

682hw5

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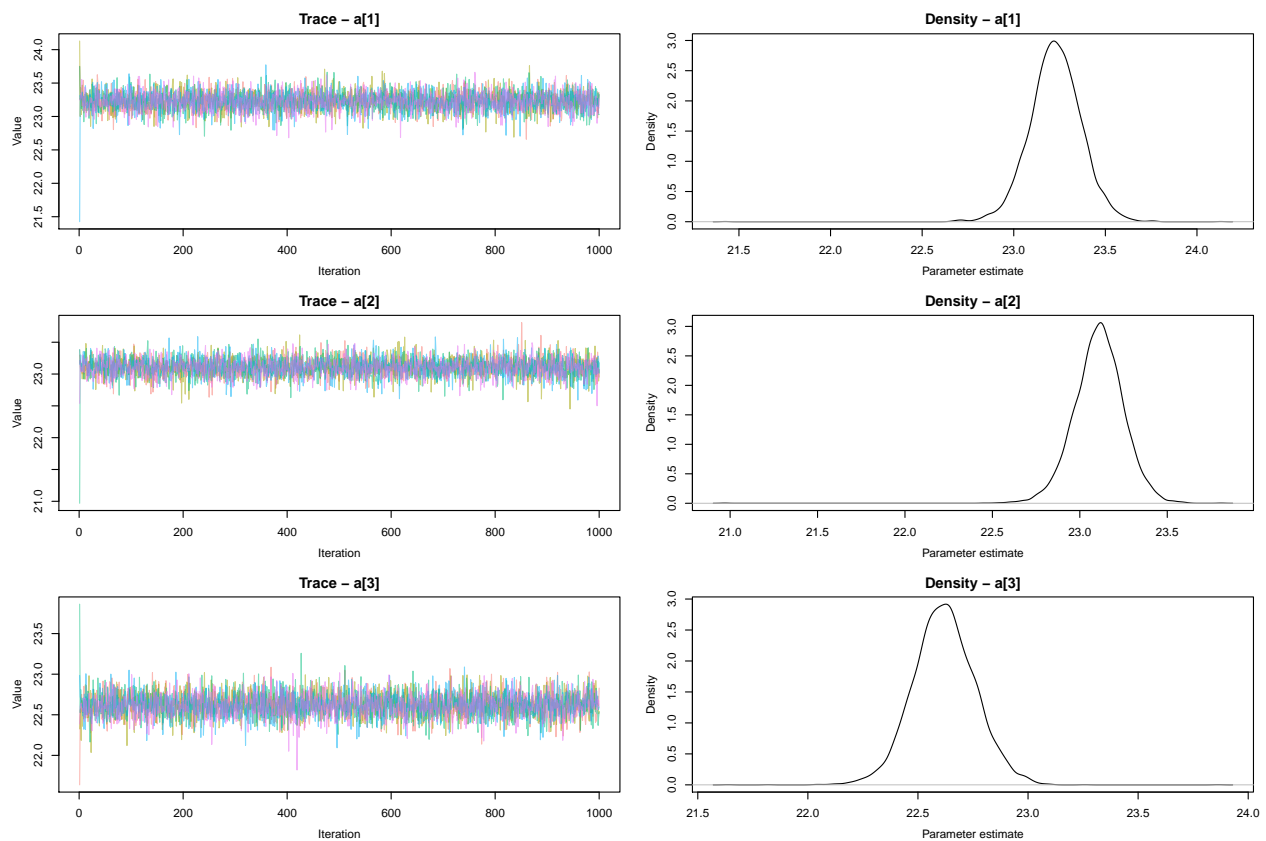
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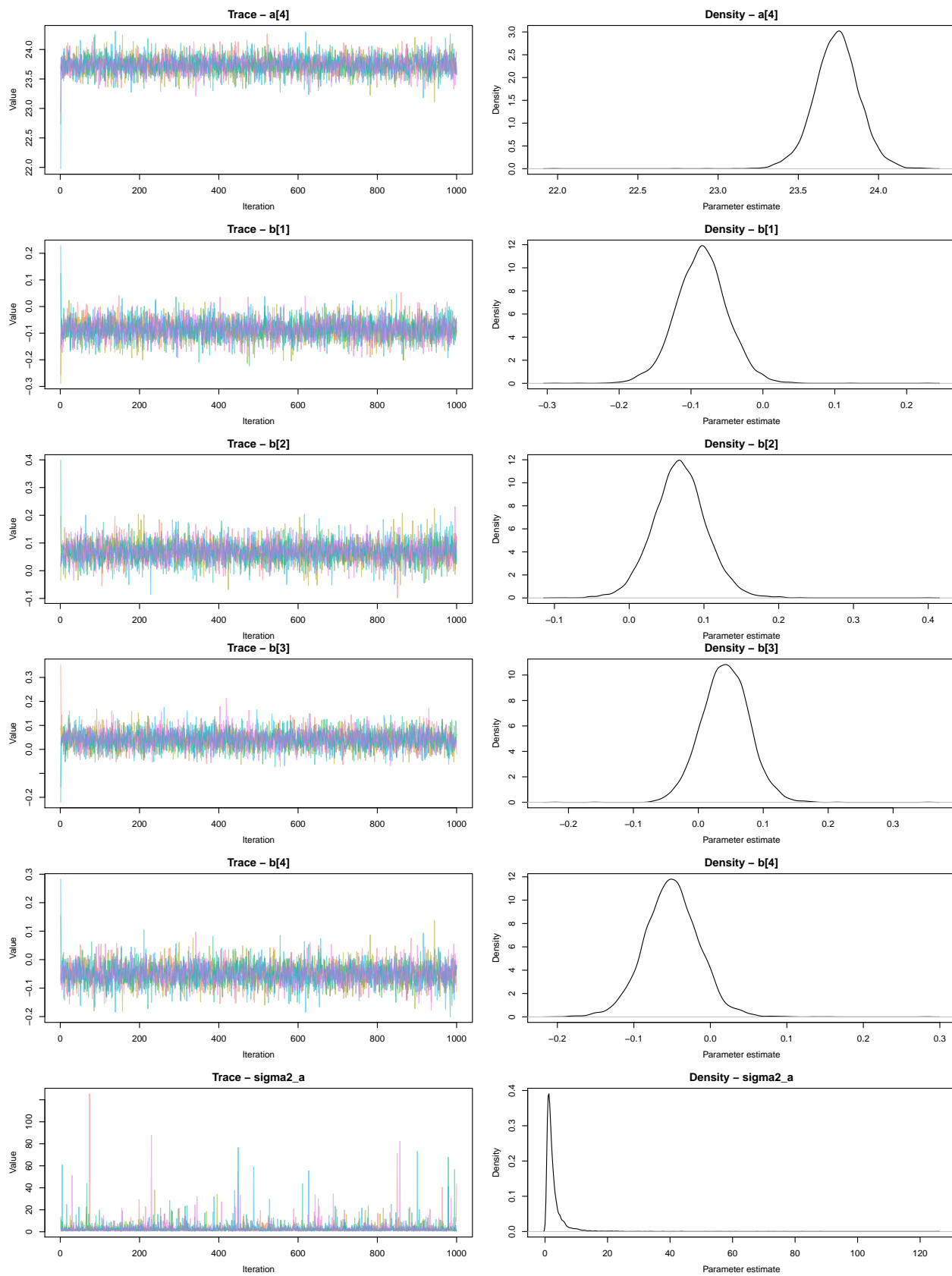
```
knitr::opts_chunk$set(fig.width=12, fig.height=8, warning=FALSE, message=FALSE)
library(R2jags)
library(MASS)
library(ggplot2)
library(mcmcplots)
library(MCMCvis)
library(geoR)

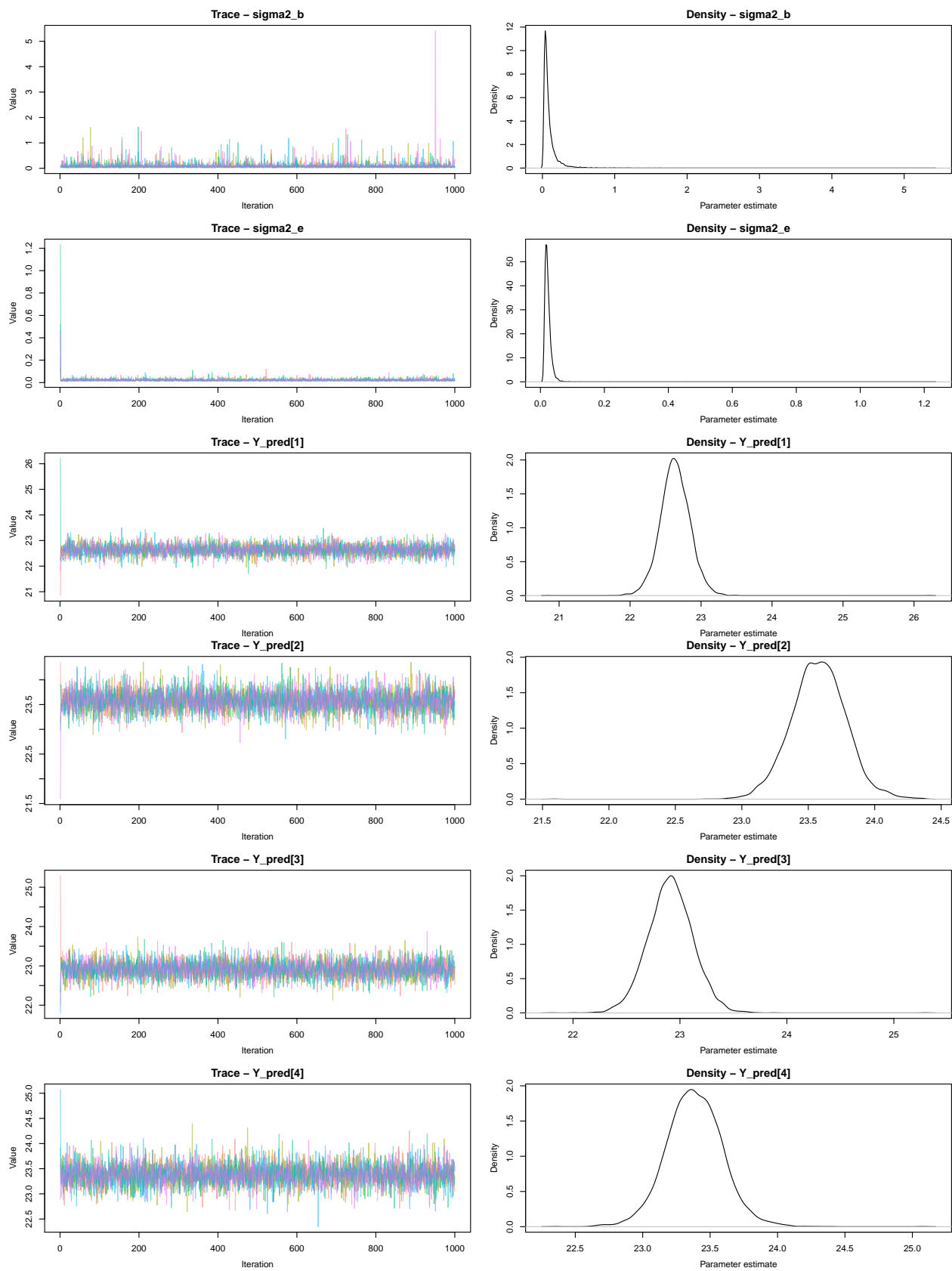
load("swim_time.RData")
ns <- nrow(Y)
nt <- ncol(Y)
JAGS_swimmer_model = function(){
  for (i in 1:ns) {
    for (j in 1:nt) {
      Y[i, j] ~ dnorm(mean[i, j], tau_e)
      mean[i, j] <- a[i] + b[i] * j
    }
  }
  for (i in 1:ns){
    a[i]~ dnorm(22, tau_a)
    b[i]~ dnorm(0, tau_b)
    Y_pred[i] ~ dnorm(a[i] + b[i] * 7, tau_e)
    z[i] <- (Y_pred[i] == min(Y_pred))
  }
  tau_a~ dgamma(0.1, 0.1)
  tau_b ~ dgamma(0.1, 0.1)
  tau_e ~ dgamma(0.1, 0.1)
  sigma2_a <- 1/tau_a
  sigma2_b <- 1/tau_b
  sigma2_e <- 1/tau_e
}

fit_swimmer_model = jags(
  data = list(Y = Y, ns = ns, nt = nt),
  inits = list(list(a = rep(20,4), b = rep(0,4)),
               list(a = rep(30,4), b = rep(1,4)),
               list(a = rep(25,4), b = rep(-1,4)),
               list(a = rep(10,4), b = rep(0,4)),
               list(a = rep(15,4), b = rep(0,4))),
  parameters.to.save = c("a","b","sigma2_a","sigma2_b", "sigma2_e", "Y_pred","z"),
  n.chains = 5,
  n.iter = 10000,
  n.burnin = 1000,
  model.file = JAGS_swimmer_model
)
```

```
## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 24
##   Unobserved stochastic nodes: 15
##   Total graph size: 158
##
## Initializing model
chains = as.mcmc(fit_swimmer_model)
MCMCtrace(chains, pdf = FALSE, params = c("a","b","sigma2_a","sigma2_b", "sigma2_e", "Y_pred"))
```







```
gelman.diag(chains, multivariate = FALSE)
```

```
## Potential scale reduction factors:
##
##           Point est. Upper C.I.
## Y_pred[1]      1.00      1.00
## Y_pred[2]      1.00      1.00
## Y_pred[3]      1.00      1.01
## Y_pred[4]      1.00      1.01
## a[1]           1.00      1.00
## a[2]           1.00      1.00
## a[3]           1.00      1.00
## a[4]           1.00      1.00
## b[1]           1.00      1.00
## b[2]           1.00      1.00
## b[3]           1.00      1.00
## b[4]           1.00      1.00
## deviance       1.00      1.01
## sigma2_a       1.01      1.01
## sigma2_b       1.09      1.10
## sigma2_e       1.00      1.00
## z[1]           1.00      1.00
## z[2]           NaN      NaN
## z[3]           1.00      1.00
## z[4]           1.06      1.06
```

