

Bayesian Statistics week 4

```
library("COUNT")

## Loading required package: msme
## Loading required package: MASS
## Loading required package: lattice
## Loading required package: sandwich

data("badhealth")
?badhealth
head(badhealth)

##   numvisit badh age
## 1      30    0  58
## 2      20    0  54
## 3      16    0  44
## 4      20    0  57
## 5      15    0  33
## 6      15    0  28

library("rjags")

## Warning: package 'rjags' was built under R version 4.0.2
## Loading required package: coda
## Linked to JAGS 4.3.0
## Loaded modules: basemod,bugs

mod_string = " model {
  for (i in 1:length(numvisit)) {
    numvisit[i] ~ dpois(lam[i])
    log(lam[i]) = int + b_badh*badh[i] + b_age*age[i]
  }

  int ~ dnorm(0.0, 1.0/1e6)
  b_badh ~ dnorm(0.0, 1.0/1e4)
  b_age ~ dnorm(0.0, 1.0/1e4)
} "

set.seed(102)

data_jags = as.list(badhealth)

params = c("int", "b_badh", "b_age")
```

```

mod = jags.model(textConnection(mod_string), data=data_jags, n.chains=3)

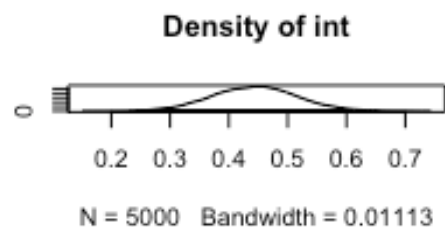
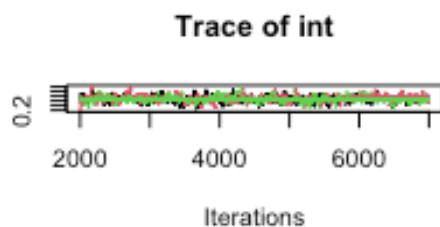
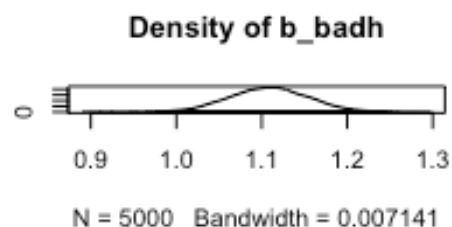
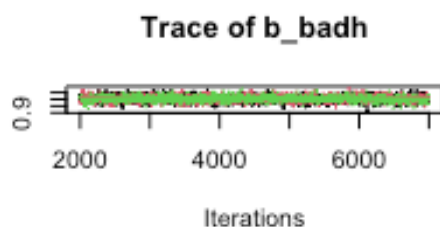
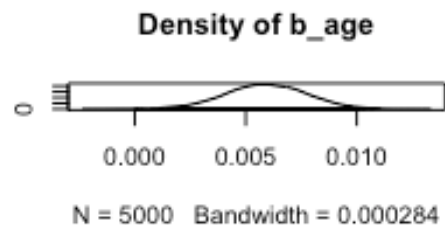
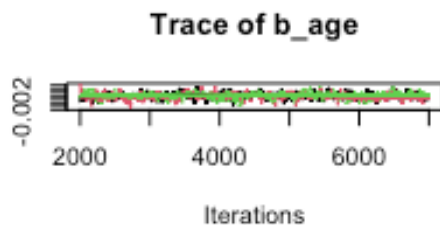
## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 1127
##   Unobserved stochastic nodes: 3
##   Total graph size: 3587
##
## Initializing model

update(mod, 1e3)

mod_sim = coda.samples(model=mod,
                       variable.names=params,
                       n.iter=5e3)
mod_csim = as.mcmc(do.call(rbind, mod_sim))

## convergence diagnostics
plot(mod_sim)

```



```
gelman.diag(mod_sim)
```

```
## Potential scale reduction factors:
```

```
##
```

```
##          Point est. Upper C.I.
```

```
## b_age      1.01      1.02
```

```
## b_badh     1.00      1.01
```

```
## int        1.01      1.02
```

```
##
```

```
## Multivariate psrf
```

```
##
```

```
## 1.01
```

```
autocorr.diag(mod_sim)
```

```
##          b_age      b_badh      int
```

```
## Lag 0  1.00000000 1.00000000 1.00000000
```

