

Newton-Raphson Method

Code

```
nextX <- function(x) {  
  m=3  
  c=5  
  x = x*(1 - 1/m + c/(m*x^m))  
  return(x)  
}  
  
newt_raphson <- function(x.i, x.next, condition) {  
  x.i = x.next  
  x.next = nextX(x.i)  
  if(abs(x.i - x.next) <= condition) {  
    return(x.next)  
  }  
  else {  
    newt_raphson(x.i, x.next, condition)  
  }  
}  
  
x0 = 5  
condition = 0.000000001  
root = newt_raphson(x0, nextX(x0), condition)
```

Results

```
#From Newton-Raphson  
root  
#[1] 1.709976
```

```
5^(1/3)  
#[1] 1.709976
```

Logistic Regression with Metropolis

Code

#printPDF function receives two vectors and a filename. It prints three plots for each vector to the specified filename: ts plot, hist, acf.

```
printPDF <- function(v1,v2,filename) {
```

```
  pdf(filename)
```

```
  #PLOTS to PDF
```

```
    par(mfrow=c(3,2)) #split plotting window into 3 rows and 2 columns
```

```
    ts.plot(v1,xlab="Iterations")
```

```
    ts.plot(v2,xlab="Iterations")
```

```
    hist(v1,probability=T, cex.lab=1.5, cex.axis=1.5)
```

```
    hist(v2,probability=T, cex.lab=1.5, cex.axis=1.5)
```

```
    acf(v1, lag.max=500)
```

```
    acf(v2, lag.max=500)
```

```
  dev.off()
```

```
}
```

#metro function receives nine parameters: two vectors "x.s" and "y.s" for the data values of interest, "a0" for the initial alpha, "b0" for the initial beta, "k" for the alpha interval, "c" for the beta interval, "N" for number of independent random normal realizations, "lag" for determining how many realizations to skip between saves, and "burnin" for determining how many realizations to skip before starting to save.

```
metro <- function(x.s,y.s,a0,b0,k,c,N,lag,burnin) {
```

```
  #Set N to be N*lag+burnin
```

```
  N <- N*lag + burnin
```

```
  #Initialize vectors to hold alpha (a.v) and beta (b.v) values
```

```
  a.s <- NULL
```

```
  b.s <- NULL
```

```
  #store the acceptance rate for alpha (a.cnt) and beta (b.cnt)
```

```
  a.cnt = 0
```

```
  b.cnt = 0
```

```
  for(i in 1:N) {
```

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#Generate an alpha star 'a.star' from the proposal density (Normal) using a0 as mu and k as std dev

```
a.star = rnorm(1,a0,k)
```

#Compute probability for alpha using Normal priors with mean (0), variance (100) -
- Use full conditional (from joint posterior which is

likelihood*prior(alpha)*prior(beta)) -- compute ratio of full conditional given a.star to the full conditional given a0

```
numerator = sum(y.s*a.star) - sum(log(1 + exp(a.star + b0*x.s))) -  
a.star^2/200
```

```
denominator = sum(y.s*a0) - sum(log(1 + exp(a0 + b0*x.s))) - a0^2/200
```

```
target.ratio = exp(numerator - denominator)
```

#Use target.ratio to determine if a.star should be accepted

```
param = 0
```

```
accept.code = 0
```

```
if(target.ratio < 1) {
```

```
  if(runif(1,0,1) < target.ratio) {
```

```
    param = a.star
```

```
    accept.code = 1
```

```
  }
```

```
  else {
```

```
    param = a0
```

```
    accept.code = 0
```

```
  }
```

```
}
```

```
else {
```

```
  param = a.star
```

```
  accept.code = 1
```

```
}
```

#if i is greater than burnin and if i is a multiple of the lag, store value

#add accept.code to total a.cnt - accept.code will be 1 if a.star was accepted and 0 otherwise

#add accepted value to vector - a0 or a.star - stored as param

```
if(i > burnin) {
```

```
  if(i %% lag == 0) {
```

```
    a.cnt = a.cnt + accept.code
```

```
    a.s = c(a.s, param)
```

#set a0 equal to param (parameter that was stored in vector -- a.star or a0) for next iteration

```
    a0 = param
```

```
}
```

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```
}
```

```
#Reset target.ratio, numerator, denominator
```

```
numerator = 0
```

```
denominator = 0
```

```
target.ratio = 0
```

```
#Generate a beta star 'b.star' from the proposal density (Normal) using b0 as mu  
and c as std dev
```