b. PPD

To obtain PPD, draw Y_pred in the JAGS model, densityplots shown for Y_pred is the posterior predictive distribution. We also obtain mean and 95% CI for Y_pred .

```
summary(chains)
```

```
##
## Iterations = 1:8992
## Thinning interval = 9
## Number of chains = 5
## Sample size per chain = 1000
##
## 1. Empirical mean and standard deviation for each variable,
##    plus standard error of the mean:
##
##           Mean      SD Naive SE Time-series SE
## Y_pred[1] 22.63421 0.21431 0.0030308      0.0031274
## Y_pred[2] 23.57707 0.20651 0.0029205      0.0028497
## Y_pred[3] 22.91268 0.21203 0.0029986      0.0029293
## Y_pred[4] 23.38329 0.20896 0.0029551      0.0029547
## a[1]      23.22786 0.14225 0.0020117      0.0019784
## a[2]      23.10830 0.14413 0.0020383      0.0020386
## a[3]      22.61924 0.14266 0.0020175      0.0020678
## a[4]      23.73894 0.14266 0.0020175      0.0020173
## b[1]      -0.08508 0.03608 0.0005102      0.0004979
## b[2]       0.06775 0.03614 0.0005111      0.0005176
## b[3]       0.04161 0.03659 0.0005174      0.0005210
```

```
## b[4]          -0.05055 0.03612 0.0005108      0.0005108
## deviance     -33.65562 8.04383 0.1137570      0.1135954
## sigma2_a      2.98535 5.00851 0.0708311      0.0708179
## sigma2_b      0.09813 0.14151 0.0020013      0.0020009
## sigma2_e      0.02346 0.02180 0.0003083      0.0003065
## z[1]          0.82980 0.37585 0.0053153      0.0052624
## z[2]          0.00020 0.01414 0.0002000      0.0002000
## z[3]          0.16680 0.37283 0.0052727      0.0052203
## z[4]          0.00320 0.05648 0.0007988      0.0008346
##
## 2. Quantiles for each variable:
##
##              2.5%      25%      50%      75%      97.5%
## Y_pred[1]  22.220277  22.50143  22.63117  22.76883  23.04825
## Y_pred[2]  23.172336  23.44412  23.57878  23.71040  23.98216
## Y_pred[3]  22.503286  22.77686  22.91214  23.04770  23.31816
## Y_pred[4]  22.971321  23.24831  23.38051  23.51728  23.79680
## a[1]       22.953355  23.14013  23.22810  23.31931  23.50335
## a[2]       22.822348  23.02151  23.11209  23.20047  23.38406
## a[3]       22.345367  22.52860  22.61770  22.71080  22.90192
## a[4]       23.452953  23.65010  23.74043  23.82952  24.01211
## b[1]       -0.156986  -0.10829  -0.08505  -0.06244  -0.01451
## b[2]       -0.001405   0.04518   0.06733   0.08976   0.13957
## b[3]       -0.029914   0.01772   0.04159   0.06588   0.11344
## b[4]       -0.120128  -0.07376  -0.05106  -0.02755   0.01954
## deviance   -46.608882 -39.19464 -34.48124 -28.93283 -16.36224
## sigma2_a    0.538198   1.13404   1.80302   3.09261  12.18865
## sigma2_b    0.019433   0.03938   0.06230   0.10741   0.40388
## sigma2_e    0.011224   0.01666   0.02096   0.02703   0.04622
## z[1]        0.000000   1.00000   1.00000   1.00000   1.00000
## z[2]        0.000000   0.00000   0.00000   0.00000   0.00000
## z[3]        0.000000   0.00000   0.00000   0.00000   1.00000
## z[4]        0.000000   0.00000   0.00000   0.00000   0.00000
```

c.

In the JAGS model, we define Z as indicator function that individual i has the fastest posterior predictive value. In this case, we output the posterior mean of Z , as the probability $Pr(Y_i^* = \min(Y_1^*, \dots, Y_4^*) | \mathbf{Y})$. Based on posterior mean of $z = (0.8364, 0.0012, 0.1594, 0.003)$, we would recommend swimmer 1.

Problem 2.

a.

