

2c

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8 May 2017

- (b) Use WinBUGS for a Bayesian analysis of (a), and find reasonable point and interval estimates for σ_1^2 and σ_2^2 . Include graphs, including one of the joint posterior. [6 marks]

```
(BT3 <- summary(aov(Sample~(I/J/K),df3)))
(m1 <- BT3[[1]]$`Mean Sq`[3])
(m2 <- (BT3[[1]]$`Mean Sq`[2]-BT3[[1]]$`Mean Sq`[3])/2)
(m3 <- (BT3[[1]]$`Mean Sq`[1]-BT3[[1]]$`Mean Sq`[2])/4)
(mAll <- m1+m2+m3)
(m1/mAll)
(m2/mAll)
(m3/mAll)

#define the model
library(R2OpenBUGS)
boxtiao3model <- function(){
  for(i in 1 : I) {
    m[i] ~ dnorm(theta, tau.I)
    for(j in 1 : J) {
      y[i , j] ~ dnorm(m[i], tau.J)
      for(k in 1: K){
        z[i,j,k] ~ dnorm(y[i,j], tau.K)
      }
    }
  }
  sigma2.I <- 1/tau.I
  tau.I ~ dgamma(0.001, 0.001)
  sigma2.J <- 1/tau.J
  tau.J ~ dgamma(0.001, 0.001)
  sigma2.K <- 1/tau.K
  tau.K ~ dgamma(0.001, 0.001)
  theta ~ dflat()
}

# write the model code out to a file
write.model(boxtiao3model, "boxtiao3model.txt")
model.file1 = paste(getwd(),"boxtiao3model.txt", sep="/")
#prepare the data for input into OpenBUGS
I <- 10
J <- 2
K <- 2
y <- as.matrix(df3)

data <- list ("I","J","K","y", .Dim = c(10,2,2))

#initialization of variables
inits <- function(){
  list(theta=1, tau.I=1, tau.J=4, tau.K=2)
}
```

```

WINE="/opt/local/bin/wine"
WINEPATH="/opt/local/bin/winepath"
OpenBUGS.pgm=paste0("/Users/benjamin/Applications/wine/",
                    "drive_c/ProgramFiles/OpenBUGS/OpenBUGS323/OpenBUGS.exe")

#these are the parameters to save
parameters = c("sigma2.I", "sigma2.J", "sigma2.K")

#run the model
bt3.sim <- bugs(data,
               inits,
               model.file = model.file1,
               parameters=parameters,
               n.burnin = 2000,
               n.chains = 5,
               n.iter = 10000,
               OpenBUGS.pgm=OpenBUGS.pgm,
               WINE=WINE,
               WINEPATH=WINEPATH,
               useWINE=T,
               codaPkg = T,
               debug = T)

out.coda <- read.bugs(dyes.sim)
plot(out.coda)
HPDinterval(out.coda)
densityplot(out.coda)
plot(out.coda)
(out.summary <- summary(out.coda, q=c(0.025, 0.975)))

sigma2_with_stat <- c(out.summary$statistics[2,1],
out.summary$q["sigma2.btw",])
sigma2_btw_stat <- c(out.summary$statistics[3,1],
out.summary$q["sigma2.with",])

summary_stats <- rbind(sigma2.with = sigma2_with_stat, sigma2.btw = sigma2_btw_stat)
colnames(summary_stats)[1] <- "Mean"
summary_stats

dfI <- rep(seq(1,10),4)
dfJ <- c(rep(1,20),rep(2,20))
dfK <- c(rep(1,10),rep(2,10),rep(1,10),rep(2,10))
df <- cbind(dfI,dfJ,dfK)

dfJ1K1 <- c(2.004,4.342,0.869,3.531,2.579,-1.404,-1.676,1.670,2.141,-1.229)
dfJ1K2 <- c(2.713,4.229,-2.621,4.185,4.271,-1.003,-0.208,2.426,3.527,-0.596)
dfJ2K1 <- c(0.602,3.344,-3.896,1.722,-2.101,-0.755,-9.139,1.834,0.462,4.471)
dfJ2K2 <- c(0.252,3.057,-3.696,0.380,0.651,-2.202,-8.653,1.200,0.665,1.606)
df2 <- c(dfJ1K1,dfJ1K2,dfJ2K1,dfJ2K2)
df <- NA
for(i in 1:length(dfJ1K1)){
  test <- c(dfJ1K1[i],dfJ1K2[i],dfJ2K1[i],dfJ2K2[i])
  df <- c(df,test)
}

```

```

# Model

model{
  for(i in 1: I){
    a[i] ~ dnorm(0.0, tauI)
    for(j in 1:J){
      b[i,j] ~ dnorm(0.0, tauJ)
      for(k in 1:K){
        e[i,j,k] <- theta + a[i] + b[i,j]
        y[i,j,k] ~ dnorm(e[i,j,k], tauK)
      }
    }
  }
}

# Priors

theta ~ dflat()
tauI ~ dgamma(0.001, 0.001)
tauJ ~ dgamma(0.001, 0.001)
tauK ~ dgamma(0.001, 0.001)

sigma2I <-1/tauI
sigma2J <-1/tauJ
sigma2K <-1/tauK

denom <- sigma2I + sigma2J + sigma2K
r1 <- sigma2I/denom
r2 <- sigma2J/denom
r3 <- sigma2K/denom
}

# Initial values

INITS

list(
  theta=0,
  a = c(0, 0, 0, 0, 0, 0, 0, 0, 0, 0),
  b = structure(
    .Data = c(0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
              0, 0, 0, 0, 0, 0, 0, 0, 0, 0),
    .Dim = c(10, 2)
  ),
  tauI = 1.0,
  tauJ = 1.0,
  tauK = 1.0
)

# Data (in tabular format)

DATA

list(

