

Homework 2

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October 5, 2016

```
library(R2jags)

## Loading required package: rjags

## Loading required package: coda

## Linked to JAGS 4.2.0

## Loaded modules: basemod,bugs

##
## Attaching package: 'R2jags'

## The following object is masked from 'package:coda':
##
##      traceplot

library(lattice) # Needed for scatterplot matrix
# Set working directory
setwd("/home/gsk/grad/courses/UCLA/biost234/homework/lab2")
getwd()

## [1] "/home/gsk/grad/courses/UCLA/biost234/homework/lab2"

# Save the data file to your working directory.

#READ IN DATA
housing=read.table("housingdata2.txt")
head(housing)    ## Always look at your data

##      V1  V2  V3 V4 V5
## 1 15.783 3.00 2.00  2  2
## 2 12.570 1.66 2.33  3  2
## 3 19.600 3.33 2.33  2  2
## 4  8.206 1.66 1.66  2  2
## 5 15.333 2.33 2.33  5  2
## 6 14.955 5.00 3.00  2  2

## Anything funny about any of the columns?
## There are no column names!
str(housing)
```

```
## 'data.frame': 21 obs. of 5 variables:
## $ V1: num 15.78 12.57 19.6 8.21 15.33 ...
## $ V2: num 3 1.66 3.33 1.66 2.33 5 4.33 2.33 1.33 3 ...
## $ V3: num 2 2.33 2.33 1.66 2.33 3 3 2.33 1.66 2.66 ...
## $ V4: int 2 3 2 2 5 2 2 3 2 2 ...
## $ V5: int 2 2 2 2 2 2 2 2 2 2 ...
```

```
colnames(housing) <- c("cost", "eaves", "windows", "yard", "roof")
```

```
#SEPARATE X & Y
```

```
y <- housing[,1]
```

```
x <- as.matrix(housing[,2:5])
```

```
## Remember: look at your data.
```

```
y
```

```
## [1] 15.783 12.570 19.600 8.206 15.333 14.955 13.710 11.388 4.802 12.547
## [11] 13.677 9.683 16.798 25.615 15.734 13.510 13.855 3.986 5.997 9.778
## [21] 10.152
```

```
head(x)
```

```
##      eaves windows yard roof
## [1,] 3.00      2.00    2    2
## [2,] 1.66      2.33    3    2
## [3,] 3.33      2.33    2    2
## [4,] 1.66      1.66    2    2
## [5,] 2.33      2.33    5    2
## [6,] 5.00      3.00    2    2
```

Classical regression

Before beginning our Bayesian analysis, we can conduct classical multiple linear regression:

```
reg = lm(y~x)
```

```
summary(reg) # classical regression
```

```
##
```

```
## Call:
```

```
## lm(formula = y ~ x)
```

```
##
```

```
## Residuals:
```

```
##      Min       1Q   Median       3Q      Max
## -3.8827 -1.8219 -0.9953  1.3467  7.4674
```

```
##
```

```
## Coefficients: (1 not defined because of singularities)
```

```
##              Estimate Std. Error t value Pr(>|t|)
```

```
## (Intercept)  -7.5055      3.9087  -1.920  0.07177 .
```

```
## xeaves        1.8981      0.9868   1.923  0.07133 .
```

```
## xwindows      3.5310      1.6703   2.114  0.04960 *
```

```
## xyard         2.5450      0.8721   2.918  0.00958 **
```

```
## xroof          NA          NA          NA          NA
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.275 on 17 degrees of freedom
## Multiple R-squared:  0.6318, Adjusted R-squared:  0.5668
## F-statistic: 9.722 on 3 and 17 DF,  p-value: 0.0005765
```

The yard and window quality seem to correlate with the contractor's estimates. We see a row of NAs for coefficients of roof, since all of the roofs in this dataset had a quality of 2- so, there is no variation in that measurement.

Bayesian approach

First we write out our model

```
# sink("housingmodel.txt")
cat("
model
{
  for(i in 1:N) {
    y[i] ~ dnorm(mu[i] , tau )
    mu[i] <- beta0 + inprod(x[i,] , beta[] )
  }

  beta0 ~ dnorm( mbeta0 , precbeta0) # mean and precision of beta0 (intercept) defined in R

for (j in 1:K) {
  beta[j] ~ dnorm( m[j] , prec[j] ) # the prior for the four different betas will be provided through
                                     # in all models we will use mean beta values estimated in an old a

  }
  tau ~ dgamma( tau.a , tau.b ) # we tend to use gamma for tau because it is always non-negative
                                # we will weigh the priors differently in a few different models by
                                # fidgeting with the precision in prior
  sigma <- 1 / sqrt(tau)        # note that we use <- here since it's a simple calculation
}
",fill = TRUE)
```

```
##
## model
## {
##   for(i in 1:N) {
##     y[i] ~ dnorm(mu[i] , tau )
##     mu[i] <- beta0 + inprod(x[i,] , beta[] )
##   }
##
##   beta0 ~ dnorm( mbeta0 , precbeta0) # mean and precision of beta0 (intercept) defined in R
##
## for (j in 1:K) {
##   beta[j] ~ dnorm( m[j] , prec[j] ) # the prior for the four different betas will be provided through
##                                     # in all models we will use mean beta values estimated in an old
##
```

```
##      }
##      tau ~ dgamma( tau.a , tau.b ) # we tend to use gamma for tau because it is always non-negative
##                                     # we will weigh the priors differently in a few different models
##                                     # fidgeting with the precision in prior
##      sigma <- 1 / sqrt(tau)        # note that we use <- here since it's a simple calculation
##  }
##
```

```
# sink()
```