Linear regression.

```
#import data
X = UScrime[,1:15]
Y = UScrime[,16]
n = nrow(UScrime)
p = ncol(UScrime) - 1
df = list()
```

```

df$X = X
df$Y = Y
df$n = n
df$p = p

JAGS_BLR_flat = function(){
  # Likelihood
  for(i in 1:n){
    Y[i] ~ dnorm(mu[i],inv_sigma2)
    mu[i] <- beta_0 + inprod(X[i,],beta)
  }
  # Prior for beta
  for(j in 1:p){
    beta[j] ~ dnorm(0,0.0001)
    #non-informative priors
  }
  # Prior for intercept
  beta_0 ~ dnorm(0, 0.0001)

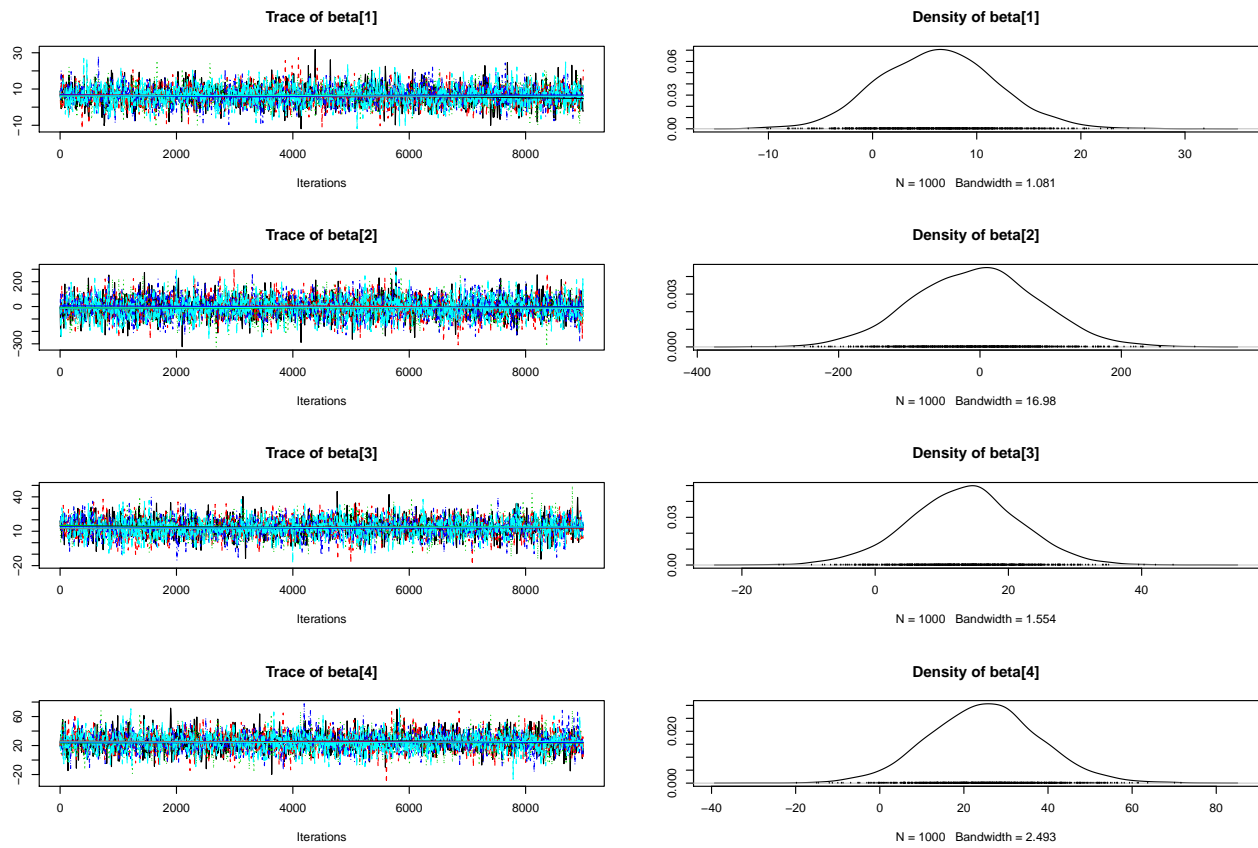
  # Prior for the inverse variance
  inv_sigma2 ~ dgamma(0.0001, 0.0001)
  sigma2 <- 1.0/inv_sigma2
}

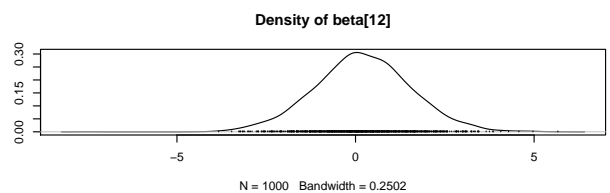
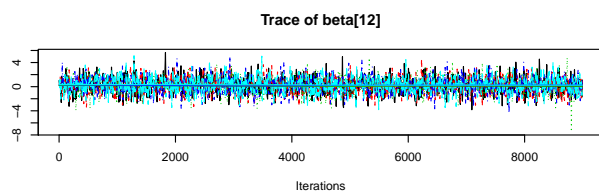
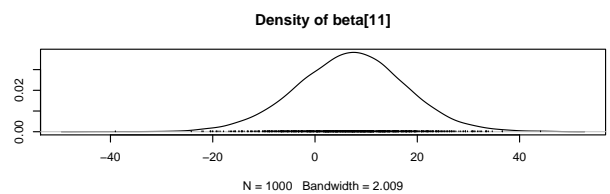
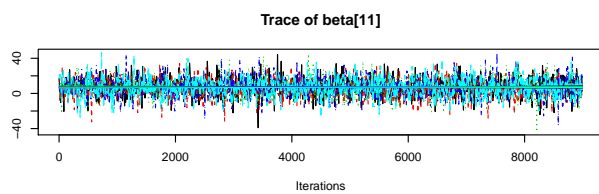
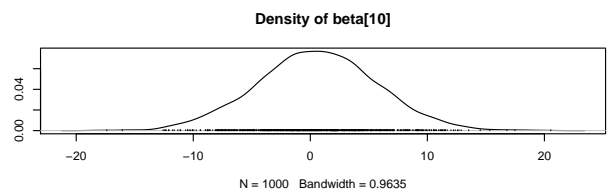
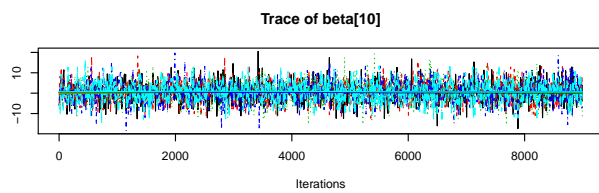
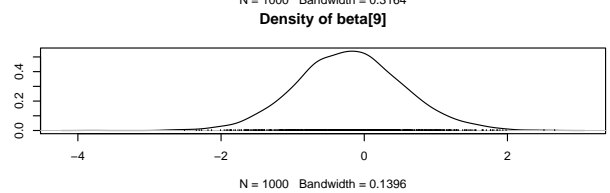
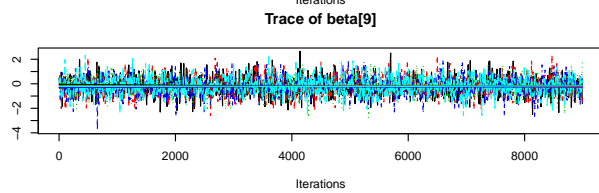