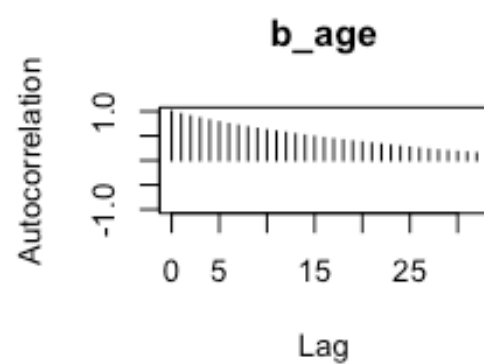
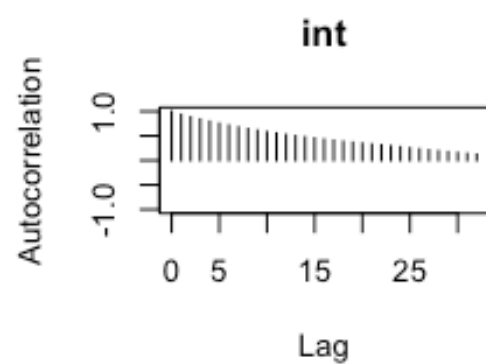
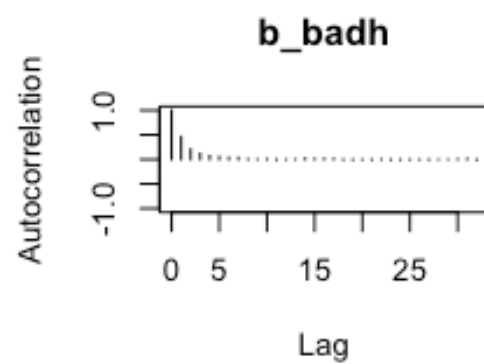
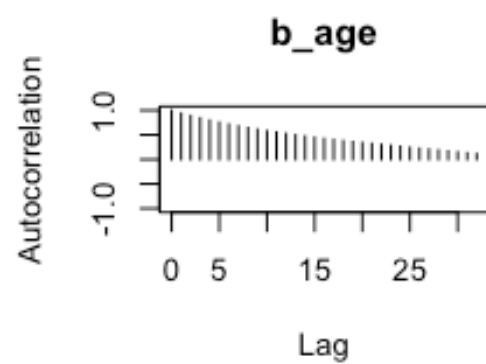
```
## Lag 1  0.94642378 0.474283632 0.94194252
```

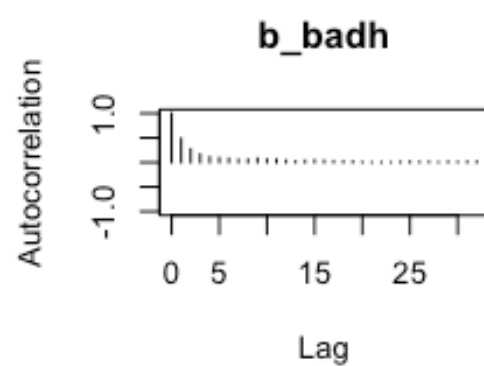
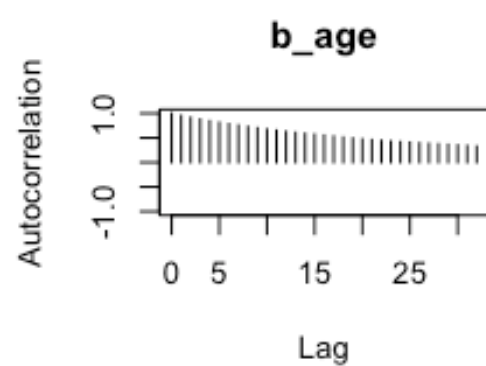
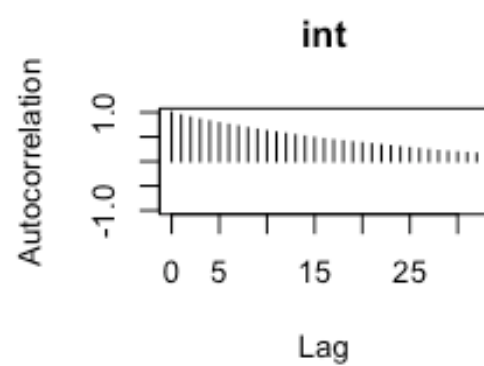
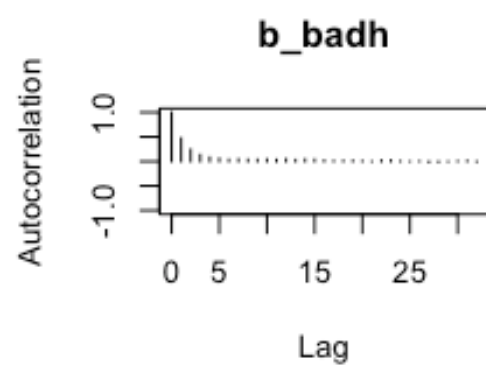
```
## Lag 5  0.78216449 0.067730795 0.77663953
```

