## Homework on Newton's methods

Leave your name and uni here

## Problem 1: Univariate optimizations.

In this problem, you will compare three common methods for univariate minimization: Newton's Method, Bisection Method and Golden-Section Search.

#### Function A:

$$f_1(x) = \ln(1+x^2) + x^2$$

#### Function B:

$$f_2(x) = x^4 - 6x^2 + 4x + 8$$

### Answer the following questions:

- 1. Computer f'(x) and f''(x)
- 2. Are  $f_1(x)$  and  $f_2(x)$  unimodal functions? If not, how many modes it contains?
- 3. Implementing Bisection, Golden Search and Newtown's methods to find global minimum of  $f_1(x)$  and  $f_2(x)$
- 4. Discuss your results which method is fastest in terms of iteration counts? which method is easist to apply if you only know a broad interval containing the mininum? which method fails or converges poorly if started badly? how the shape of the function influences the performance of these algorithms?
- 5. Please show all relevant R code for bisection, golden-section, and Newton's methods.

### Answer:

```
# 2.
# install.packages("rootSolve")
library(rootSolve)

f1 <- function(x) log(1 + x^2) + x^2
f1_first <- function(x) (2*x)/(1 + x^2) + 2*x
f1_second <- function(x) 2*(x^4 + x^2 + 2) / (1 + x^2)^2

f2 <- function(x) x^4 - 6*x^2 + 4*x + 8
f2_first <- function(x) 4*x^3 - 12*x + 4
f2_second <- function(x) 12*x^2 - 12
critical_points_f2 <- uniroot.all(f2_first, lower = -10, upper = 10)

# 3. find local minimum
# bisection</pre>
```

```
# we need dfa * dfb < 0
bisection <- function(df, a, b, tol = 1e-10){</pre>
  cur \leftarrow (a + b) / 2
  i <- 0
  while (abs(df(cur)) > tol){
    i <- i + 1
    if (df(a) * df(cur) > 0){
     a <- cur
   }else {
      b <- cur
    cur <- (a + b) / 2
  res <- c(cur, i)
  return(res)
}
# golden search
goldenSearch <- function(f, a, b, tol=1e-10){</pre>
  gr \leftarrow (sqrt(5) - 1) / 2
  x1 \leftarrow b - gr * (b - a)
  x2 \leftarrow a + gr * (b - a)
  i <- 0
  while(abs(b - a) > tol){
    i <- i + 1
   if (f(x1) < f(x2)){
      b <- x2
      x2 <- a + gr * (b - a) # can also be x1
      x1 \leftarrow b - gr * (b - a)
    }else{
      a <- x1
      x1 \leftarrow b - gr * (b - a) # can also be x2
      x2 <- a + gr * (b - a)
    }
  }
  res <- c((a + b) / 2, i)
 return (res)
# newton method
newtonMethod <- function(f, df, start, tol=1e-10){</pre>
 i <- 0
 x <- start
  while (abs(f(x)) > tol){
   i <- i + 1
   x \leftarrow x - f(x) / df(x)
 res \leftarrow c(x, i)
 return(res)
# f1
a <- -2
```

```
b <- 1
res_f1_bisec <- bisection(f1_first, a, b)
res_f1_gs <- goldenSearch(f1, a, b)
res_f1_nt <- newtonMethod(f1_first, f1_second, a)
# f2
as <-c(-2, 0.9)
bs <-c(0, 2)
res_f2_bisec1 <- bisection(f2_first, as[1], bs[1])
res_f2_bisec2 <- bisection(f2_first, as[2], bs[2])
res_f2_gs1 <- goldenSearch(f2, as[1], bs[1])
res_f2_gs2 <- goldenSearch(f2, as[2], bs[2])
res_f2_nt1 <- newtonMethod(f2_first, f2_second, as[1])
res_f2_nt2 <- newtonMethod(f2_first, f2_second, as[2])
comparison1 <- as.data.frame(rbind(c("bisection", res_f1_bisec),</pre>
                                    c("golden search", res_f1_gs),
                                    c("newton method", res_f1_nt)))
comparison1 <- comparison1 %>%
  rename(name = "V1", result = "V2", iteration = "V3")
comparison1
##
                                    result iteration
              name
## 1
         bisection 1.45519152283669e-11
                                                   35
## 2 golden search -5.79591141089835e-12
                                                   51
## 3 newton method 4.58039842491341e-16
                                                   5
comparison2 <- as.data.frame(rbind(c("bisection root 1", res_f2_bisec1),</pre>
                                     c("bisection root 2", res_f2_bisec2),
                                     c("golden search root 1", res_f2_gs1),
                                     c("golden search root 2", res_f2_gs2),
                                     c("newton method root 1", res_f2_nt1),
                                     c("newton method root 2", res_f2_nt2)))
comparison2 <- comparison2 %>%
  rename(name = "V1", result = "V2", iteration = "V3")
comparison2
##
                                       result iteration
                      name
## 1
         bisection root 1 -1.87938524156925
         bisection root 2 1.53208888624067
                                                      35
## 3 golden search root 1 -1.87938523016814
                                                      50
## 4 golden search root 2 1.53208889529803
                                                      49
## 5 newton method root 1 -1.87938524157182
                                                       4
## 6 newton method root 2 1.53208888623796
                                                       7
  1. function A: f_1'(x) = \frac{2x}{1+x^2} + 2x, f_1''(x) = \frac{2(x^4+x^2+2)}{(1+x^2)^2}
function B: f_2'(x) = 4x^3 - 12x + 4, f_2''(x) = 12x^2 - 12
```