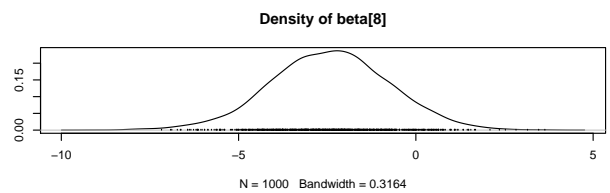
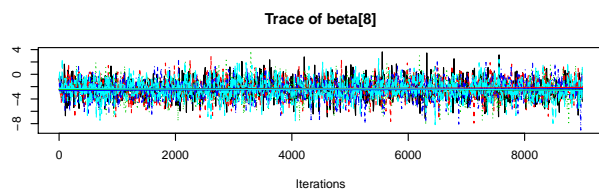
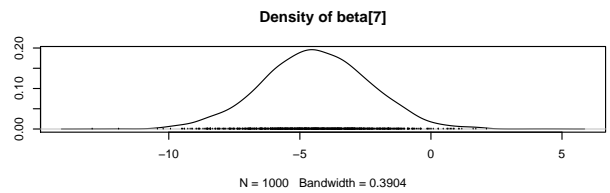
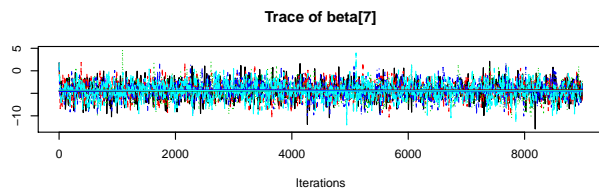
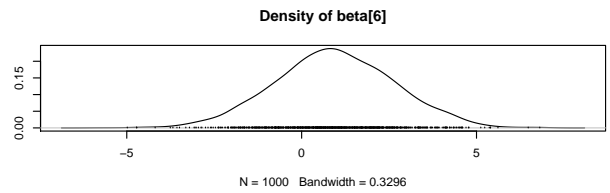
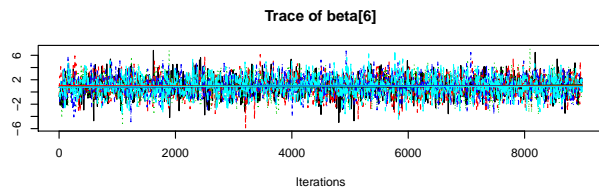
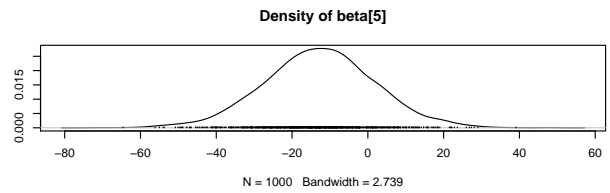
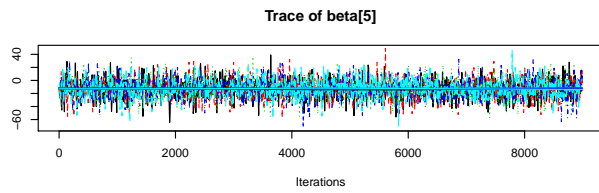
fit_JAGS_flat = jags(data=df,
  inits=list(list(beta = rnorm(p),
    beta_0 = 0,
    inv_sigma2 = 1),
    list(beta = rnorm(p),
    beta_0 = 1,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 2,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 10,
    inv_sigma2 = 5),
    list(beta = rnorm(p),
    beta_0 = 20,
    inv_sigma2 = 1)),
  parameters.to.save = c("beta_0","beta","sigma2"),
  n.chains=5,
  n.iter=10000,
  n.burnin=1000,
  model.file=JAGS_BLR_flat)

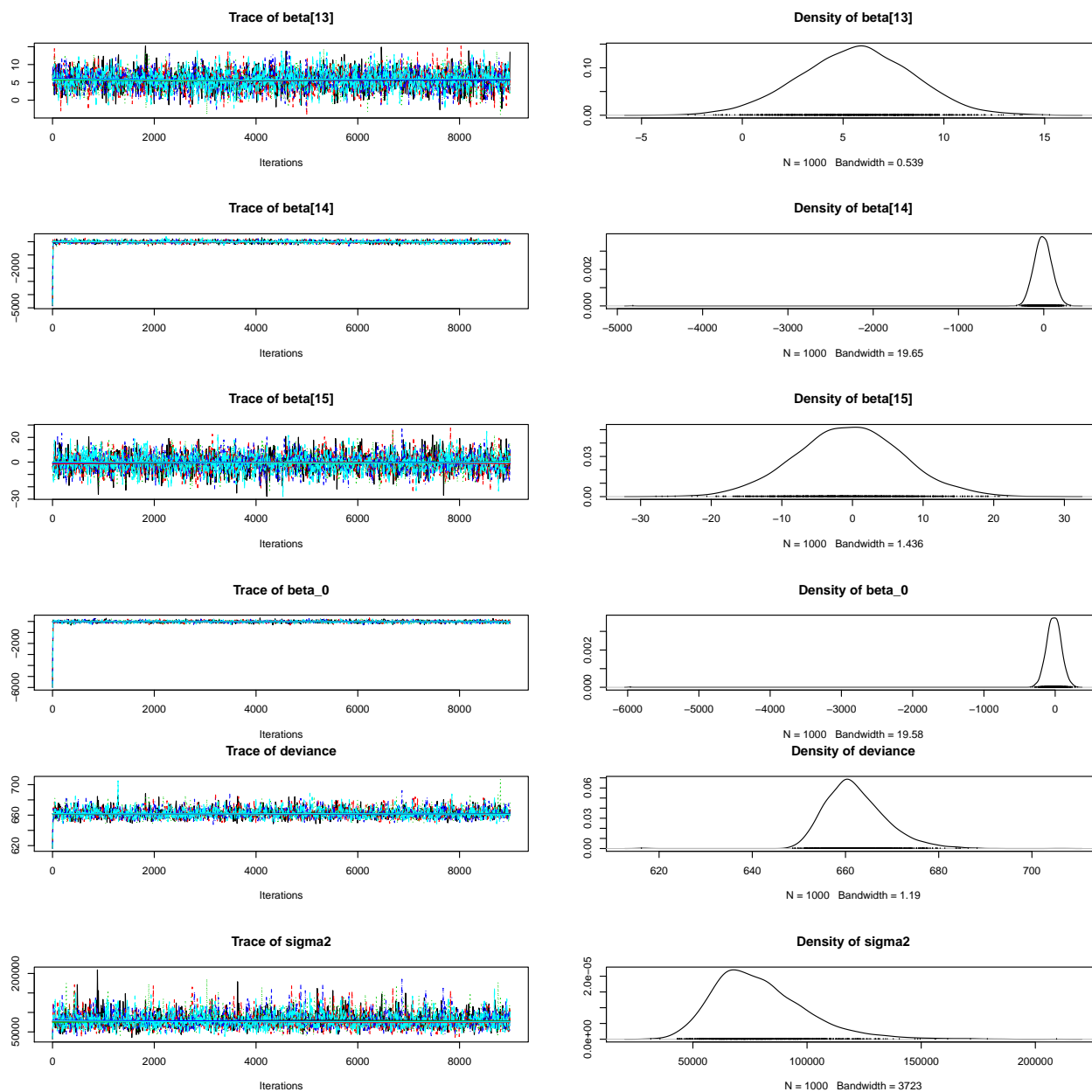
## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 47
##   Unobserved stochastic nodes: 17
##   Total graph size: 950

