows:

$$\begin{pmatrix} n \ln(\lambda) + \sum_{i=1}^n \ln(x_i) - n\psi(\alpha) \\ \frac{n\alpha}{\lambda} - \sum_{i=1}^n x_i \end{pmatrix}$$

Then we can get the Hessian matrix:

$$\begin{pmatrix} -n\psi'(\alpha) & \frac{n}{\lambda} \\ \frac{n}{\lambda} & -\frac{n\alpha}{\lambda^2} \end{pmatrix}$$

So the Fisher information matrix is:

$$\begin{pmatrix} n\psi'(\alpha) & -\frac{n}{\lambda} \\ -\frac{n}{\lambda} & \frac{n\alpha}{\lambda^2} \end{pmatrix}$$

The following function “newton_mle” implements the Newton-Raphson algorithm to compute the MLEs of α and λ based on x_1, \dots, x_n .

It also outputs an estimate of the variance-covariance matrix of the MLEs. (\$cov_matrix)

```
newton_mle <- function(x,eps=1.e-8,max.iter=50) {
  n <- length(x)
  alpha <- mean(x)^2/var(x) #estimate of alpha
  lambda <- mean(x)/var(x) #estimate of lambda
  theta <- c(alpha,lambda)
  # compute the scores based on the initial estimates
  score1 <- sum(log(x)) + n*(log(lambda) - digamma(alpha))
  score2 <- n*alpha/lambda-sum(x)
  score <- c(score1,score2) # score vector
  iter <- 1
  while (max(abs(score))>eps && iter<=max.iter) {
    # compute observed Fisher information
    info.11 <- n*trigamma(alpha)
    info.12 <- -n/lambda #same as the info.21
    info.22 <- n*alpha/lambda^2
    info <- matrix(c(info.11,info.12,info.12,info.22),ncol=2) # Fisher information matrix
    # Newton-Raphson iteration
    theta <- theta + solve(info,score)
    # update alpha and lambda
    alpha <- theta[1]
    lambda <- theta[2]
    iter <- iter + 1
    # update score vector
    score1 <- sum(log(x)) + n*(log(lambda) - digamma(alpha))
    score2 <- n*alpha/lambda-sum(x)
    score <- c(score1,score2)
  }
  if (max(abs(score))>eps) print("No convergence")
  else {
    print(paste("Number of iterations =",iter-1))
    loglik <- n*alpha*log(lambda) + (alpha -1)*sum(log(x)) - lambda*sum(x) - n*log(gamma(alpha))
    # log-likelihood of alpha and lambda
    info.11 <- n*trigamma(alpha)
    info.12 <- -n/lambda
    info.22 <- n*alpha/lambda^2
    info <- matrix(c(info.11,info.12,info.12,info.22),ncol=2) # Fisher information matrix
    cov_matrix <- solve(info) # estimated variance-covariance matrix
    r <- list(alpha=alpha,lambda=lambda,loglik=loglik,info=info,cov_matrix=cov_matrix)
    r
  }
}
```

Then we can test the above function on data generated from a Gamma distribution.

```
x <- rgamma(n=100, shape=1) # 100 observations from a gamma distribution with shape=1
r <- newton_mle(x) # use "newton_mle" estimates of alpha and lambda
```

```
## [1] "Number of iterations = 5"
```

```
r
```

```
## $alpha
## [1] 1.009051
##
## $lambda
## [1] 0.8832763
##
## $loglik
## [1] -113.3101
##
## $info
##          [,1]      [,2]
## [1,] 162.3438 -113.2149
## [2,] -113.2149 129.3361
##
## $cov_matrix
##          [,1]      [,2]
## [1,] 0.01581259 0.01384161
## [2,] 0.01384161 0.01984810
```