2. To find whether functions are unimodal, we need to find root where first prime of function is 0 and second prime of function is positive.

For function A, it is obvious that the first derivative of function is 0 only when x = 0, and  $f_2''(0) = 4 > 0$ .

Therefore, function A is unimodal.

For function B, we use library(rootSolve) and find 3 root for function B. is not unimodal.

3. The comparison of results for function A and B are presented above in tables.

By question 2, we know that root of function B are: -1.8794479, 0.3472831, 1.5320901, and their second derivatives are 30.3878946, -10.5527331, 16.1676027. Therefore, we only have 2 local minimum.

The newton's method is fastest in terms of iteration.

I think the Golden Search is the easiest method to apply if we only know a broad range because we don't need to know the expression of derivatives of function. Also, bisection and newton methods are root finding methods, but for golden search, we can implement it to obtain either maximum or minimum.

Newton's Method is most prone to failure or poor convergence when initial guess is bad. if initial guess is too far or its second derivative is too close to 0, then the method might diverge to another search region or crash(denominator=0).

Shape of function influences:

- Bisection because it requires the signs of first derivative of the range we choose to be different. Therefore, if we don't properly choose the interval, other critical points might be included.
- Golden Search because it assumes unimodality of our function. If there is multiple modals, we need to
  isolate each of them.
- Newton Method because the second derivative of the function(local curvature) can influence convergence significantly.

### Problem 2: Newton's Method in Two Dimensions

q(x,y) is a 2D function

$$g(x,y) = x^2 + xy + y^2 - 4x - 3y + 7$$

## Answer the folling questions:

- 1. Derive the gradient  $\Delta g(x,y)$  and the Hessian matrix  $\Delta^2 g(x,y)$ .
- 2. Implementation a Newton's algorithm to find its minimizer.
- 3. Choose two different starting values, and compare the resulting solutions, iteration counts and path to convergence
- 4. Create a coutour plot of g around its minium, and overlay the sequence of iterates from Newton's method to show the path to the minimum.