```

```
##
## Initializing model
chains_2a = as.mcmc(fit_JAGS_flat)
plot(chains_2a)
```







```
gelman.diag(chains_2a)
```

```
## Potential scale reduction factors:
##
##      Point est. Upper C.I.
## beta[1]      1.003      1.01
## beta[2]      0.999      1.00
## beta[3]      1.002      1.01
## beta[4]      1.001      1.00
## beta[5]      1.001      1.01
## beta[6]      1.000      1.00
## beta[7]      1.001      1.00
## beta[8]      1.001      1.00
## beta[9]      1.001      1.00
## beta[10]     1.000      1.00
```

```
## beta[11]      1.000      1.00
## beta[12]      1.001      1.00
## beta[13]      1.001      1.00
## beta[14]      1.000      1.00
## beta[15]      1.000      1.00
## beta_0        1.000      1.00
## deviance      1.001      1.00
## sigma2        1.005      1.01
##
## Multivariate psrf
##
## 1.01
```

```
##output 95% credible interval
summary(chains_2a)
```

```
##
## Iterations = 1:8992
## Thinning interval = 9
## Number of chains = 5
## Sample size per chain = 1000
##
## 1. Empirical mean and standard deviation for each variable,
##    plus standard error of the mean:
```

```
##
##           Mean      SD Naive SE Time-series SE
## beta[1]      6.2665    5.600  0.07919      0.07773
## beta[2]     -6.2477   87.991  1.24438      1.22961
## beta[3]     13.2462    8.368  0.11835      0.11490
## beta[4]     24.8543   13.238  0.18721      0.18716
## beta[5]    -12.7524   14.686  0.20769      0.20759
## beta[6]      0.9395    1.712  0.02421      0.02328
## beta[7]     -4.3958    2.046  0.02894      0.02829
## beta[8]     -2.4215    1.678  0.02373      0.02372
## beta[9]     -0.1999    0.745  0.01054      0.01004
## beta[10]      0.5547    5.136  0.07264      0.07169
## beta[11]      7.0196   10.712  0.15148      0.15147
## beta[12]      0.2036    1.353  0.01913      0.01952
## beta[13]      5.6415    2.805  0.03966      0.03790
## beta[14]    -13.6725  182.689  2.58361      2.58445
## beta[15]     -0.4990    7.564  0.10696      0.10700
## beta_0     -23.4653   213.148  3.01437      3.01544
## deviance    662.2543    6.606  0.09343      0.09352
## sigma2   79533.4483 20590.852 291.19863     290.99404
```

```
##
## 2. Quantiles for each variable:
```

