pset 1 631

```
# ==== Inustrial Organization Problem Set 1 ====
# clear objects
rm(list = ls(pos = ".GlobalEnv"), pos = ".GlobalEnv")
options(scipen = 999)
cat("\f")
# Load packages
library(data.table)
library(stats4)
library(broom)
library(AER)
library(xtable)
library(Matrix)
library(BLPestimatoR)
library(SQUAREM)
library(BB)
# set save option
opt_save <- TRUE
# set path for output
f_out <- "C:/Users/Nmath_000/Documents/Code/courses/econ 631/ps1/tex/"
#=====#
# ==== Question 1 ====
#=====#
# laod data
q1dt <- fread("file:///C:/Users/Nmath_000/Documents/MI_school/Third Year/Econ 631/ps1/ps1.dat")
# create variable names
setnames(q1dt, colnames(q1dt), c("y", "x1", "x2", "z"))
#=====#
# ==== part 1 ====
#=====#
# write log likelihood function
Probit_llf <- function(th0, th1, th2){</pre>
```

```
mu <- q1dt[, pnorm(th0 + th1*x1 + th2*x2)]</pre>
  -sum(q1dt[, y*log(mu) + (1-y)*log(1-mu)])
# create starting values
prob_start <- list(th0 = 0,</pre>
                   th1 = .01,
                   th2 = .01)
# run the mle function
probit_res <- mle(Probit_llf, start = prob_start)</pre>
# get the results I need
probit_res <- data.table(variable = rownames(summary(probit_res)@coef),</pre>
                         summary(probit_res)@coef)
# check my results
check \leftarrow glm( y ~ x1+ x2,
              family = binomial(link = "probit"),
              data = q1dt)
# looks good
summary(check)
probit_res
#----#
# ==== Part2 ====
#=====#
probit_res$Estimate[3]*dnorm(probit_res$Estimate[1] + probit_res$Estimate[2]*mean(q1dt$x1) + probit_res
#=====#
# ==== Part3 ====
#=====#
# define logit log likelihood
logit_llf <- function(th0, th1, th2){</pre>
 mu <- q1dt[, plogis(th0 + th1*x1 + th2*x2)]</pre>
  -sum(q1dt[, y*log(mu) + (1-y)*log(1-mu)])
# create startign values
logit_start <- list(th0 = 0,</pre>
                    th1 = .01,
                    th2 = .01)
# run the mle function
logit_res <- mle(logit_llf, start = logit_start)</pre>
```

```
# get the results I need
logit_res <- data.table(variable = rownames(summary(logit_res)@coef),</pre>
                        summary(logit res)@coef)
# check my results
check \leftarrow glm( y ~ x1+ x2,
              family = binomial(link = "logit"),
              data = q1dt)
# looks good
summary(check)
logit_res
#======#
# ==== Part 6 ====
#----#
p6_llf <- function(th0, th1, th2, th3, th4, th5, p, sig){
  # define some intermediate terms
  q1dt[, psi := (p/sig^2)*(x2 - (th3 + th4*x1 + th5*z))]
  q1dt[, tau := 1-(p^2/sig^2)]
  q1dt[, m:= (-th0 - th1*x1 - th2*x2 - psi)/(tau^.5)]
  # get prob y equals one conditional on x1, x2, z, theta
  q1dt[, p_y_1 := 1- pnorm(m)]
  # get pro x2 = x2i given x1 z theta
  q1dt[, p_x := dnorm((x2 - th3 - th4*x1 - th5*z)/sig)]
  # qet log likelihoods
  q1dt[, llf := y*log(p_y_1) + (1-y)*log(1-p_y_1) + log(p_x) - log(sig)]
  # sum them
  sum_llf <- q1dt[, sum(llf)]</pre>
  # remove extra vars
  q1dt[ , := ` (psi = NULL, tau = NULL, m = NULL, p_y_1 = NULL, p_x = NULL, llf = NULL)]
  # return negative sum
 return(-sum_llf)
}
second_stage <- glm(y ~ x1+ x2,
                    family = binomial(link = "probit"),
                    data = q1dt)
first_stage <- lm(x2 \sim x1 + z, data = q1dt)
p6_start <- as.list(c(second_stage$coefficients, first_stage$coefficients, .1, 1))
names(p6_start) <- c( paste0("th", 0:5), "p", "sig")</pre>
```

```
# run the mle function
p6_res <- mle(p6_llf, start = p6_start)
