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4. Horseshoe Prior

Findings:

1. By changing the prior to horseshoe prior, the acceptance rate for beta decreases a lot.
2. Try many hyper-parameter taus, but hard to increase the acceptance rate.

The horseshoe prior is as follows (from Juho & AKi paper):

$$\beta_j | \lambda_j, \tau \sim N(0, \tau^2 \lambda_j^2)$$

$$\lambda_j \sim C^+(0, 1), \quad j = 1, \dots, p$$

In our function, we define:

$$\sigma_j = \tau \lambda_j$$

```
MH_horseshoe_Sampling <- function(Y,delta,tau,
                                   A,beta0,sigma0,var.prop,
                                   m,B,eta,
                                   Wmat_option=0){

  accept_beta = 0
  accept_lambda = 0
  beta = beta0
  lambda = lambda0
  sigma = sigma0

  # What we want to record
  BETA = matrix(0,m,dim(A)[2])
  LAMBDA = matrix(0,m,dim(A)[2])
  ThetaRecord <- matrix(0, m, length(Y))
  C_stat = c()
```

```

# For safety m>B
if (B>m){
  B = 0
}

# 0 means we use Harrell C statistics
# 1 means we use Uno C statistics
if (Wmat_option==0){
  Wmat <- HarrellC_Wmat(Y, delta, tau)
}else if (Wmat_option==1){
  Wmat <- UnoC_Wmat(Y, delta, tau)
}else{ # Other Possible C index...
  Wmat <- HarrellC_Wmat(Y, delta, tau)
}

for (i in 1:m){

  # Sample beta from proposal distribution
  beta.p = t(rmvnorm(1,beta,var.prop))

  # Compute theta from current and last iteration
  theta.p = THETA(A,beta.p)
  theta = THETA(A,beta)

  # Record theta from last iteration
  ThetaRecord[i,] <- theta

  # Compute C-statistics from current and last iteration
  HC.p = HarrellC(theta.p, Wmat)
  HC = HarrellC(theta, Wmat)

  # Record C-statistics from last iteration
  C_stat = c(C_stat,HC)

  #####
  # Compute log of MH ratio

  lrMH = eta*log(HC.p) +
    sum(dnorm(beta.p,beta0,sigma,log=T))-
    eta*log(HC) -
    sum(dnorm(beta,beta0,sigma,log=T))

  if (log(runif(1))<lrMH){
    beta = beta.p
    accept_beta = accept_beta + 1
  }
  BETA[i,] = beta
  #####

```

```
#####
# Compute log of MH_lambda ratio
lambda.p = exp(t(rnorm(dim(A)[2],log(lambda),rep(1,dim(A)[2]))))
sigma.p = lambda.p*beta_tau

lrMH_lambda = sum(dnorm(beta,beta0,sigma.p,log=T))-
               sum(dnorm(beta,beta0,sigma,log=T))

if (log(runif(1))<lrMH_lambda){
  lambda = lambda.p
  sigma = sigma.p
  accept_lambda = accept_lambda + 1
}
LAMBDA[i,] = lambda
}
#####

if (B == 0){
  return(list(BETA=BETA,
             LAMBDA = LAMBDA,
             accept_beta=accept_beta/m,
             accept_lambda=accept_lambda/m,
             THETA = ThetaRecord,
             C_stat = C_stat))
}else{
  return(list(BETA=BETA[-c(1:B),],
             LAMBDA = LAMBDA[-c(1:B),],
             accept_beta=accept_beta/m,
             accept_lambda=accept_lambda/m,
             THETA = ThetaRecord[-c(1:B),],
             C_stat = C_stat[-c(1:B)]))
}

}
```