	2.5%	25%	50%	75%	97.5%
## beta[1]	-4.327e+00	2.391e+00	6.2105	9.9408	1.747e+01
## beta[2]	-1.797e+02	-6.666e+01	-4.6218	51.5321	1.662e+02
## beta[3]	-3.784e+00	7.825e+00	13.4183	18.6130	2.945e+01
## beta[4]	-9.218e-01	1.599e+01	24.8460	33.3070	5.166e+01
## beta[5]	-4.161e+01	-2.221e+01	-12.7385	-3.1869	1.700e+01
## beta[6]	-2.428e+00	-1.824e-01	0.9265	2.1062	4.268e+00
## beta[7]	-8.468e+00	-5.753e+00	-4.4165	-3.0418	-4.739e-01

```
## beta[8] -5.781e+00 -3.527e+00 -2.4091 -1.3297 8.129e-01
## beta[9] -1.640e+00 -6.926e-01 -0.2052 0.2767 1.312e+00
## beta[10] -9.464e+00 -2.802e+00 0.5451 3.8889 1.066e+01
## beta[11] -1.439e+01 -5.498e-02 7.2358 13.8972 2.812e+01
## beta[12] -2.517e+00 -6.659e-01 0.1811 1.0713 2.925e+00
## beta[13] 9.149e-02 3.803e+00 5.7142 7.5454 1.104e+01
## beta[14] -2.036e+02 -7.807e+01 -10.3631 58.3941 1.889e+02
## beta[15] -1.551e+01 -5.458e+00 -0.4843 4.5134 1.476e+01
## beta_0 -2.134e+02 -8.536e+01 -16.9771 50.5969 1.781e+02
## deviance 6.519e+02 6.577e+02 661.4326 666.0076 6.771e+02
## sigma2 4.852e+04 6.488e+04 76334.0844 90729.4792 1.284e+05
```

b.

Cross validation

```
#split data into training and test set
split_data = function(df,train_test_ratio = 1,random=TRUE){
  n_train = floor(df$n*train_test_ratio/(1+train_test_ratio))
  n_test = df$n - n_train
  if(random){
    train_idx = sample(1:n,n_train,replace = FALSE)
    test_idx = setdiff(1:n,train_idx)
  }
  else{
    train_idx = 1:n_train
    test_idx = n_train+1:n_test
  }

  df_t = list()
  df_t$Y_train = df$Y[train_idx]
  df_t$X_train = df$X[train_idx,,drop=FALSE]
  df_t$X_test = df$X[test_idx,,drop=FALSE]
  df_t$n_train = n_train
  df_t$n_test = n_test
  df_t$p = df$p

  return(list(df_t=df_t,Y_test=df$Y[test_idx]))
}

pred = split_data(df, random = FALSE)

##define a predictive JAGS
JAGS_BLR_flat_pred = function(){
  # Likelihood
  for(i in 1:n_train){
    Y_train[i] ~ dnorm(mu_train[i],inv_sigma2)
    mu_train[i] <- beta_0 + inprod(X_train[i,],beta)
    # same as beta_0 + X[i,1]*beta[1] + ... + X[i,p]*beta[p]
  }
  # Prior for beta
  for(j in 1:p){
    beta[j] ~ dnorm(0,0.0001)
    #non-informative priors
  }
}
```

```

}
# Prior for intercept
beta_0 ~ dnorm(0, 0.0001)

# Prior for the inverse variance
inv_sigma2 ~ dgamma(0.0001, 0.0001)
sigma2 <- 1.0/inv_sigma2

#prediction
# Predictions
for(i in 1:n_test){
  Y_test[i] ~ dnorm(mu_test[i],inv_sigma2)
  mu_test[i] <- beta_0 + inprod(X_test[i,],beta)
}
}

fit_JAGS_flat_pred = jags(data=pred$df_t,
  inits=list(list(beta = rnorm(p),
    beta_0 = 0,
    inv_sigma2 = 1),
    list(beta = rnorm(p),
    beta_0 = 1,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 2,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 10,
    inv_sigma2 = 5),
    list(beta = rnorm(p),
    beta_0 = 20,
    inv_sigma2 = 1)),
  parameters.to.save = c("beta_0","beta","sigma2", "Y_test"),
  n.chains=5,
  n.iter=10000,
  n.burnin=1000,
  model.file=JAGS_BLR_flat_pred)

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 23
##   Unobserved stochastic nodes: 41
##   Total graph size: 951
##
## Initializing model

chains_2b = as.mcmc(fit_JAGS_flat_pred)

##plot the predictive values
result = summary(chains_2b)
q = result$quantiles
dtfr = as.data.frame(cbind(result$quantiles))

```

```
##compare
pred$Y_test

## [1] 968 523 1993 342 1216 1043 696 373 754 1072 923 653 1272 831
## [15] 566 826 1151 880 542 823 1030 455 508 849

fit_JAGS_flat_pred$BUGSoutput$median$Y_test

## [1] 702.3189 911.1052 1694.7808 190.2616 861.1977 2098.1610 798.2761
## [8] 289.2185 1290.1620 658.4909 761.7653 1272.8585 764.8326 562.4828
## [15] 745.2396 995.5814 1403.8654 534.4566 503.4575 887.8674 580.6554
## [22] 550.1389 1331.0586 592.6950
```