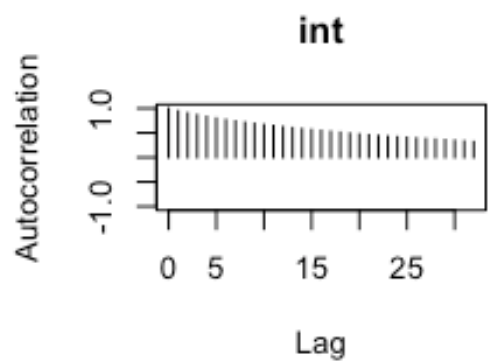
```
## Lag 10 0.62445905 0.037836302 0.62202174
```

```
## Lag 50 0.07126978 0.004483109 0.06460705
```

```
autocorr.plot(mod_sim)
```







```
effectiveSize(mod_sim)

##      b_age      b_badh      int
## 357.4101 4589.2172 369.2683

## compute DIC
dic = dic.samples(mod, n.iter=1e3)

X = as.matrix(badhealth[, -1])
head(X)

##      badh age
## [1,]    0  58
## [2,]    0  54
## [3,]    0  44
## [4,]    0  57
## [5,]    0  33
## [6,]    0  28

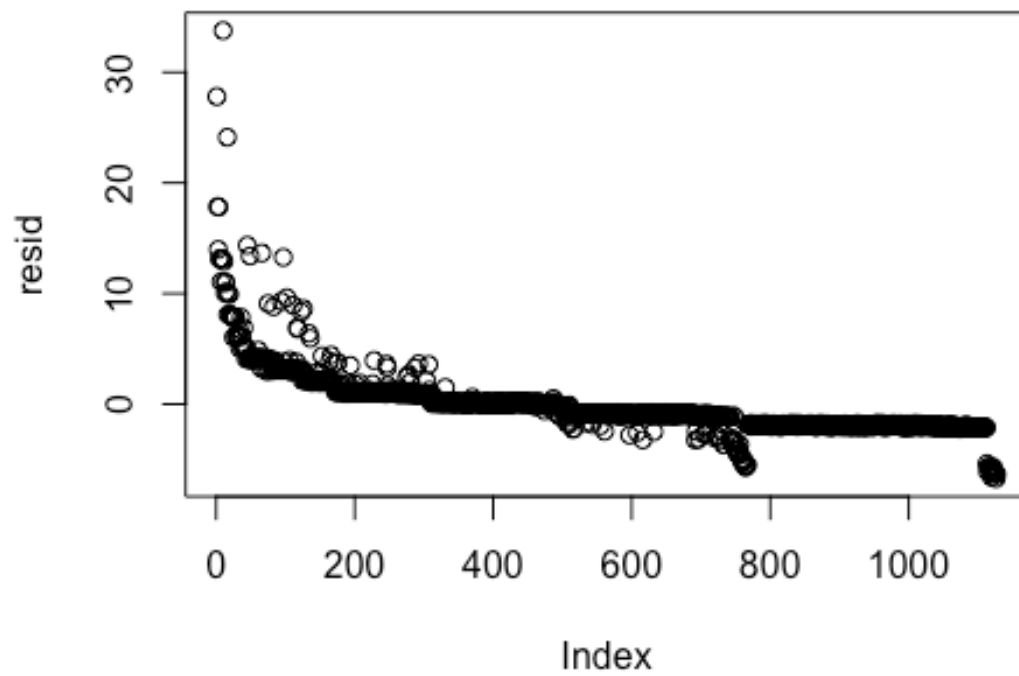
(pmed_coef = apply(mod_csim, 2, median))

##      b_age      b_badh      int
## 0.005900392 1.108531028 0.444110580
```

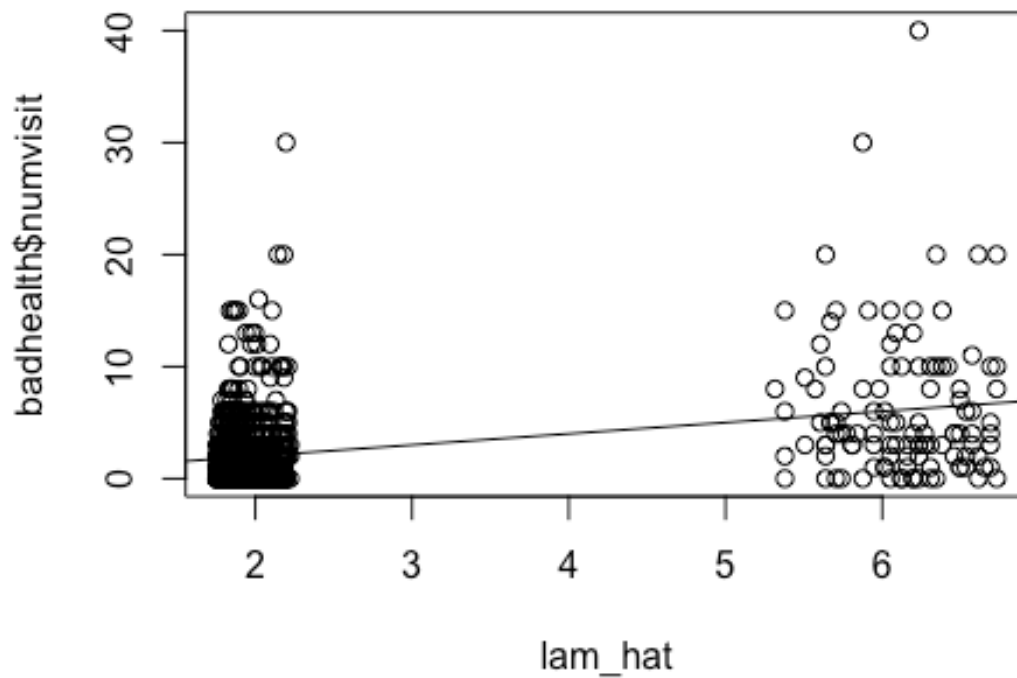
```
l1lam_hat = pmed_coef["int"] + X %*% pmed_coef[c("b_badh", "b_age")]
lam_hat = exp(l1lam_hat)
hist(lam_hat)
```



```
resid = badhealth$numvisit - lam_hat
plot(resid) # the data were ordered
```



```
plot(lam_hat, badhealth$numvisit)  
abline(0.0, 1.0)
```