logLik(p6_res)
# get the results I need
p6_res_tab <- data.table(variable = rownames(summary(p6_res)@coef),</pre>
                       summary(p6_res)@coef)
p6_res_tab
#======#
# ==== Question 2 ====
#======#
  #=====#
  # ==== part 3 ====
  #=====#
   # load cereal data
   cereal <- data.table(readxl::read_excel("C:/Users/Nmath_000/Documents/MI_school/Third Year/Econ 631</pre>
    # get total market share by city year
   cereal[, s0 := 1-sum(share), c("city", "year", "quarter" )]
    # create column for mean utility
   cereal[, m_u := log(share) - log(s0)]
    #=====#
    # ==== a ====
   #=====#
   # run ols, tidy it up, make it a data.table
   q2_p3_ols <- data.table(tidy(lm(m_u ~ mushy + sugar + price , data = cereal)))
   # round p value
   q2_p3_ols[, p.value := round(p.value, 6)]
   #=====#
    # ==== b ====
    #=====#
   # create sugar instroment
   cereal[, (.N-1), c('firm_id', "city", "quarter")]
   cereal[, i1_sugar := (sum(sugar) - sugar)/ (.N-1), c('firm_id', "city", "quarter")]
   # create mush intrument
```

```
cereal[, i1_mushy := (sum(mushy) - mushy)/(.N-1), c('firm_id', "city", "quarter")]
   # create price instrument
   cereal[, i1_price := (sum(price) - price)/ (.N-1), c('firm_id', "city", "quarter")]
   # now do 2sls, tidy it up, make it a data.table
   q2_p3_iv1 <- data.table(tidy(ivreg(m_u ~ mushy + sugar + price | i1_sugar + i1_mushy + mushy + sug
   q2_p3_iv1[, p.value := round(p.value,6)]
   #=====#
   # ==== c ====
   #=====#
   # now create second set of instroments
   # would be ideal to write functino for this, but who has the time
   # create sugar instroment
   # start by getting sum of sugar for firm
   cereal[, f_sugar_sum := sum(sugar), c( "city", "quarter", "firm_id")]
   # get number of products by firm
   cereal[, f_nprod := .N, c( "city", "quarter", "firm_id")]
   # get sum of sugar in market, subtract off sum of sugar for the firm
   # divide by number of products minus the products for this firm that we subtracted off
   cereal[, i2_sugar := (sum(sugar) - f_sugar_sum)/ (.N-f_nprod), c("city", "quarter")]
   # now do the same for the other
   cereal[, f_mushy_sum := sum(mushy), c( "city", "quarter", "firm_id")]
   cereal[, i2_mushy := (sum(mushy) - f_mushy_sum)/ (.N-f_nprod), c("city", "quarter")]
   # #note dont actually need this
   # cereal[, f_price_sum := sum(price), c( "city", "quarter", "firm_id")]
   \# cereal[, i2\_price := (sum(price) - f\_price\_sum)/(.N-f\_nprod), c("city", "quarter")]
   # now do 2sls
   ivreg_out <- ivreg(m_u ~ mushy + sugar + price | i2_sugar + i2_mushy + mushy + sugar , data = cerea
   q2_p3_iv2 <- data.table(tidy(ivreg(m_u ~ mushy + sugar + price | i2_sugar + i2_mushy + mushy + sugar
   q2_p3_iv2[, p.value := round(p.value,6)]
#=====#
# ==== Question 3 ====
#=====#
 #=======#
 # ==== set up data for functions ====
 #=======#
```

```
# load cereal data
cereal <- data.table(readxl::read_excel("C:/Users/Nmath_000/Documents/MI_school/Third Year/Econ 631</pre>
#note these are data manipulations that can happen outside the outer loop. We
# just need to convert some things to matrices since we actually need to
# use matrix algebra in this question
# create a single market variale. Don't have to referece two variables then
cereal[, mkt := paste0(city, "_", quarter)]
# get total market share by city year
#note: should we do this before or after dropping obs with missing instruments?