```

```

I=10,
J=2,
K=2,
y = structure(
  .Data=c(2.004, 2.713, 0.602, 0.252,
          4.342, 4.229, 3.344, 3.057,
          0.869, -2.621, -3.896, -3.696,
          3.531, 4.185, 1.722, 0.380,
          2.579, 4.271, -2.101, 0.651,
          -1.404, -1.003, -0.755, -2.202,
          -1.676, -0.208, -9.139, -8.653,
          1.670, 2.426, 1.834, 1.200,
          2.141, 3.527, 0.462, 0.665,
          -1.229, -0.596, 4.471, 1.606),
  .Dim=c(10, 2, 2)
)
)

```

```

#define the model
library(R2OpenBUGS)
threecompmode1 <- function(){
  for( i in 1 : I ) {
    a[i] ~ dnorm(theta, tauI)
    for( j in 1 : J ) {
      b[i , j] ~ dnorm(a[i], tauJ)
      for( k in 1 : K ) {
        e[i , j , k] <- mu0 + a[i] + b[i , j]
        y[i , j , k] ~ dnorm(e[i , j , k], tauK)
      }
    }
  }
}

## Priors

theta ~ dnorm(0.0,1.00E-04) # Batch Mean
mu0 ~ dnorm(0.0,1.00E-04) # Sample Mean
tauI ~ dgamma(0.001, 0.001) # Between Batch variation
tauJ ~ dgamma(0.001, 0.001) # Between Sample variation
tauK ~ dgamma(0.001, 0.001) # Between Test variation

sigma2I <- 1 / tauI
sigma2J <- 1 / tauJ
sigma2K <- 1 / tauK

sigma2JK <- sigma2J / sigma2K

denom <- sigma2I + sigma2J + sigma2K
r1 <- sigma2I / denom
r2 <- sigma2J / denom
r3 <- sigma2K / denom
}

# write the model code out to a file
write.model(threecompmode1, "threecompmode1model.txt")
model.file.3comp = paste(getwd(),"threecompmode1model.txt", sep="/")

```

```

#prepare the data for input into OpenBUGS

data <-
  list(I=10,J=2,K=2,y = structure(
    .Data=c(2.004, 2.713, 0.602, 0.252,
            4.342, 4.229, 3.344, 3.057,
            0.869, -2.621, -3.896, -3.696,
            3.531, 4.185, 1.722, 0.380,
            2.579, 4.271, -2.101, 0.651,
            -1.404, -1.003, -0.755, -2.202,
            -1.676, -0.208, -9.139, -8.653,
            1.670, 2.426, 1.834, 1.200,
            2.141, 3.527, 0.462, 0.665,
            -1.229, -0.596, 4.471, 1.606),
    .Dim=c(10,2,2)
  ))

#initialization of variables
inits <- function(){
  list(
    mu0 = 0,
    theta=0,
    a = c(0,0,0,0,0,0,0,0,0,0),
    b = structure(
      .Data = c(0,0,0,0,0,0,0,0,0,0,
                0,0,0,0,0,0,0,0,0,0),
      .Dim = c(10,2)
    ),
    tauI = 1.0,
    tauJ = 1.0,
    tauK = 1.0
  )
}

WINE="/opt/local/bin/wine"
WINEPATH="/opt/local/bin/winepath"
OpenBUGS.pgm=paste0("/Users/benjamin/Applications/wine/",
                    "drive_c/ProgramFiles/OpenBUGS/OpenBUGS323/OpenBUGS.exe")

#these are the parameters to save
parameters = c("sigma2I","sigma2J","sigma2K", "sigma2JK", "r1", "r2", "r3")

#run the model
threecomp.sim <- bugs(data,
  inits,
  model.file = model.file.3comp,
  parameters=parameters,
  n.burnin = 5000,
  n.chains = 5,
  n.iter = 50000,
  OpenBUGS.pgm=OpenBUGS.pgm,
  WINE=WINE,
  WINEPATH=WINEPATH,

```

```

        useWINE=T,
        codaPkg = F,
        debug = F)

library(ggplot2)
library(ggtern)
attach.bugs(threecomp.sim)

threecompsimlist <- tbl_df(threecomp.sim$sims.list)

# densityplot(threecompsimlist$sigma2I)
# densityplot(threecompsimlist$sigma2J)
# densityplot(threecompsimlist$sigma2K)

# sig2I <- threecompsimlist$sigma2I
# sig2J <- threecompsimlist$sigma2J
# sig2K <- threecompsimlist$sigma2K

df <- tbl_df(threecompsimlist$r1)
df <- tbl_df(cbind(rBatches = threecompsimlist$sigma2I, rSamples = threecompsimlist$sigma2J, rTests = threecompsimlist$sigma2K))
denom <- sum(threecomp.sim$summary[1:3,1])

rI <- threecomp.sim$summary[1,1]/denom
rJ <- threecomp.sim$summary[2,1]/denom
rK <- threecomp.sim$summary[3,1]/denom

ggtern(data = df, aes(x=rBatches,z=rSamples,y=rTests)) +
  theme_rgbw() +
  geom_point(size=0.1, colour = "red") +
  geom_confidence_tern(breaks = c(0.95)) +
  theme_clockwise() +
  labs(x="Batches",z="Samples",y="Tests",title="Contours of the posterior distribution of (r1, r2, r3)")

```