# Answer: your answer starts here...

```
1. \Delta g(x,y) = \binom{2x+y-4}{x+2y-3}, \Delta^2 g(x,y) = \begin{pmatrix} 2 & 1 \\ 1 & 2 \end{pmatrix}
```

```
y <- y0
  hess_inv <- solve(hess)
  g_val <- grad(x, y)</pre>
  res \leftarrow c(x, y, g_val, i)
  prev_x <- -Inf
  prev_y <- -Inf</pre>
  while(i < max_iter && (x-prev_x > tol) && (y-prev_y > tol)){\#sqrt(sum(g_val^2)) > tol){
    i <- i + 1
    curr <- hess_inv %*% g_val</pre>
    prev_x <- x
    prev_y <- y
    x \leftarrow x - curr[1]
    y <- y - curr[2]
    g_val <- grad(x, y)</pre>
    res <- rbind(res, c(x, y, g_val, i))
  return(res)
}
res_table1 <- newtonRalphson(0.8, 0.7, grad, hess)
res_table2 <- newtonRalphson(10, -5, grad, hess)
res_table1
##
            [,1]
                       [,2] [,3] [,4] [,5]
## res 0.800000 0.7000000 -1.7 -0.8
       1.666667 0.6666667 0.0 0.0
res_table2
##
                                       [,3] [,4] [,5]
             [,1]
                         [,2]
## res 10.000000 -5.0000000 1.100000e+01
        1.666667 0.6666667 1.776357e-15
                                                     1
```

### Problem 3

Suppose we have data  $(x_i, y_i), i = 1, ..., n$ , with  $y_i$  follows a conditional expotential distribution

$$Y_i \mid x_i \sim \exp(\lambda_i)$$
, where  $\log(\lambda_i) = \alpha + \beta x_i$ .

### Please complete the following tasks:

- 1. Derive the log-likelihood of  $(x_i, y_i)$ , as well as its Gradient and Hession Matrix
- 2. Generate a synthesic data with true  $\alpha = 0.5$ , true  $\beta = 1.2$  and sample size n = 200.
- 3. Implement an Newton's algorithm to its MLE
- 4. Implement a modified Newton's algorithm, where you incoporate both of the step-having and ascent direction check. If a direction is not ascent, you can swith to a simpler gradient descent.
- 5. For a generalized linear models, one can replace the observed Hessian with the expected Hessian (the Fisher information), which might lead to stable updates akin to a Fisher scoring approach. Implement another modified newton's algorithm with Fisher scoring toestimate MLE.

6. Compare the final parameter estimates, iteration counts and convergences of optimization, and summarize your findings.