cereal[, s0 := 1-sum(share), mkt]
# create column for mean utility
cereal[, m_u := log(share) - log(s0)]
# create sugar instrument
cereal[, (.N-1), c('firm_id', "city", "quarter")]
cereal[, i1_sugar := (sum(sugar) - sugar)/ (.N-1), c('firm_id', "city", "quarter")]
# create mush instrument
cereal[, i1_mushy := (sum(mushy) - mushy)/(.N-1), c('firm_id', "city", "quarter")]
# drop observations with missing instruments. Missing becasue firm only has one product
cereal <- cereal[!is.na(i1_sugar)]</pre>
# defint some variables
# I'm not crazy about doing this. I think this either needs to be written as a function
# or else we should just hard code the column names. This is like a weird middle ground
# of generalizability that isn't that helpful
                "share"
share.fld =
prod.id.fld = "product id"
                "mkt"
mkt.id.fld =
prc.fld =
                "price"
x.var.flds =
              c("sugar",
                  "mushv")
# order data and get nrows
cereal <- setorder(cereal, "city", "product_id")</pre>
JM <- nrow(cereal)</pre>
# make matrix of controls X
X <- as.matrix(cereal[, c(x.var.flds), with = FALSE])</pre>
K \leftarrow ncol(X)
# make a matrix that includes price
cereal[, n_price := -price]
XP <- as.matrix(cereal[, c("sugar", "mushy", "n_price"), with = FALSE])</pre>
# make a vector for price
```

```
P <- as.matrix(cereal[, price])</pre>
  # put market into a vector
 mkt.id <- cereal[, get(mkt.id.fld)]</pre>
 # put shares into a vector
 s.jm <- as.vector(cereal[, get(share.fld)]);</pre>
 # put shates into a vector
 s.j0 <- cereal[, s0]
 # set number of simulated consumers
 n.sim = 20
 # Standard normal distribution draws, one for each characteristic in X
 # columns are simulated consumers, rows are variables in X (including constant and price)
 # as Ying pointed out we want to do this once outside the outer loop to get faster convergence
 v = matrix(rnorm(K * n.sim), nrow = K, ncol = n.sim)
 # Z matrix needed for qmm function
 Z <- as.matrix(cereal[!is.na(i1_sugar),c("sugar", "mushy", "i1_sugar", "i1_mushy"), with = FALSE])
 # PZ matrix needed for GMM function
 PZ <- Z %*% solve(t(Z) %*% Z) %*% t(Z)
 # make weighting matrix. Also used in GMM function
 W.inv \leftarrow diag(1, 4, 4)
#======#
# ==== functions for search ====
#=======#
 # these are all the functions we will need for a given sd_mush sd_sug guess
 # hard code estimates for variance paremter to test funcitons
  # this is what the outside loop will be optimizing over
 sd_mush_test <- .1
 sd_sug_test <- .2
 # get inital delta guess
 #note doesn't need to come from cereal data.table, this is just an initial guess
 delta.initial <- cereal[, m_u]</pre>
 # this function takes the data.table, V sims, and the sd parameter estimates and returns
 # the mu (individual utility) estimates
 \#note I am hard coding this for mushy and sugar as the relavent variables.
 # could write this in a more general way if we wanted and have it take a list of theta parms
  # and a list of corresponding variabe names
 find_mu <- function(X, sd_sug.in, sd_mush.in, v.in){</pre>
    X \% \% diag(x = c(sd_sug.in, sd_mush.in), 2, 2) \% \% v.in
 }
```

```
# test it out
mu_mat <- find_mu(X, sd_sug_test, sd_mush_test, v )</pre>
# This computes a matrix of share estiamtes. rows are estimates for every product
# columns are estimates for every simulated V
ind_sh <- function(delta.in, mu.in, mkt.id.in){</pre>
  # This function computes the "individual" probabilities of choosing each brand
  numer <- exp(mu.in) * matrix(rep(exp(delta.in), n.sim), ncol = n.sim)</pre>
  denom <- as.matrix(do.call("rbind", lapply(mkt.id.in, function(tt){</pre>
    1 + colSums(numer[mkt.id.in %in% tt, ])
  })))
 return(numer / denom);
