library(haven)

library(BMA)

data <- read\_sav("data.sav")

head(data)

str(data)

mydata <- as.matrix(data)

x<-mydata[,-1]

y<-mydata[,1]

bayes1<-bicreg(x,y,maxCol=7)

bayes1

summary(bayes1)

plot(bayes1)