### Answer:

1. Since the conditional distribution is exponential, we use its pdf and then we have  $L(\alpha, \beta) = ln \prod_{i=1}^{n} [\lambda_i] e^{-\lambda_i y_i} = \sum_{i=1}^{n} [ln(\lambda_i) - \lambda_i y_i] = \sum_{i=1}^{n} [\alpha + \beta x_i - y_i e^{\alpha + \beta x_i}]$ 

```
Gradient: \nabla L(\alpha, \beta) = \begin{pmatrix} n - \sum_{i=1}^{n} y_i \, e^{\alpha + \beta x_i} \\ \sum_{i=1}^{n} x_i - \sum_{i=1}^{n} (x_i \, y_i \, e^{\alpha + \beta x_i}) \end{pmatrix}.

Hessian: \nabla^2 L(\alpha, \beta) = \begin{pmatrix} -\sum_{i=1}^{n} y_i \, e^{\alpha + \beta x_i} & -\sum_{i=1}^{n} x_i \, y_i \, e^{\alpha + \beta x_i} \\ -\sum_{i=1}^{n} x_i \, y_i \, e^{\alpha + \beta x_i} & -\sum_{i=1}^{n} x_i^2 \, y_i \, e^{\alpha + \beta x_i} \end{pmatrix}.
# 2
n <- 200
alpha_true <- 0.5
beta_true <- 1.2
set.seed(123)
# Generate predictor values (x i)
x \leftarrow rnorm(n)
lambda <- exp(alpha_true + beta_true * x)</pre>
# Generate y_i ~ Exp(rate = lambda_i)
y <- rexp(n, rate = lambda)
df \leftarrow data.frame(x = x, y = y)
# 3. Newton method for MLE of conditional dist
loglik_expreg <- function(alpha, beta, x, y){</pre>
   lambda <- exp(alpha + beta * x)</pre>
   val <- sum((alpha + beta * x) - y * lambda)</pre>
   return (val)
}
grad_expreg <- function(alpha, beta, x, y) {</pre>
   lambda <- exp(alpha + beta * x)
   d_alpha <- length(x) - sum(y * lambda)</pre>
   d_{beta} \leftarrow sum(x) - sum(x * y * lambda)
   return(c(d_alpha, d_beta)) # 2x1 vector
hess_expreg <- function(alpha, beta, x, y) {
   lambda <- exp(alpha + beta * x)</pre>
   term <- y * lambda # common term y_i e^{alpha+beta x_i}
   h11 <- -sum(term)
                                              # d^2/dalpha^2
   h22 \leftarrow -sum(x^2 * term)
                                            # d^2/dbeta^2
   h12 \leftarrow -sum(x * term)
                                               # d^2/dalpha dbeta = d^2/dbeta dalpha
   H <- matrix(c(h11, h12,</pre>
                        h12, h22),
                     nrow = 2, byrow = TRUE)
   return(H)
```

```
}
newton_expreg <- function(x, y, alpha_init = 0, beta_init = 0,</pre>
                            tol = 1e-8, max_iter = 100) {
  alpha <- alpha_init</pre>
  beta <- beta_init
  i <- 0
  prev_loglik <- -Inf</pre>
  curr_loglik <- loglik_expreg(alpha, beta, x, y)</pre>
  while (i < max_iter && abs(curr_loglik - prev_loglik) > tol) { # or compute gradient norm sqrt(sum(g^
    i <- i+1
    g <- grad_expreg(alpha, beta, x, y)</pre>
    H <- hess_expreg(alpha, beta, x, y)</pre>
    # Update alpha, beta via Newton step
    update <- solve(H, g) # H^{-1} * g
    alpha <- alpha - update[1]</pre>
    beta <- beta - update[2]
    # Update log-likelihood
    prev_loglik <- curr_loglik</pre>
    curr_loglik <- loglik_expreg(alpha, beta, x, y)</pre>
  return (list(alpha = alpha,
               beta = beta,
               loglik = curr_loglik,
               iter = i))
}
res_newtonRalphson <- newton_expreg(x, y,</pre>
                             alpha_init = 0, beta_init = 0,
                             tol = 1e-10, max_iter = 200)
# 4. step halving at the end when ascent direction
modified_newton_expreg <- function(x, y,</pre>
                                      alpha_init = 0,
                                     beta_init = 0,
                                     tol = 1e-8,
                                     max_iter = 100) {
  alpha <- alpha_init
  beta <- beta init
        <- 0
  prev_loglik <- -Inf</pre>
  curr_loglik <- loglik_expreg(alpha, beta, x, y)</pre>
  while (i < max_iter && abs(curr_loglik - prev_loglik) > tol) {
    i \leftarrow i + 1
    # Compute gradient and Hessian
    g <- grad_expreg(alpha, beta, x, y)