c. Slab and spike

```
JAGS_BLR_SpikeSlab = function(){
  # Likelihood
  for(i in 1:n){
    Y[i] ~ dnorm(mu[i],inv_sigma2)
    mu[i] <- beta_0 + inprod(X[i,],beta)
    # same as beta_0 + X[i,1]*beta[1] + ... + X[i,p]*beta[p]
  }
  #Prior for beta_j
  for(j in 1:p){
    beta[j] ~ dnorm(0,inv_tau2[j])
    inv_tau2[j] <- (1-gamma[j])*1000+gamma[j]*0.01
    gamma[j] ~ dbern(0.5)
  }
  # Prior for intercept
  beta_0 ~ dnorm(0, 0.0001)

  # Prior for the inverse variance
  inv_sigma2 ~ dgamma(0.0001, 0.0001)
  sigma2 <- 1.0/inv_sigma2
}

fit_JAGS_SpikeSlab = jags(data=df,
  inits=list(list(beta = rnorm(p),
    beta_0 = 0,
    inv_sigma2 = 1),
    list(beta = rnorm(p),
    beta_0 = 1,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 2,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 10,
    inv_sigma2 = 5),
    list(beta = rnorm(p),
    beta_0 = 20,
    inv_sigma2 = 1)),
  parameters.to.save = c("beta_0","beta","sigma2"),
```

```

n.chains=5,
n.iter=10000,
n.burnin=1000,
model.file=JAGS_BLR_flat)

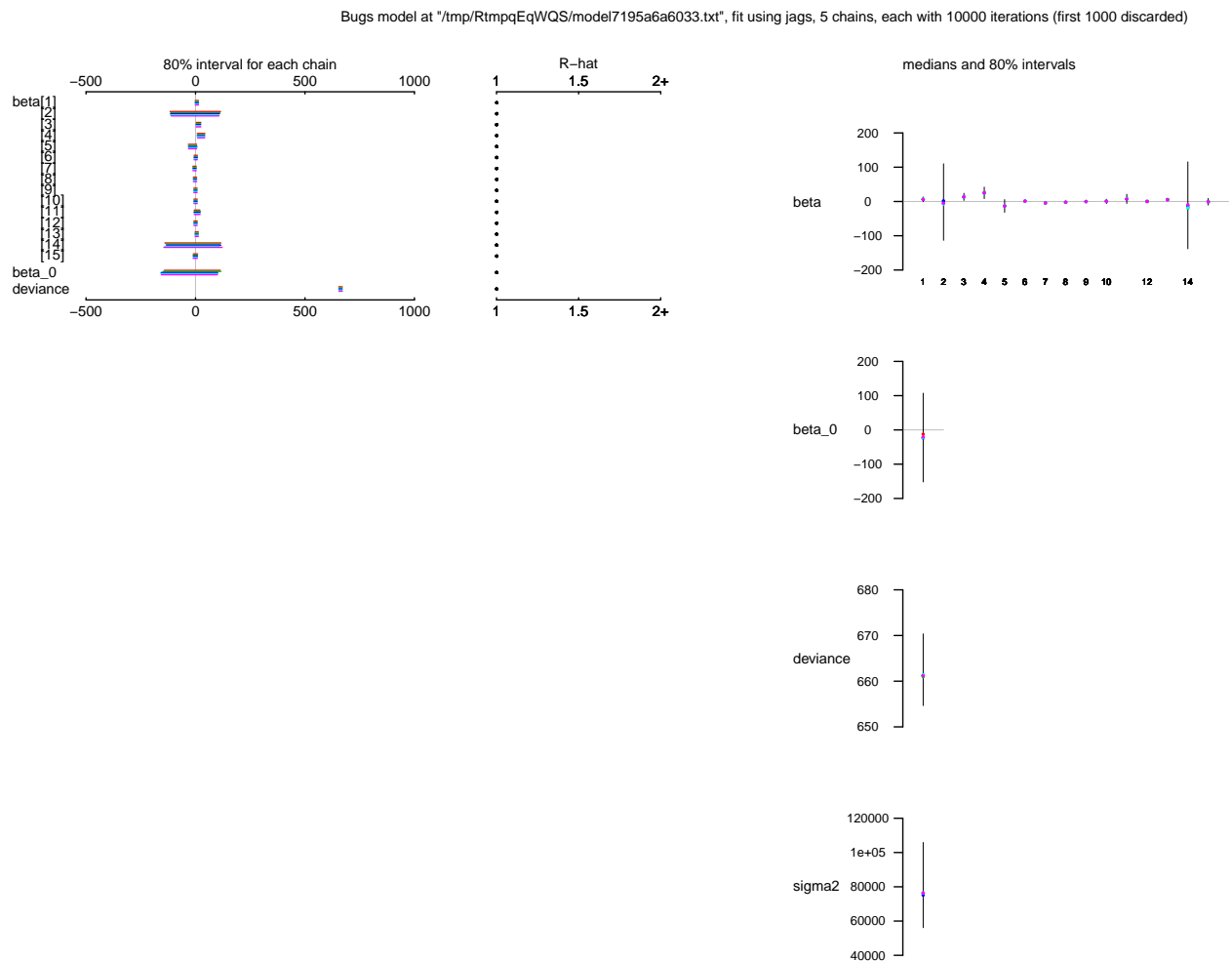
```

```

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 47
##   Unobserved stochastic nodes: 17
##   Total graph size: 950
##
## Initializing model

```

```
plot(fit_JAGS_SpikeSlab)
```



```

chains_2c = as.mcmc(fit_JAGS_SpikeSlab)
summary(chains_2c)

```

```

##
## Iterations = 1:8992
## Thinning interval = 9
## Number of chains = 5

```

```

## Sample size per chain = 1000
##
## 1. Empirical mean and standard deviation for each variable,
##    plus standard error of the mean:
##
##          Mean          SD Naive SE Time-series SE
## beta[1]      6.3092      5.400  0.07637      0.07628
## beta[2]     -2.6117     88.419  1.25043      1.25034
## beta[3]     13.2802      8.259  0.11680      0.11681
## beta[4]     25.1446     13.178  0.18636      0.18134
## beta[5]    -13.0849     14.577  0.20615      0.20049
## beta[6]      0.9506      1.692  0.02392      0.02419
## beta[7]     -4.3264      1.991  0.02815      0.02816
## beta[8]     -2.3521      1.654  0.02339      0.02338
## beta[9]     -0.2120      0.739  0.01045      0.01053
## beta[10]     0.3355      5.068  0.07167      0.07170
## beta[11]     7.4999     10.496  0.14843      0.14843
## beta[12]     0.1314      1.325  0.01874      0.01846
## beta[13]     5.4855      2.782  0.03934      0.03935
## beta[14]    -17.3998    182.078  2.57497      2.57593
## beta[15]     -0.7892      7.525  0.10642      0.10578
## beta_0     -27.1134    213.578  3.02045      3.02106
## deviance    662.0127      6.519  0.09219      0.09116
## sigma2    79143.6611 20658.342 292.15307     306.47839
##
## 2. Quantiles for each variable:
##
##          2.5%          25%          50%          75%          97.5%
## beta[1] -4.554e+00      2.7418      6.3351      9.9140     1.670e+01
## beta[2] -1.759e+02    -61.6151     -4.4024     55.8237     1.725e+02
## beta[3] -3.055e+00      7.8237     13.3335     18.6867     2.982e+01
## beta[4]  1.121e-01     16.3637     25.0514     33.8903     5.097e+01
## beta[5] -4.198e+01    -22.7576    -13.1229     -3.4576     1.527e+01
## beta[6] -2.346e+00     -0.1727      0.9546      2.0783     4.264e+00
## beta[7] -8.164e+00     -5.6202     -4.3302     -3.0469    -4.070e-01
## beta[8] -5.645e+00     -3.4453     -2.3590     -1.2744     9.474e-01
## beta[9] -1.653e+00     -0.6958     -0.1988      0.2833     1.220e+00
## beta[10] -9.754e+00    -3.0409      0.4276      3.7375     1.029e+01
## beta[11] -1.295e+01      0.6334      7.4273     14.4106     2.804e+01
## beta[12] -2.517e+00     -0.7261      0.1227      1.0188     2.759e+00
## beta[13]  9.279e-02      3.5854      5.4443      7.3662     1.102e+01
## beta[14] -2.079e+02    -80.9641    -14.8048     53.8219     1.900e+02
## beta[15] -1.534e+01     -5.6004     -0.9122      4.1580     1.398e+01
## beta_0  -2.216e+02    -89.2852    -19.9943     46.1587     1.732e+02
## deviance  6.518e+02    657.5149    661.2607    665.6451     6.771e+02
## sigma2   4.816e+04  64411.3123  76117.5539  89902.4117     1.282e+05
#####
##Prediction
#####

JAGS_BLR_SpikeSlab_pred = function(){
  # Likelihood

