Lab 2 Ravinder

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1. Linear and polynomial regression

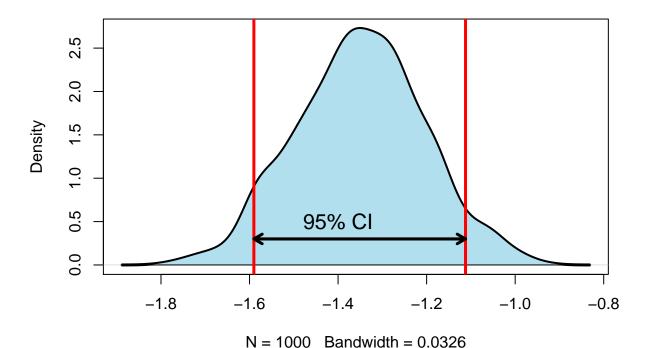
```
library(mvtnorm)
# Scaled Inverse chi square distribution
pdfinvchisquare <- function(x,n,tau_sq){</pre>
  res <- (((tau_sq*(n-1))/2)^(n-1)/2 * exp(-(tau_sq*(n-1))/(2*x)))/((x^(1+(n-1)/2)) * gamma((n-1)/2))
  return(res)
}
# Function for sampling from inverse chi square distribution
rinvchisquare <- function(num_draws, n, tau_sq){</pre>
  set.seed(1234)
  x <- rchisq(num_draws,df = n-1)</pre>
  x_{inv} \leftarrow ((n-1)*tau_{sq})/x
  return(x_inv)
# Prior parameters initialization
sisquare0 <- 1
v0 <- 4
mu0 <- c(-10, 100, -100)
si0 <- 0.01*diag(3)
si0_inv <- solve(si0)</pre>
num_draws = 1000
# Prior hierarchy
# Prior 1: si^2
var_prior <- rinvchisquare(num_draws, v0, sisquare0)</pre>
# Prior 2: beta/si^2
beta_prior <- matrix(NA, nrow = num_draws, ncol = 3)</pre>
temp <- c()
for(i in 1:num draws){
  beta_prior[i,] <- rmvnorm(1,mean = mu0, sigma = var_prior[i]*si0_inv)</pre>
  \#temp[i] \leftarrow
}
```

2. Posterior approximation for classification with logistic regression

```
ww_data <- read.table('WomenWork.dat', header = TRUE)</pre>
rows <- nrow(ww_data)</pre>
cols <- ncol(ww_data)</pre>
y <- as.matrix(ww_data[1])</pre>
X <- as.matrix(ww_data[,2:cols])</pre>
params \leftarrow dim(X)[2]
mu <- as.matrix(rep(0,params))</pre>
tau = 10
Sigma = (tau^2)*diag(params)
LogPostLogistic <- function(betas,y,X,mu,Sigma){</pre>
  linPred <- X%*%betas;</pre>
  logLik <- sum(linPred*y - log(1 + exp(linPred)))</pre>
  logPrior <- dmvnorm(betas, mu, Sigma, log=TRUE)</pre>
  return(logLik + logPrior)
initValue <- matrix(0,params,1)</pre>
OptimRes <- optim(initValue,
                   LogPostLogistic, gr=NULL, y, X, mu, Sigma, method=c("BFGS"),
                   control=list(fnscale=-1), hessian=TRUE)
print('Posterior Mode: ')
## [1] "Posterior Mode: "
print(OptimRes$par)
##
                [,1]
## [1,] 0.62672884
## [2,] -0.01979113
## [3,] 0.18021897
## [4,] 0.16756670
## [5,] -0.14459669
## [6,] -0.08206561
## [7,] -1.35913317
## [8,] -0.02468351
print('Inverse of hessian matrix')
## [1] "Inverse of hessian matrix"
inversehessian <- solve(OptimRes$hessian)</pre>
print(inversehessian)
                 [,1]
                                [,2]
                                               [,3]
                                                              [,4]
## [1,] -2.266022568 -3.338861e-03 6.545121e-02 1.179140e-02 -0.0457807243
## [2,] -0.003338861 -2.528045e-04 5.610225e-04 3.125413e-05 -0.0001414915
## [3,] 0.065451206 5.610225e-04 -6.218199e-03 3.558209e-04 -0.0018962893
## [4,] 0.011791404 3.125413e-05 3.558209e-04 -4.351716e-03 0.0142490853
## [5,] -0.045780724 -1.414915e-04 -1.896289e-03 1.424909e-02 -0.0555786706
```

```
## [6,] 0.030293450 3.588562e-05 3.240448e-06 1.340888e-04 0.0003299398
## [7,]
        0.188748354 -5.066847e-04 6.134564e-03 1.468951e-03 -0.0032082535
## [8,]
        0.098023929 1.444223e-04 -1.752732e-03 -5.437105e-04 -0.0005120144
                 [,6]
                                [,7]
##
                                              [,8]
## [1,]
        3.029345e-02 0.1887483542 0.0980239285
## [2,]
        3.588562e-05 -0.0005066847 0.0001444223
## [3.]
        3.240448e-06 0.0061345645 -0.0017527317
## [4,]
         1.340888e-04 0.0014689508 -0.0005437105
## [5,] 3.299398e-04 -0.0032082535 -0.0005120144
## [6,] -7.184611e-04 -0.0051841611 -0.0010952903
## [7,] -5.184161e-03 -0.1512621814 -0.0067688739
## [8,] -1.095290e-03 -0.0067688739 -0.0199722657
beta_post <- OptimRes$par</pre>
names(beta_post) <- colnames(ww_data[,2:cols])</pre>
approx_par_NSC <- rnorm(1000,beta_post['NSmallChild'],-inversehessian[7,7])
lowerInterval <- quantile(approx_par_NSC,0.05)</pre>
upperInterval <- quantile(approx_par_NSC, 0.95)</pre>
plot(density(approx_par_NSC), lwd = 3, main = '95% Posterior Probability Interval of NSmallChild variable
polygon(density(approx_par_NSC), col = 'lightblue2')
abline(v = lowerInterval, col = 'red', lwd = 3)
abline(v = upperInterval, col = 'red', lwd = 3)
arrows(lowerInterval, 0.3, upperInterval, 0.3, length = 0.1, col = 'black', lwd = 3)
arrows(upperInterval, 0.3, lowerInterval, 0.3, length = 0.1, col = 'black', lwd = 3)
text(-1.4,0.5, '95\% CI', lwd = 3, cex = 1.3)
```