```
    b.star = rnorm(1,b0,c)
```

```
#Compute probability for beta using Normal priors with mean (0), variance (100) --
```

```
Use full conditional (from joint posterior which is
```

```
likelihood*prior(alpha)*prior(beta)) -- compute ratio of full conditional given b.star  
to the full conditional given b0
```

```
    numerator = sum(y.s*b.star*x.s) - sum(log(1 + exp(a.star + b.star*x.s))) -  
b.star^2/200
```

```
    denominator = sum(y.s*b0*x.s) - sum(log(1 + exp(a.star + b0*x.s))) -  
b0^2/200
```

```
    target.ratio = exp(numerator - denominator)
```

```
#Use target.ratio to determine if b.star should be accepted
```

```
param = 0
```

```
accept.code = 0
```

```
if(target.ratio < 1) {
```

```
    if(runif(1,0,1) < target.ratio) {
```

```
        param = b.star
```

```
        accept.code = 1
```

```
    }
```

```
    else {
```

```
        param = b0
```

```
        accept.code = 0
```

```
    }
```

```
}
```

```
else {
```

```
    param = b.star
```

```
    accept.code = 1
```

```
}
```

```
#if i is greater than burnin and if i is a multiple of the lag, store value
```

```
#add accept.code to total b.cnt - accept.code will be 1 if b.star was accepted and 0  
ow
```

```
#add accepted value to vector - b0 or b.star - stored as param
```

```
    if(i > burnin) {
```

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```
        if(i %% lag == 0) {
            b.cnt = b.cnt + accept.code
            b.s = c(b.s, param)
            #set b0 equal to param (parameter that was stored in vector -- b.star
or b0) for next iteration
            b0 = param
        }
    }
}
```

```
filename = sprintf("Documents/R-FILES/HW9-lag-%s-k-%s-c-%s.pdf",lag,k,c)
printPDF(a.s,b.s,filename)
```

```
vectors <- list("alpha" = a.s, "aCount" = a.cnt, "beta" = b.s, "bCount" = b.cnt)
return(vectors)
}
```

```
#Set metro parameters
```

```
N = 5000
```

```
lag = 500
```

```
burnin = 0
```

```
k=.5
```

```
c=.011
```

```
a0 = -3.82
```

```
b0 = .086
```

```
x.s = c(21,24,25,26,28,31,33,34,35,37,43,49,51,55,25,29,43,44,46,46,51,55,56,58)
```

```
y.s = c(0,0,0,0,0,0,0,0,0,0,0,0,0,0,1,1,1,1,1,1,1,1,1,1)
```

```
#Obtain alpha and beta from data to use for a0 and b0
```

```
#output = summary(glm(y.s~x.s,family=binomial))
```

```
vectors = metro(x.s,y.s,a0,b0,k,c,N,lag,burnin)
```

```
#Obtain acceptance probabilities
```

```
alpha.accept = vectors$aCount/N
```

```
beta.accept = vectors$bCount/N
```

```
alpha.accept
```

```
#[1] 0.6864
```

```
beta.accept
```

```
#[1] 0.6946
```

```
#Obtain credible intervals for alpha and beta
```

```
quantile(vectors$alpha, 0.025)
```

```
#-9.030698
```

