Methods-for-Utility-Evaluation

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```
library(ggplot2)
CEdata <- read.csv("CEdata.csv")</pre>
CEdata$LogIncome <- log(CEdata$Income)</pre>
CEdata$LogExpenditure <- log(CEdata$Expenditure)</pre>
require(runjags)
require(coda)
modelString <-"
model {
## sampling
for (i in 1:N){
y[i] ~ dnorm(beta0 + beta1*x[i], invsigma2)
## priors
beta0 ~ dnorm(mu0, g0)
beta1 ~ dnorm(mu1, g1)
invsigma2 ~ dgamma(a, b)
sigma <- sqrt(pow(invsigma2, -1))</pre>
}"
y <- as.vector(CEdata$LogIncome)</pre>
x <- as.vector(CEdata$LogExpenditure)</pre>
N <- length(y)
the_data <- list("y" = y, "x" = x, "N" = N,
                 "mu0" = 0, "g0" = 0.0001,
                 "mu1" = 0, "g1" = 0.0001,
                 a'' = 1, b'' = 1
initsfunction <- function(chain){</pre>
.RNG.seed \leftarrow c(1,2) [chain]
.RNG.name <- c("base::Super-Duper",
"base::Wichmann-Hill")[chain]
return(list(.RNG.seed=.RNG.seed,
.RNG.name=.RNG.name))
}
posterior <- run.jags(modelString,</pre>
                       n.chains = 1,
                        data = the_data,
                       monitor = c("beta0", "beta1", "sigma"),
                       adapt = 1000,
                       burnin = 5000,
                       sample = 5000,
                       thin = 50,
                        inits = initsfunction)
```

```
## Calling the simulation...
## Welcome to JAGS 4.3.0 on Sun Feb 23 22:02:27 2020
## JAGS is free software and comes with ABSOLUTELY NO WARRANTY
## Loading module: basemod: ok
## Loading module: bugs: ok
## . . Reading data file data.txt
## . Compiling model graph
     Resolving undeclared variables
##
##
     Allocating nodes
## Graph information:
     Observed stochastic nodes: 994
##
     Unobserved stochastic nodes: 3
     Total graph size: 3990
## . Reading parameter file inits1.txt
## . Initializing model
## . Adaptation skipped: model is not in adaptive mode.
## . Updating 5000
## -----| 5000
## ************* 100%
## . . . . Updating 250000
## -----| 250000
## ********** 100%
## . . . . Updating 0
## . Deleting model
## Note: the model did not require adaptation
## Simulation complete. Reading coda files...
## Coda files loaded successfully
## Calculating summary statistics...
## Finished running the simulation
library(coda)
post <- as.mcmc(posterior)</pre>
synthesize <- function(X, index, n){</pre>
mean_Y <- post[index, "beta0"] + X * post[index, "beta1"]</pre>
synthetic_Y <- rnorm(n, mean_Y, post[index, "sigma"])</pre>
data.frame(X, synthetic_Y)
}
n <- dim(CEdata)</pre>
m < -20
synthetic_m <- vector("list", m)</pre>
for (1 in 1:m){
synthetic one <- synthesize(CEdata$LogExpenditure, 4980+1, n)
names(synthetic_one) <- c("logExpenditure", "logIncome_syn")</pre>
synthetic_m[[1]] <- synthetic_one</pre>
}
mean_syn_m <- vector("list", m)</pre>
for (1 in 1:m){
  mean_syn_m[[1]] <- mean(synthetic_m[[1]]$logIncome_syn)</pre>
}
```

```
median_syn_m <- vector("list", m)
for (l in 1:m){
    median_syn_m[[1]] <- median(synthetic_m[[1]]$logIncome_syn)
}

regression_m <- vector("list", m)
for (l in 1:m){
    income <- synthetic_m[[1]]$logIncome_syn
    expenditure <- synthetic_m[[1]]$logExpenditure
    linearMod <- lm(logIncome_syn ~ logExpenditure, data = synthetic_m[[1]])
    regression_m[[1]] <- linearMod$coefficients[2]
}</pre>
```

Fully synthetic data

```
mean_syn <- unlist(mean_syn_m, use.names = FALSE)</pre>
median_syn <- unlist(median_syn_m, use.names = FALSE)</pre>
regression_syn <- unlist(regression_m, use.names = FALSE)</pre>
u_mean <- var(mean_syn)
u_median <- var(median_syn)</pre>
u_regression <- var(regression_syn)</pre>
q_bar_mean <- mean(mean_syn)</pre>
q_bar_median <- mean(median_syn)</pre>
q_bar_regression <- mean(regression_syn)</pre>
b_m = - sum((mean_syn - q_bar_mean)^2) / (m - 1)
b_m_median <- sum((median_syn - q_bar_median)^2) / (m - 1)</pre>
b_m_regression <- sum((regression_syn - q_bar_regression)^2) / (m - 1)</pre>
u_bar_mean <- sum(u_mean) / m</pre>
u bar median <- sum(u median) / m
u_bar_regression <- sum(u_regression) / m</pre>
T_{mean} \leftarrow (1 + m^{(-1)}) * b_{m_{mean}} - u_{bar_{mean}}
T_{median} \leftarrow (1 + m^{(-1)}) * b_{m_{median}} - u_{bar_{median}}
T_{regression} \leftarrow (1 + m^{-1}) * b_{m_{regression}} - u_{bar_{regression}}
```

For the mean of the synthesized log income,