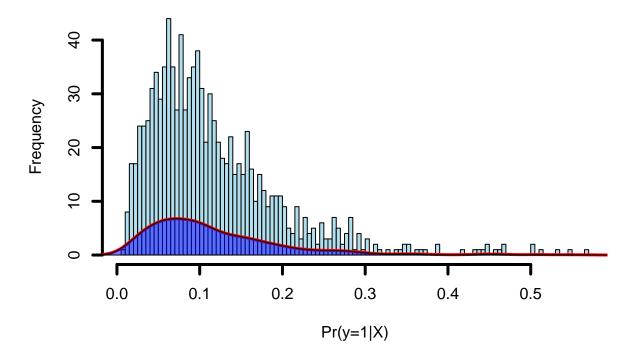
95% Posterior Probability Interval of NSmallChild variable



b.

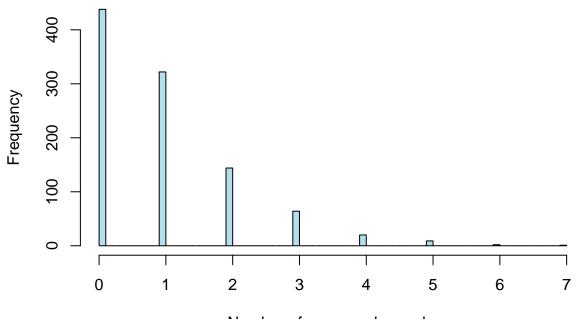
```
posteriorPredictive <- function(x, beta_post, inversehessian){</pre>
  post_sample <- rmvnorm(1, beta_post,-inversehessian)</pre>
  #print(x)
  #print(x %*% t(post_sample))
  logist_prob <- (exp(x %*% t(post_sample)))/(1 + exp(x %*% t(post_sample)))</pre>
  #print(logist_prob)
  return(logist_prob)
x \leftarrow c(1,13, 8, 11, (11/10)^2, 37, 2, 0)
nsamples = 1000
post_predict <- c(rep(0,nsamples))</pre>
for(i in 1:nsamples){
  post_predict[i] <- posteriorPredictive(x, beta_post, inversehessian)</pre>
#print(post_predict)
hist(post_predict, breaks = 100,
     col = 'lightblue2', lwd = 3,
     xlab = 'Pr(y=1|X)',
     main = 'Posterior Predictive Plot')
lines(density(post_predict), col = 'red', lwd = 3)
polygon(density(post_predict), col = rgb(red = 0, green = 0, blue = 1, alpha = 0.5))
```

Posterior Predictive Plot



c.

Histogram of number of women who works



Number of women who works