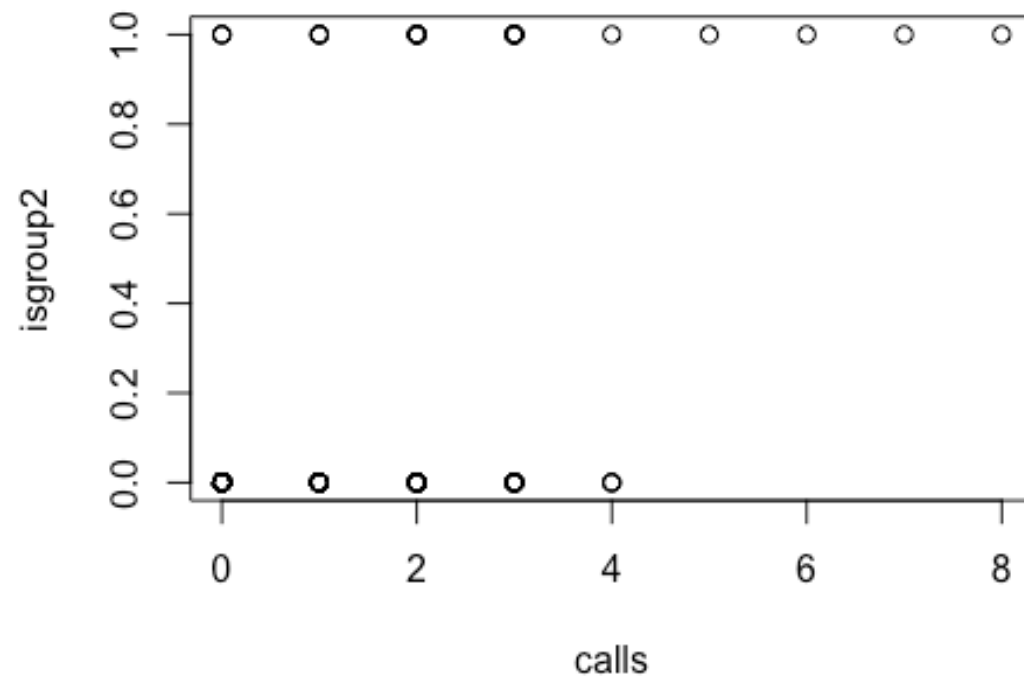
```
dat = read.csv(file="data.csv", header=TRUE)
```

```
attach(dat)
```