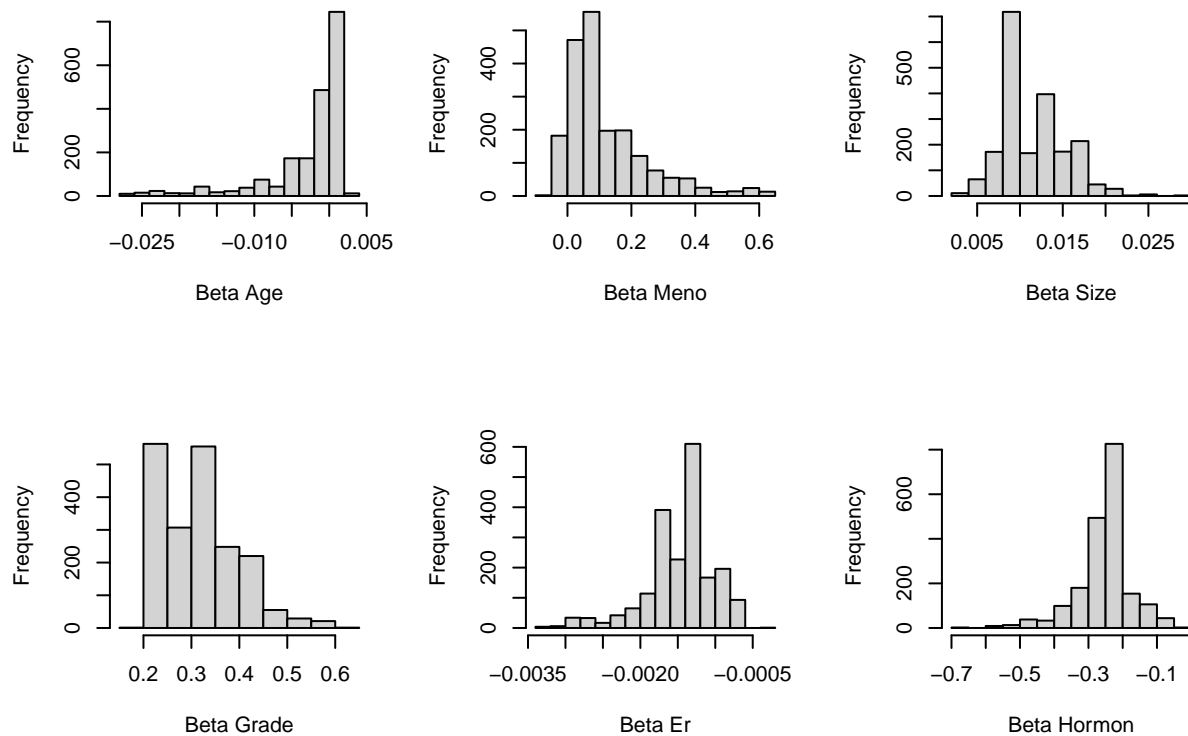
```
m = 2200
B = 200

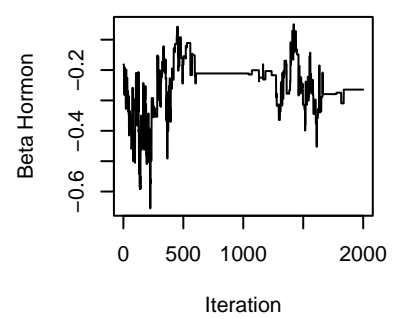
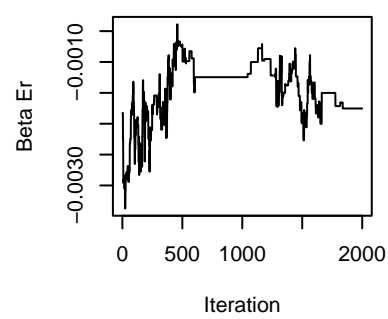
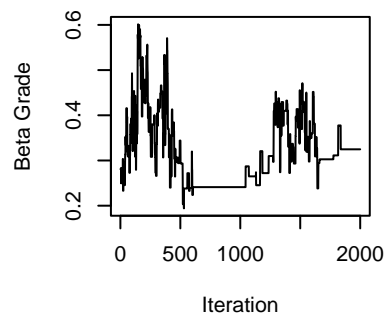
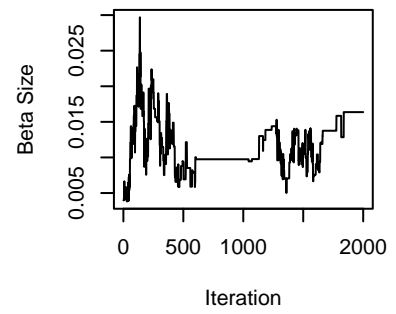
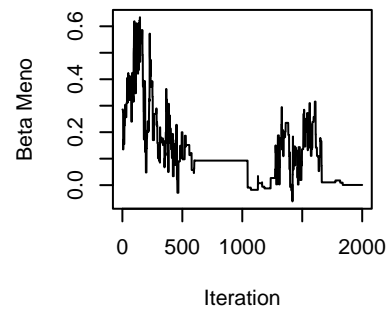
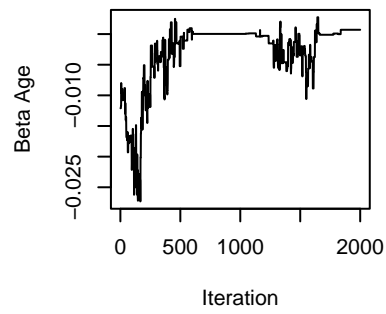
beta0 = rep(0,dim(A)[2])
lambda0 = rep(1,dim(A)[2])
beta_tau = 100000000
sigma0 = beta_tau*lambda0

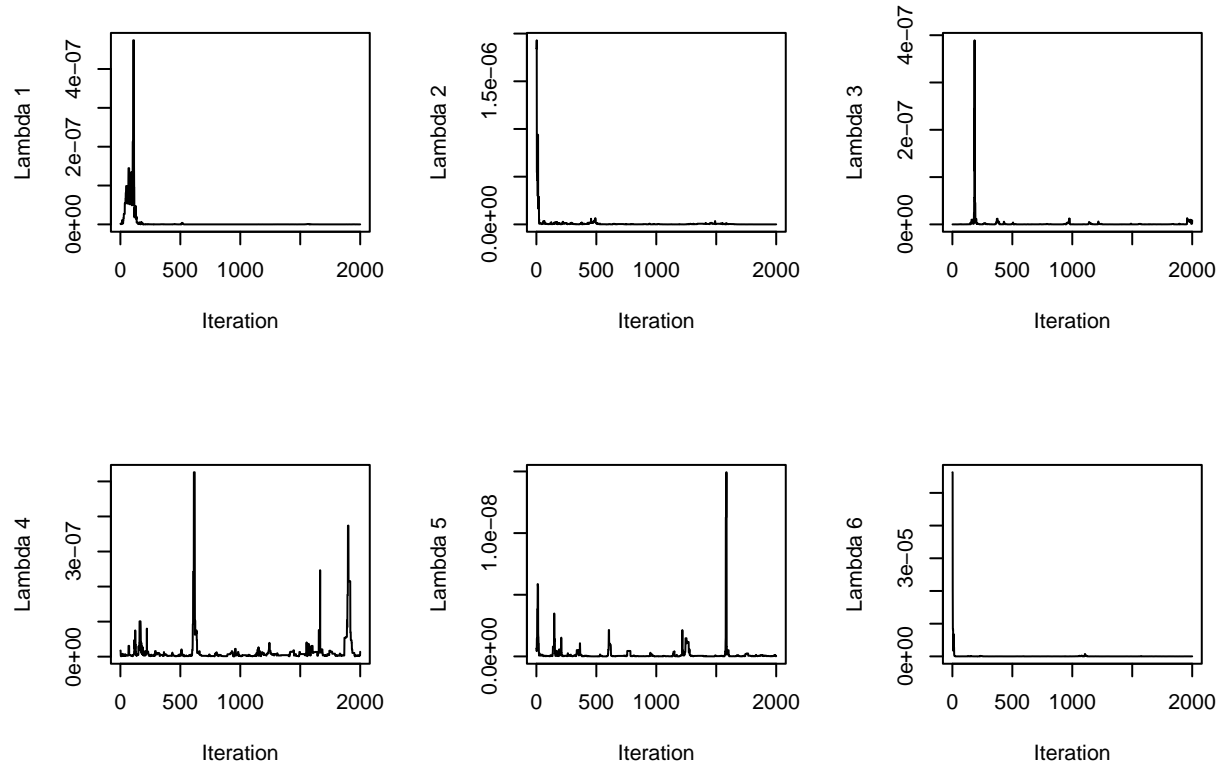
system.time({
result_horseshoe = MH_horseshoe_Sampling(Y,delta,tau,
                                         A,beta0,sigma0,var.prop,
                                         m,B,eta,
                                         Wmat_option)
})
```