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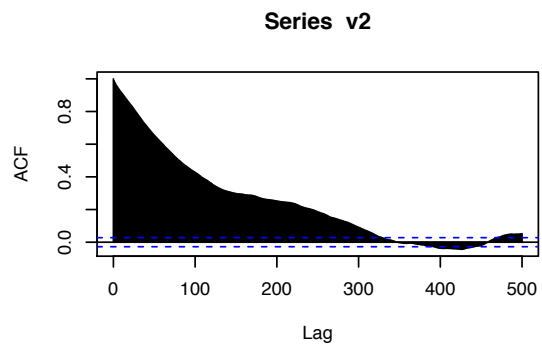
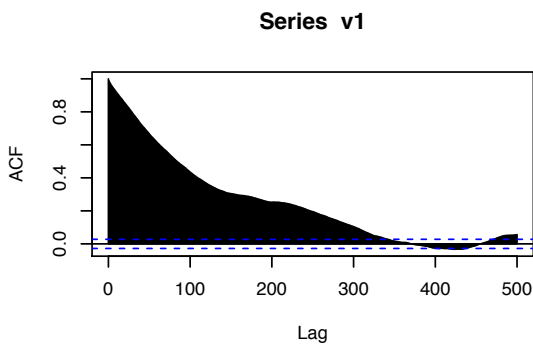
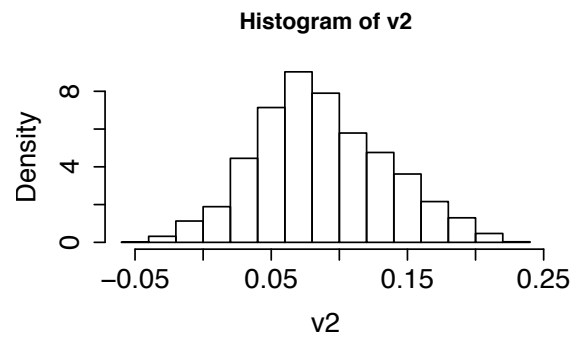
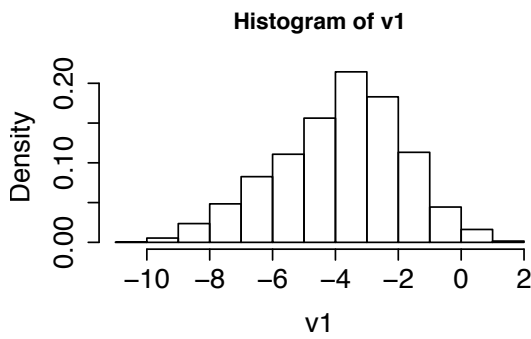
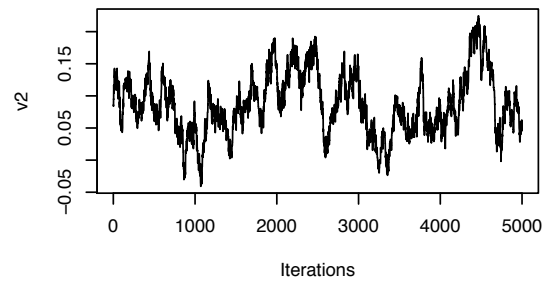
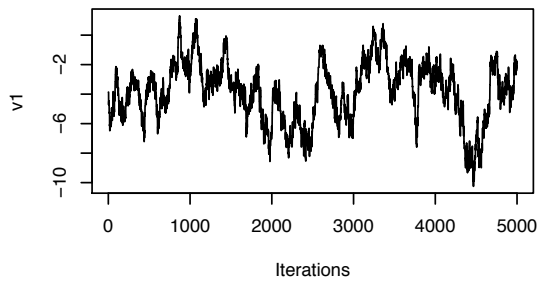
```
quantile(vectors$alpha, 0.975)  
#-0.1419091  
quantile(vectors$beta, 0.025)  
#-0.0021883  
quantile(vectors$beta, 0.975)  
#0.2096534
```

#Produce a scatterplot of x and y data with posterior mean curve and credible
bounds superposed on the plot

```
x = seq(min(x.s),max(x.s),length=250)  
m <- matrix(, nrow = 5000, ncol = 250)  
for(i in 1:250){  
  vector = exp(vectors$alpha + vectors$beta*x[i])/(1+exp(vectors$alpha +  
vectors$beta*x[i]))  
  m[, i] <- vector  
}  
m.mean = apply(m, 2, mean)  
q97.5 = apply(m, 2, quantile, probs=0.975)  
q2.5 = apply(m, 2, quantile, probs=0.025)  
  
plot(x.s,y.s)  
lines(x, m.mean)  
lines(x, q97.5)  
lines(x, q2.5)
```

Results

- $\pi(\alpha, \beta | (\vec{x}, \vec{y})) = \frac{1}{200\pi} e^{\sum_{i=1}^n y_i \alpha + \sum_{i=1}^n y_i \beta x_i - \sum_{i=1}^n \ln(1 + e^{\alpha + \beta x_i}) - \frac{\alpha^2}{200} - \frac{\beta^2}{200}}$
- Trace Plots



- Credible Intervals

Credible Interval for α
(-9.030698, -0.1419091)

Credible Interval for β
(-0.0021883, 0.2096534)

- Scatterplot with posterior mean curve and credible bounds superposed