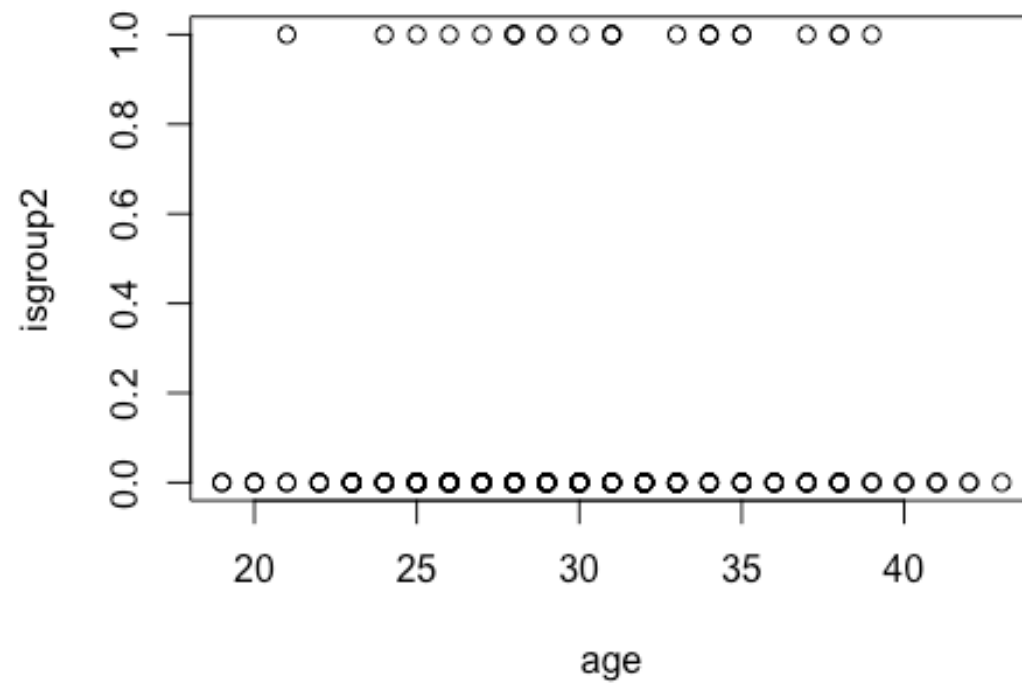
```
head(dat)
```

```
##   calls days_active isgroup2 age
## 1     2          32         0  27
## 2     4          81         0  32
## 3     0          41         0  22
## 4     1          36         0  28
## 5     0          55         0  31
## 6     0          25         0  33
```