                                             # a 2-vector
```

```
H <- hess_expreg(alpha, beta, x, y) # a 2x2 matrix</pre>
    # Newton update is effectively -H^-1 * g
    update <- solve(H, g)</pre>
    # The "Newton direction" in terms of "move" is -update:
    d_newton <- -update</pre>
    # Check if d newton is ascent
    dir_check <- sum(g * d_newton)</pre>
    if (dir check <= 0) {</pre>
      # fallback: use gradient direction
      d <- g
    } else {
      d <- d_newton
    # Step-halving line search
    step <- 1
    old_loglik <- curr_loglik</pre>
    # Proposed new parameters
    new_alpha <- alpha + step * d[1]</pre>
    new_beta <- beta + step * d[2]</pre>
    new_loglik <- loglik_expreg(new_alpha, new_beta, x, y)</pre>
    # While new log-likelihood <= old, reduce step
    while (step > 1e-14 && new_loglik <= old_loglik) {</pre>
      step <- step / 2
      new_alpha <- alpha + step * d[1]</pre>
      new_beta <- beta + step * d[2]</pre>
      new_loglik <- loglik_expreg(new_alpha, new_beta, x, y)</pre>
    # Update alpha, beta, loglik
    prev_loglik <- curr_loglik</pre>
    alpha <- new_alpha
    beta <- new_beta
    curr_loglik <- new_loglik</pre>
  list(alpha = alpha,
       beta = beta,
       loglik = curr_loglik,
       iter = i)
}
# 5. Fisher score modified
newton_expreg_fisher <- function(x, y,</pre>
                                    alpha_init = 0, beta_init = 0,
                                    tol = 1e-8, max_iter = 100) {
  # Initialize parameters
  alpha <- alpha_init
  beta <- beta_init</pre>
```

```
i <- 0
  prev_loglik <- -Inf</pre>
  curr loglik <- loglik expreg(alpha, beta, x, y)
  while (i < max_iter && abs(curr_loglik - prev_loglik) > tol) {
    i <- i + 1
    g <- grad_expreg(alpha, beta, x, y)</pre>
    # Compute the Fisher Information matrix.
    I_mat <- matrix(c(length(x), sum(x),</pre>
                       sum(x),
                                    sum(x^2),
                    nrow = 2, byrow = TRUE)
    # Compute the Fisher scoring update:
    # update = I^{-1} * g. (For maximization, we add this update.)
    update <- solve(I_mat, g)</pre>
    # Step-halving (line search) to ensure an increase in log-likelihood.
    step <- 1
    old_loglik <- curr_loglik</pre>
    new_alpha <- alpha + step * update[1]</pre>
    new_beta <- beta + step * update[2]</pre>
    new_loglik <- loglik_expreg(new_alpha, new_beta, x, y)</pre>
    while (step > 1e-14 && new loglik < old loglik) {
      step <- step / 2
      new_alpha <- alpha + step * update[1]</pre>
      new_beta <- beta + step * update[2]</pre>
     new_loglik <- loglik_expreg(new_alpha, new_beta, x, y)</pre>
    }
    # Update the parameters and log-likelihood
    prev_loglik <- curr_loglik</pre>
    alpha <- new_alpha
    beta <- new_beta
    curr_loglik <- new_loglik</pre>
 return(list(alpha = alpha,
              beta = beta,
              loglik = curr_loglik,
              iter = i))
}
res_orig <- newton_expreg(x, y, alpha_init = 0, beta_init = 0, tol = 1e-10, max_iter = 200)
           <- modified_newton_expreg(x, y, alpha_init = 0, beta_init = 0, tol = 1e-10, max_iter = 200)</pre>
res_fisher <- newton_expreg_fisher(x, y, alpha_init = 0, beta_init = 0, tol = 1e-10, max_iter = 200)
# Create a comparison table
results <- data.frame(Method = c("Original Newton", "Modified Newton", "Fisher Scoring"),
                       Alpha = c(res_orig$alpha, res_mod$alpha, res_fisher$alpha),
                       Beta = c(res_orig$beta, res_mod$beta, res_fisher$beta),
```

```
## 1 Original Newton 0.5441476 1.153412 -93.14753 6 6 ## 2 Modified Newton 0.5441476 1.153412 -93.14753 6 ## 3 Fisher Scoring 0.5441476 1.153412 -93.14753 7
```