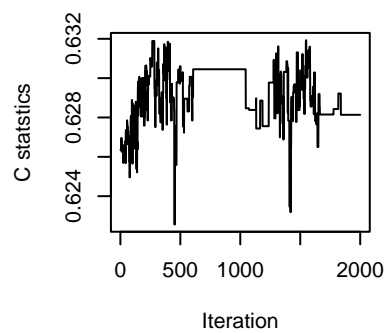
```
## user system elapsed
## 52.56 14.37 112.23
```

```
lambda_iter <- function(result,m,B){
  par(mfrow=c(2,3))
  plot(1:(m-B),result$LAMBDA[,1],xlab = "Iteration",
       ylab = "Lambda 1",type = "l")
  plot(1:(m-B),result$LAMBDA[,2],xlab = "Iteration",
       ylab = "Lambda 2",type = "l")
  plot(1:(m-B),result$LAMBDA[,3],xlab = "Iteration",
       ylab = "Lambda 3",type = "l")
  plot(1:(m-B),result$LAMBDA[,4],xlab = "Iteration",
       ylab = "Lambda 4",type = "l")
  plot(1:(m-B),result$LAMBDA[,5],xlab = "Iteration",
       ylab = "Lambda 5",type = "l")
  plot(1:(m-B),result$LAMBDA[,6],xlab = "Iteration",
       ylab = "Lambda 6",type = "l")
}
```









```
# Acceptance Rate
```

```
result_horseshoe$accept_beta
```

```
## [1] 0.1313636
```

```
result_horseshoe$accept_lambda
```

```
## [1] 0.2222727
```