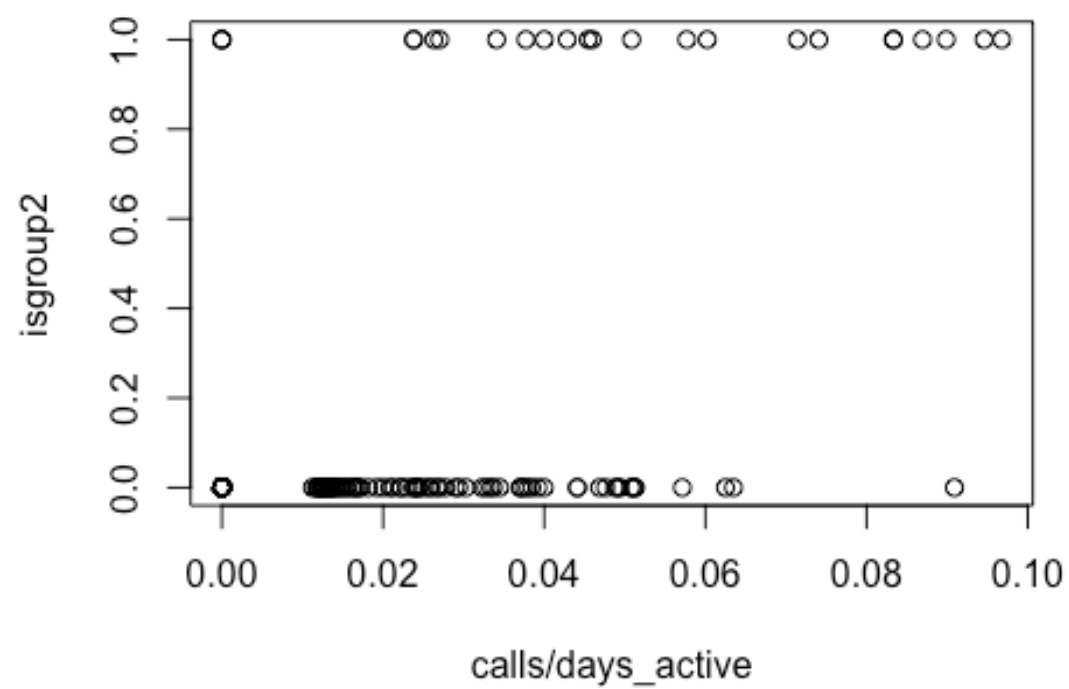
```
plot(calls, isgroup2)
```



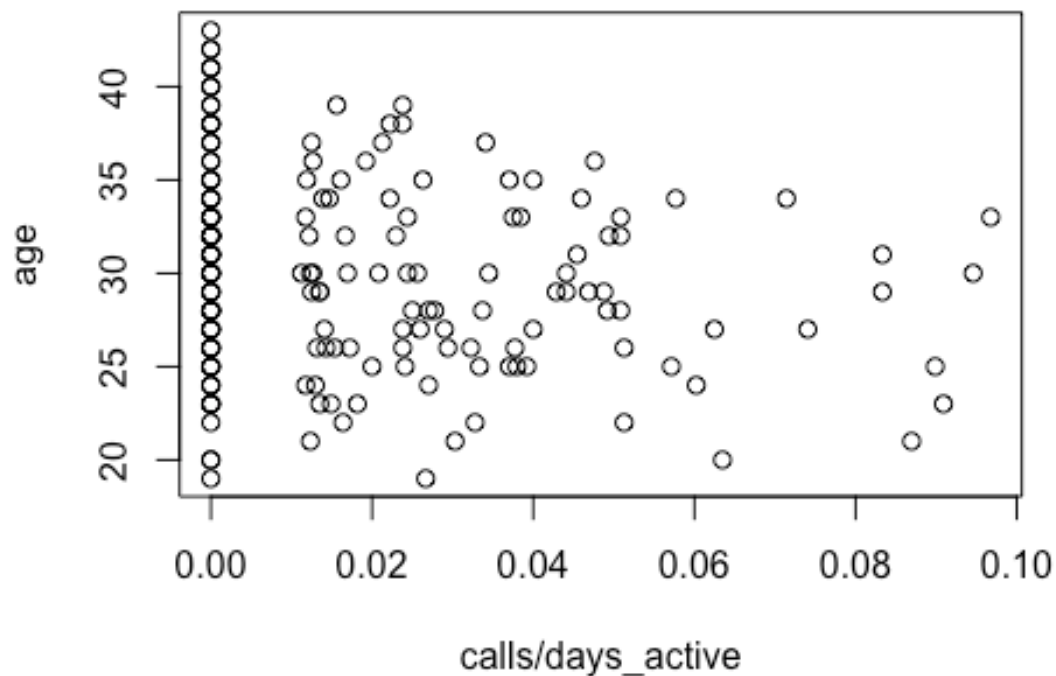
```
plot(age, isgroup2)
```



```
plot(calls/days_active, isgroup2)
```



```
plot(calls/days_active,age)
```



```
mod_string_callers = " model {
  for (i in 1:length(calls)) {
    calls[i] ~ dpois( lam[i] )
    log(lam[i]) = b0 + b1*age[i] + b2*isgroup2[i]
  }

  b0 ~ dnorm(0.0, 1.0/1e2)
  b1 ~ dnorm(0.0, 1.0/1e2)
  b2 ~ dnorm(0.0, 1.0/1e2)
} "
```

```
set.seed(102)
```

```
data_jags_callers = as.list(dat)
```

```
params = c("int", "b0", "b1", "b2")
```

```
mod_c = jags.model(textConnection(mod_string_callers), data=data_jags_callers,
  n.chains=3)
```

```
## Warning in jags.model(textConnection(mod_string_callers), data =
## data_jags_callers, : Unused variable "days_active" in data
```

```

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 224
##   Unobserved stochastic nodes: 3
##   Total graph size: 786
##
## Initializing model

update(mod_c, 1e3)

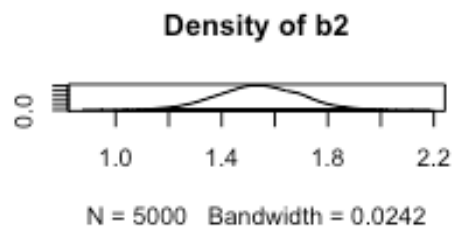
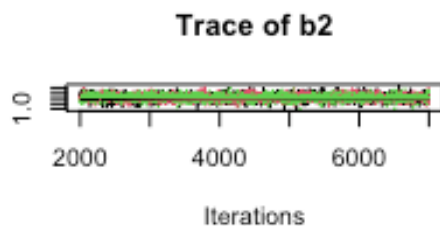
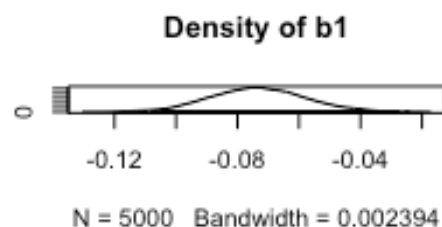
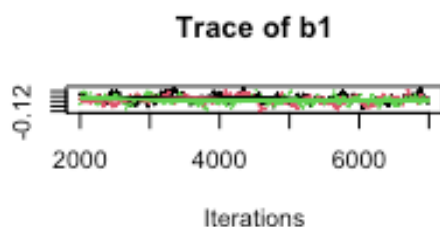
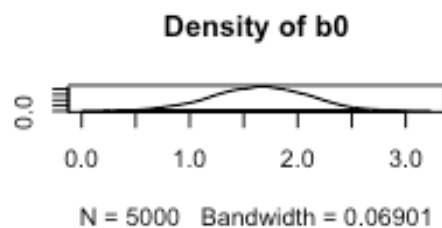
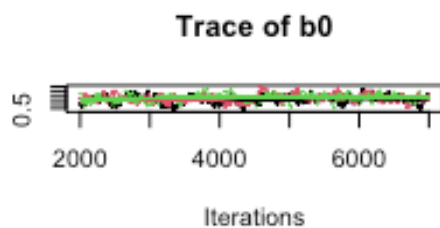
mod_sim_c = coda.samples(model=mod_c,
                        variable.names=params,
                        n.iter=5e3)

## Warning in FUN(X[[i]], ...): Failed to set trace monitor for int
## Variable int not found

mod_csim_c = as.mcmc(do.call(rbind, mod_sim_c))

## convergence diagnostics
plot(mod_sim_c)