Now, we set up three different models:

```
# define some variables
N = nrow(housing)
K = ncol(x) # number of coefficients to estimate betas for
m = c(1.6053, 1.2556, 2.3413, 3.6771) # mean betas from previous analysis

dataA <- list(N=N, K=4, m = m,
             prec = c(.2164, .1105, .2061, .1337), tau.a=17,
             tau.b = 1128, mbeta0= -5.682, precbeta0=.05464, x=x, y=y)

dataB <- list(N=N, K=4, m=m,
             prec=c(.02774, .014160, .02642, .01714), tau.a=2.1795,
             tau.b=144.6, mbeta0= -5.682, precbeta0=.007005, x=x, y=y)

dataC <- list(N=N, K=4, m=m,
             prec=c(.005549, .002832, .005284, .003428), tau.a=.4359,
             tau.b=28.92, mbeta0= -5.682, precbeta0=.00140, x=x, y=y)

inits <- rep(list(list(beta0=0, beta=c(1,1,1,1),tau=1)),5) # 5 equal to the n.chains in jags call

#DEFINE PARAMETERS TO MONITOR
parameters <- c("beta0", "beta" , "tau", "sigma")
```

Then we run jags:

```
#RUN THE JAGS PROGRAM, SAVING DATA TO LAB2.SIM
lab2.simA <- jags (dataA, inits, parameters, "housingmodel.txt", n.chains=5,
                 n.iter=5100, n.burnin=100, n.thin=1, DIC=FALSE) # DIC = F - deviance not calcuated
```

```
## module glm loaded

## module dic loaded

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 21
##   Unobserved stochastic nodes: 6
##   Total graph size: 191
##
## Initializing model
```

```
lab2.simB <- jags (dataB, inits, parameters, "housingmodel.txt", n.chains=5,
  n.iter=5100, n.burnin=100, n.thin=1, DIC=FALSE)
```

```
## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 21
##   Unobserved stochastic nodes: 6
##   Total graph size: 191
##
## Initializing model
```

```
lab2.simC <- jags (dataC, inits, parameters, "housingmodel.txt", n.chains=5,
  n.iter=5100, n.burnin=100, n.thin=1, DIC=FALSE)
```

```
## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
## Graph information:
##   Observed stochastic nodes: 21
##   Unobserved stochastic nodes: 6
##   Total graph size: 191
##
## Initializing model
```

```
lab2.sims = list(lab2.simA, lab2.simB, lab2.simC)
```

```
knitr::kable(lab2.simA$BUGSoutput$summary[,c("mean", "sd", "2.5%", "97.5%")], digits = 3)
```

| | mean | sd | 2.5% | 97.5% |
|---------|--------|-------|---------|-------|
| beta[1] | 1.786 | 1.366 | -0.897 | 4.449 |
| beta[2] | 1.787 | 1.997 | -2.115 | 5.735 |
| beta[3] | 2.233 | 1.316 | -0.351 | 4.796 |
| beta[4] | 2.586 | 2.258 | -1.895 | 7.008 |
| beta0 | -7.010 | 3.830 | -14.576 | 0.497 |
| sigma | 6.931 | 0.688 | 5.749 | 8.440 |
| tau | 0.021 | 0.004 | 0.014 | 0.030 |

```
knitr::kable(lab2.simB$BUGSoutput$summary[,c("mean", "sd", "2.5%", "97.5%")], digits = 3)
```

| | mean | sd | 2.5% | 97.5% |
|---------|--------|-------|---------|--------|
| beta[1] | 1.911 | 1.413 | -0.892 | 4.710 |
| beta[2] | 3.220 | 2.319 | -1.426 | 7.754 |
| beta[3] | 2.480 | 1.263 | -0.025 | 4.961 |
| beta[4] | 1.155 | 4.992 | -8.719 | 10.944 |
| beta0 | -8.876 | 9.604 | -27.656 | 9.930 |
| sigma | 4.840 | 0.771 | 3.611 | 6.591 |

| | mean | sd | 2.5% | 97.5% |
|-----|-------|-------|-------|-------|
| tau | 0.046 | 0.014 | 0.023 | 0.077 |

```
knitr::kable(lab2.simC$BUGSoutput$summary[,c("mean", "sd", "2.5%", "97.5%")], digits = 3)
```

| | mean | sd | 2.5% | 97.5% |
|---------|--------|--------|---------|--------|
| beta[1] | 1.904 | 1.165 | -0.421 | 4.185 |
| beta[2] | 3.480 | 1.972 | -0.402 | 7.388 |
| beta[3] | 2.543 | 1.034 | 0.515 | 4.607 |
| beta[4] | 0.839 | 10.690 | -20.247 | 21.808 |
| beta0 | -9.059 | 21.239 | -50.290 | 32.592 |
| sigma | 3.829 | 0.689 | 2.758 | 5.437 |
| tau | 0.074 | 0.025 | 0.034 | 0.131 |

Question 2

2. Give a table of inferences for the coefficient of roofs for the three priors. Briefly explain why it comes out as it does.

```
temp3 <- t(sapply(lab2.sims, function(x) x$BUGSoutput$summary["beta[4]",c("mean", "sd")]))
rownames(temp3) <- c("Model 1", "Model 2", "Model 3")
knitr::kable(temp3)
```

| | mean | sd |
|---------|-----------|-----------|
| Model 1 | 2.5864595 | 2.257813 |
| Model 2 | 1.1547660 | 4.992266 |
| Model 3 | 0.8393065 | 10.689721 |