```



```

for(i in 1:n_train){
  Y_train[i] ~ dnorm(mu_train[i],inv_sigma2)
  mu_train[i] <- beta_0 + inprod(X_train[i,],beta)
  # same as beta_0 + X[i,1]*beta[1] + ... + X[i,p]*beta[p]
}
# Prior for beta
for(j in 1:p){
  beta[j] ~ dnorm(0,inv_tau2[j])
  inv_tau2[j] <- (1-gamma[j])*1000+gamma[j]*0.01
  gamma[j] ~ dbern(0.5)
}
# Prior for intercept
beta_0 ~ dnorm(0, 0.0001)

# Prior for the inverse variance
inv_sigma2 ~ dgamma(0.0001, 0.0001)
sigma2 <- 1.0/inv_sigma2

#prediction
# Predictions
for(i in 1:n_test){
  Y_test[i] ~ dnorm(mu_test[i],inv_sigma2)
  mu_test[i] <- beta_0 + inprod(X_test[i,],beta)
}
}

fit_JAGS_SpikeSlab_pred = jags(data = pred$df_t,
  inits=list(list(beta = rnorm(p),
    beta_0 = 0,
    inv_sigma2 = 1),
    list(beta = rnorm(p),
    beta_0 = 1,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 2,
    inv_sigma2 = 2),
    list(beta = rnorm(p),
    beta_0 = 10,
    inv_sigma2 = 5),
    list(beta = rnorm(p),
    beta_0 = 20,
    inv_sigma2 = 1)),
  parameters.to.save = c("beta_0","beta","sigma2","Y_test"),
  n.chains=5,
  n.iter=10000,
  n.burnin=1000,
  model.file=JAGS_BLR_SpikeSlab_pred)

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 23
##   Unobserved stochastic nodes: 56

```

```
## Total graph size: 1071
```

```
##
```

```
## Initializing model
```

```
cbind(pred$Y_test, fit_JAGS_SpikeSlab_pred$BUGSoutput$median$Y_test)
```

```
##      [,1]      [,2]
```

```
## [1,]  968  796.8714
```

```
## [2,]  523  674.8399
```

```
## [3,] 1993 1440.6227
```

```
## [4,]  342  680.0654
```

```
## [5,] 1216  835.9979
```

```
## [6,] 1043 1936.1407
```

```
## [7,]  696  746.9942
```

```
## [8,]  373  611.9618
```

```
## [9,]  754 1119.7948
```

```
## [10,] 1072  740.4464
```

```
## [11,]  923  979.5285
```

```
## [12,]  653 1220.9262
```

```
## [13,] 1272 1035.2173
```

```
## [14,]  831  722.1332
```

```
## [15,]  566  591.3444
```

```
## [16,]  826  773.2865
```

```
## [17,] 1151 1080.8901
```

```
## [18,]  880  692.0879
```

```
## [19,]  542  579.3675
```

```
## [20,]  823  867.2079
```

```
## [21,] 1030  985.5025
```

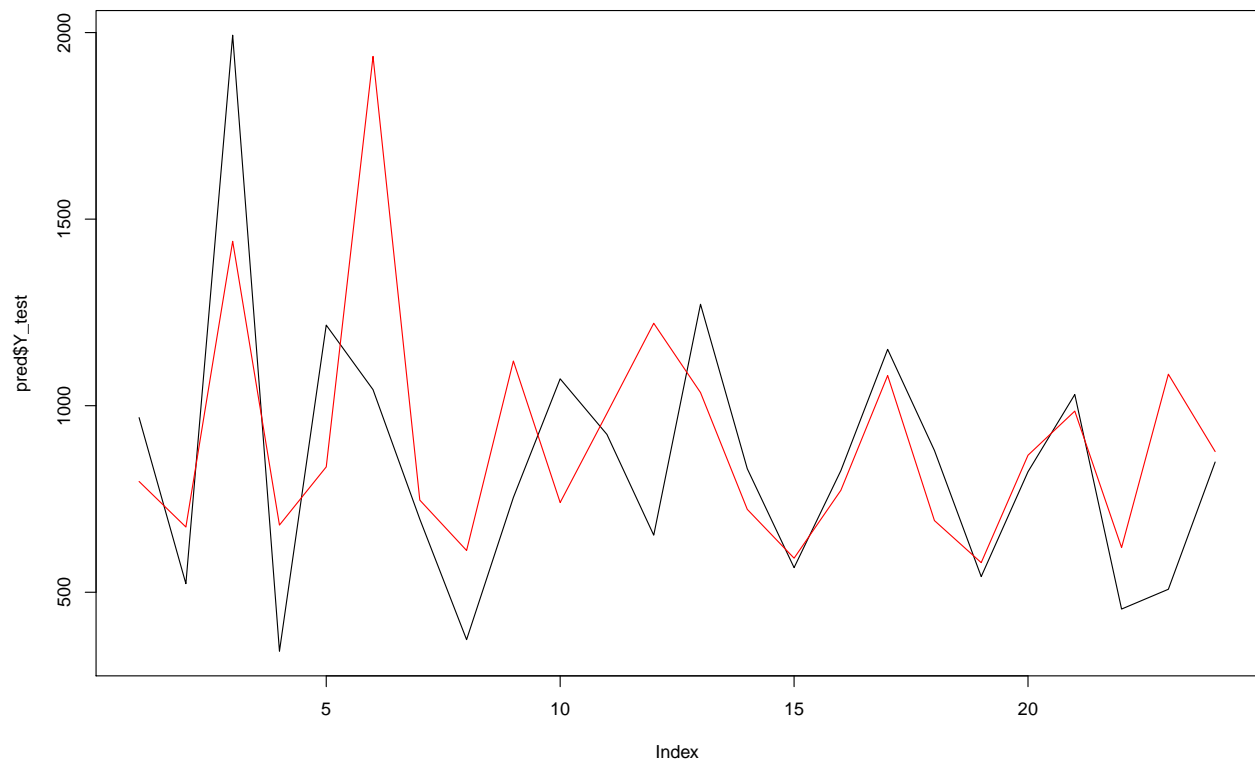
```
## [22,]  455  619.7641
```

```
## [23,]  508 1084.4153
```

```
## [24,]  849  877.1035
```

```
plot(pred$Y_test, type = 'l')
```

```
lines(fit_JAGS_SpikeSlab_pred$BUGSoutput$median$Y_test, col= 'red')
```



```
v_loc = unique(gambia[, "x"])
v = match(gambia[, "x"], v_loc)
```