```



```
gelman.diag(mod_sim_c)
```

```
## Potential scale reduction factors:
```

```
##
```

```
##      Point est. Upper C.I.
```

```
## b0      1.03      1.12
```

```
## b1      1.04      1.12
```

```
## b2      1.00      1.00
```

```
##
```

```
## Multivariate psrf
```

```
##
```

```
## 1.03
```

```
autocorr.diag(mod_sim_c)
```

```
##              b0              b1              b2
```

```
## Lag 0  1.0000000 1.0000000 1.0000000000
```

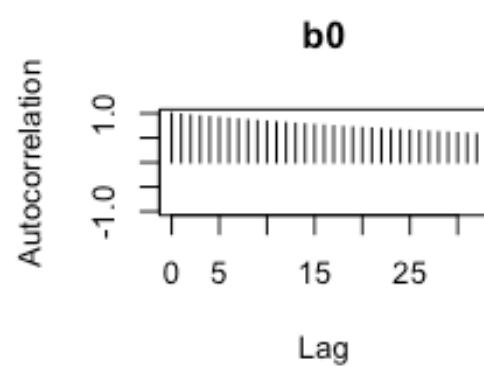
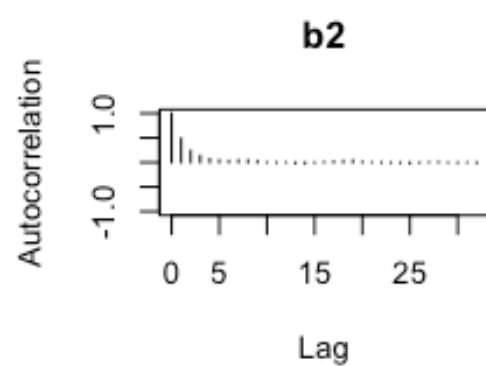
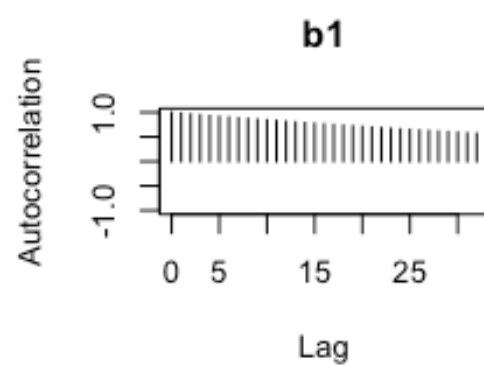
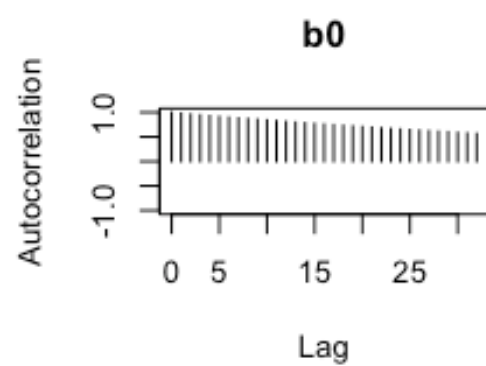
```
## Lag 1  0.9784843 0.9790483 0.499051191
```

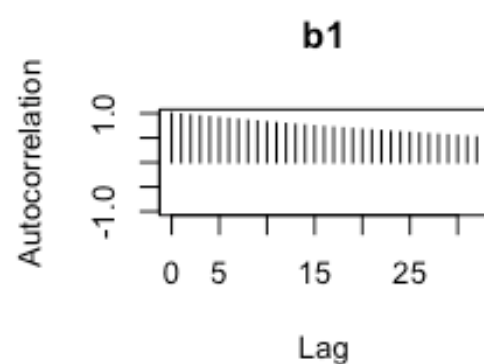
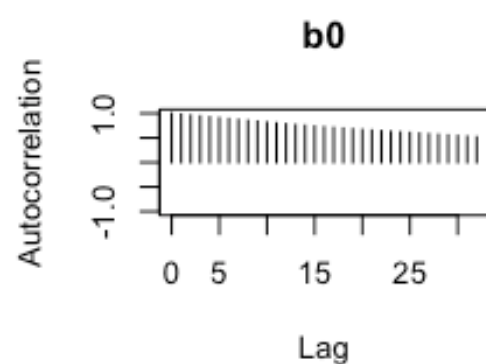
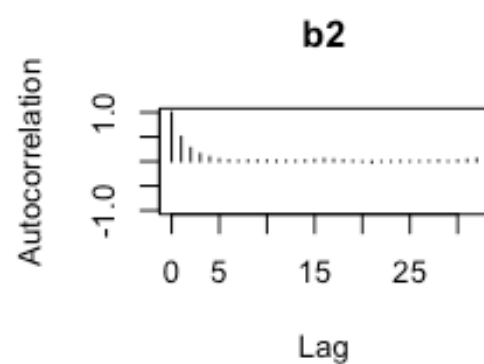
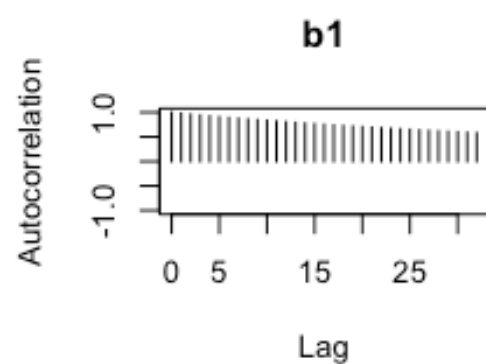
```
## Lag 5  0.9086386 0.9096017 0.041799320
```