```
q_bar_mean

## [1] 10.1876

T_mean

## [1] 0.4372374

For the median of the synthesized log income,
```

q_bar_median

```
## [1] 10.1876
T_median
```

```
## [1] 0.4372374
```

For the regression coefficients between the synthesized log income and the original logged expenditure,

```
q_bar_regression
## [1] -0.002719087
T_regression
## [1] 0.003523324
Partial synthetic data
T mean p <- b m mean/m + u bar mean
T_median_p <- b_m_median/m + u_bar_median</pre>
T_regression_p <- b_m_regression/m + u_bar_regression</pre>
For the mean of the synthesized log income,
q_bar_mean
## [1] 10.1876
T_mean_p
## [1] 0.04372374
For the median of the synthesized log income,
q_bar_median
## [1] 10.1876
T_median_p
## [1] 0.04372374
For the regression coefficients between the synthesized log income and the original logged expenditure,
q_bar_regression
## [1] -0.002719087
T_regression_p
```

[1] 0.0003523324

Overlay interval

```
syn_data <- synthetic_m[[1]]
logincome_syn <- syn_data$logIncome_syn
logincome_original <- CEdata$LogIncome
L_s <- unname(quantile(logincome_syn, 0.025))
U_s <- unname(quantile(logincome_syn, 0.975))
L_o <- unname(quantile(logincome_original, 0.025))
U_o <- unname(quantile(logincome_original, 0.975))
L_i <- max(L_s, L_o)
U_i <- min(U_s, U_o)
I <- (U_i - L_i) / (2 * (U_o - L_o)) + (U_i - L_i) / (2 * (U_s - L_s))
I</pre>
```

[1] 0.7618307

Since data with high utility would have a score of 1 and no data with no utility will have a score of 0, our synthesized data has a medium utility.

Project synthesize

```
bnbData <- read.csv("AB_NYC_2019.csv")</pre>
length <- dim(bnbData)</pre>
avail <- bnbData$availability 365
Category <- c()</pre>
for(i in 1:length){
  if (avail[i] > 330){
    Category <- c(Category, 1)</pre>
  }else if( avail[i] <= 330 & avail[i] > 270){
    Category <- c(Category, 2)</pre>
  }else if( avail[i] <= 270 & avail[i] > 60){
    Category <- c(Category, 3)</pre>
  }else{
    Category <- c(Category, 4)</pre>
}
## Warning in 1:length: numerical expression has 2 elements: only the first
## used
room_type <- bnbData$room_type</pre>
room_category <- c()</pre>
for(i in 1:length){
  if (room_type[i] == "Private room"){
    room_category <- c(room_category, 1)</pre>
  }else if( room_type[i] == "Entire home/apt"){
    room_category <- c(room_category, 2)</pre>
  }else{
    room_category <- c(room_category, 3)</pre>
  }
}
## Warning in 1:length: numerical expression has 2 elements: only the first
neigh <- bnbData$neighbourhood_group</pre>
neigh_category <- c()</pre>
for(i in 1:length){
  if (neigh[i] == "Brooklyn"){
    neigh_category <- c(neigh_category, 1)</pre>
  }else if(neigh[i] == "Manhattan"){
    neigh_category <- c(neigh_category, 2)</pre>
  }else if(neigh[i] == "Queens"){
    neigh_category <- c(neigh_category, 3)</pre>
  }else if(neigh[i] == "Staten Island"){
    neigh_category <- c(neigh_category, 4)</pre>
```

```
}else{
    neigh_category <- c(neigh_category, 5)</pre>
  }
}
## Warning in 1:length: numerical expression has 2 elements: only the first
## used
bnbData_cat <- data.frame(bnbData, category = Category, room_category = room_category, neigh_category =</pre>
neigh_unique <- unique(bnbData_cat$neighbourhood_group)</pre>
room_unique <- unique(bnbData_cat$room_type)</pre>
price <- bnbData$price</pre>
modelString <-"
model {
## sampling
for (i in 1:N){
  ez<- t (beta)%*%X[i, ]
  a \leftarrow max(-Inf, g[y[i] -1], na.rm=TRUE)
  b<-min( g[y[i]] , Inf , na . rm=TRUE)</pre>
  u<-runif(1 , pnorm( a-ez ) , pnorm(b-ez ) )</pre>
  z[i] \leftarrow ez + qnorm(u)
  a < -max(z[y==k])
  b < -min(z[y==k+1])
  sig[i] \sim dnorm(0,1)
  u<-runif(1 , pnorm( ( a - ez ) / sig[i] ) , pnorm(( b - ez ) / sig[i] ))
  g[i] \leftarrow ez + sig[i] * qnorm(u)
beta ~ dnorm(n/((n+1) * (t(X)%*%X)) %*% t(X) %*% z, n/((n+1) * (t(X)%*%X)))
X <- cbind(room_category, neigh_category, price)</pre>
y <- bnbData_cat$category</pre>
```