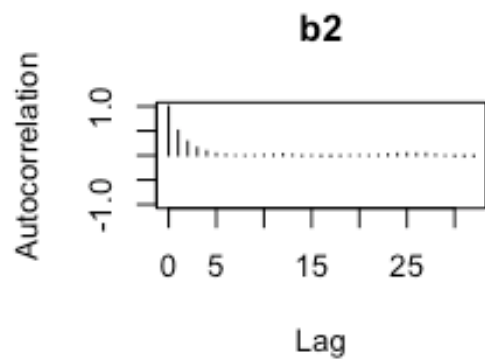
```
## Lag 10 0.8295925 0.8315188 0.009287233
```

```
## Lag 50 0.4092002 0.4152839 0.001305422
```

```
autocorr.plot(mod_sim_c)
```







```
effectiveSize(mod_sim_c)

##          b0          b1          b2
## 141.0077 139.0833 4812.7578

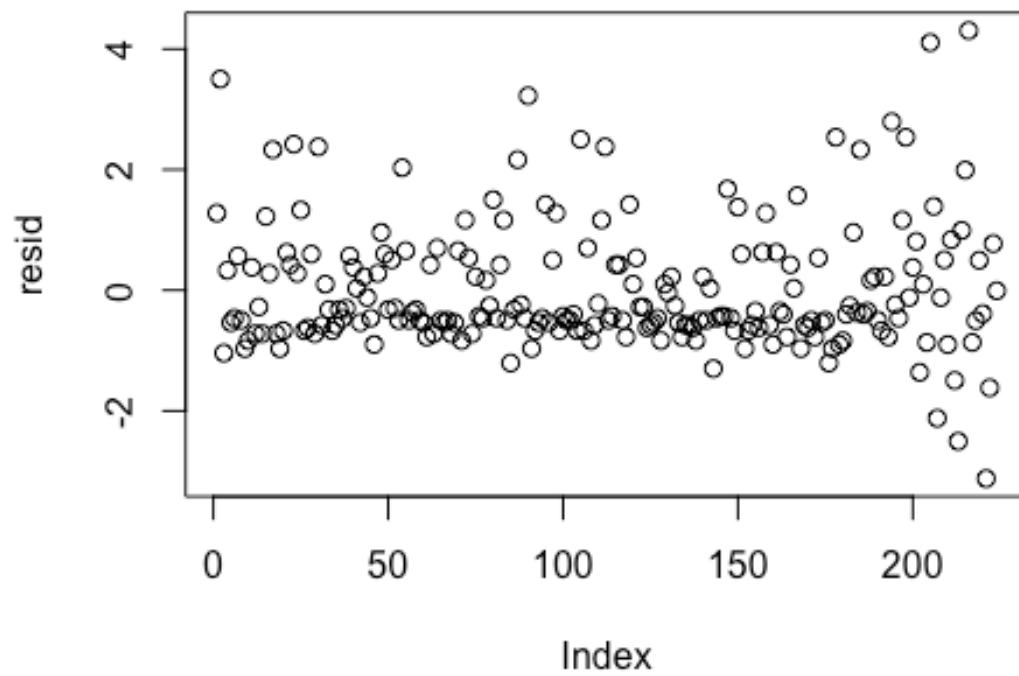
## compute DIC

dic_c = dic.samples(mod_c, n.iter=1e3)

X = as.matrix(dat[,c(4,3)])
(pmed_coef = apply(mod_csim_c, 2, median))

##          b0          b1          b2
## 1.65976285 -0.07359971 1.53960069

llam_hat = pmed_coef["b0"] + X %*% pmed_coef[c("b1", "b2")]
lam_hat = exp(llam_hat)
resid = dat$calls - lam_hat
plot(resid)
```



```
mean(mod_csim_c[, "b2"] > 0)
```

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