}
# test it out
sj_mat <- ind_sh(delta.initial, mu_mat, mkt.id)</pre>
blp_inner <- function(delta.in, mu.in, s.jm.in, mkt.id.in) {</pre>
  # Computes a single update of the BLP (1995) contraction mapping.
  # of market level predicted shares.
  # This single-update function is required by SQUAREM, see Varadhan and
  # Roland (SJS, 2008), and Roland and Varadhan (ANM, 2005)
  # INPUT
  # delta.in : current value of delta vector
  # mu.in: current mu matrix
  # OUTPUT
    delta.out : delta vector that equates observed with predicted market shares
  pred.s <- rowMeans(ind_sh(delta.in, mu.in, mkt.id.in));</pre>
  delta.out <- delta.in + log(s.jm.in) - log(pred.s)</pre>
 return(delta.out)
}
# test it out
delta_new <- blp_inner(delta.initial, mu_mat, s.jm, mkt.id)</pre>
# The function to iterate the contraction mapping is genereic. WE don't need to write it
# here is an example #note it is dependent on output of above examples
#note I have to pass additional variables so that we are not pulling objects from global enviorment
  squarem.output <- squarem(par</pre>
                                       = delta.initial,
                             fixptfn = blp_inner,
                                     = mu_mat,
                             mu.in
                             s.jm.in = s.jm,
                             mkt.id.in = mkt.id,
                             control = list(trace = TRUE))
  delta <- squarem.output$par</pre>
  summary(delta.initial - delta)
```

```
# function to runn GMM
 # a lot of inputs here but this is how you get around using global objects
 # This is supposed to be better practice but it doe sqet a bit wild with all these
 gmm_obj_f <- function(parm_vector.in, delta.in, X.in, XP.in, Z.in, PZ.in, W.inv.in, s.jm.in, mkt.
   sd_sug.in <- exp(parm_vector.in[[1]])</pre>
   sd_mush.in <- exp(parm_vector.in[[2]])</pre>
    # get new MU matrix
   mu <- find_mu(X.in, sd_sug.in = sd_sug.in, sd_mush.in = sd_mush.in, v.in)</pre>
   # update delta
   squarem.output <- squarem(par</pre>
                                       = delta.in,
                             fixptfn = blp_inner,
                             mu.in
                                       = mu,
                             s.jm.in = s.jm.in,
                             mkt.id.in = mkt.id.in,
                             control = list(trace = TRUE))
   delta_new <- squarem.output$par</pre>
   # first step
   PX.inv <- solve(t(XP.in) %*% PZ.in %*% XP.in)
   # finsih getting theta
   theta1 <- PX.inv %*% t(XP.in) %*% PZ.in %*% delta_new
   # get xi hat
   xi.hat <- delta_new - XP.in %*% theta1
   result <- t(xi.hat) %*% Z.in %*% W.inv.in %*% t(Z.in) %*% xi.hat
   # get function value
   return(result)
 }
 # make sd_vector
 parm_vector \leftarrow c(3, 3)
 # test it out
 f <- gmm_obj_f(parm_vector.in = parm_vector,</pre>
                delta.in = delta.initial,
                X.in
                           = X
                XP.in
                          = XP,
                Z.in
                            = Z,
                PZ.in
                           = PZ,
                W.inv.in = W.inv,
                s.jm.in = s.jm,
                mkt.id.in = mkt.id,
                v.in
                           = v)
Results <- optim(par
                              = c(3,3),
```

```
fn = gmm_obj_f,
                  # # uncomment if you want fast convergence
                   \# control = list(reltol = 5),
                   delta.in = delta.initial,
                   X.in = X,
                   XP.in
                             = XP,
                  Z.in = Z,
PZ.in = PZ,
                   W.inv.in = W.inv,
                   s.jm.in = s.jm,
                   mkt.id.in = mkt.id,
                   v.in
                             = v)
# save these real quick in case something happens
save(Results, file = paste0(f_out, "results_save.R"))
# grab out values
sd_final <- exp(Results$par)</pre>
sd_final
# get betas
# get new MU matrix
mu <- find_mu(X, sd_sug.in = sd_final[[1]], sd_mush.in = sd_final[[2]], v)</pre>
# update delta
squarem.output <- squarem(par = delta.initial,</pre>
                          fixptfn = blp_inner,
                          mu.in = mu,
                          s.jm.in = s.jm,
                          mkt.id.in = mkt.id,
                          control
                                   = list(trace = TRUE))
delta_final <- squarem.output$par</pre>
# first step
PX.inv <- solve(t(XP) %*% PZ %*% XP)</pre>
# finsih getting theta
theta_final <- PX.inv %*% t(XP) %*% PZ %*% delta_new
# qet m(parms) moment condition
temp <- delta_final - XP%*%theta_final</pre>
temp <- cbind(temp, temp, temp, temp)</pre>
m_final <- Z*(temp)</pre>
S_final <- (1/nrow(m_final))*t(m_final)%*%m_final
# get derivative of moment condition for Beta
temp <- cbind(X[,1],X[,1],X[,1],X[,1])
dmdb1 <- -Z*temp
```

```
temp <- cbind(X[,2],X[,2],X[,2],X[,2])
dmdb2 <- -Z*temp
# now for p
temp <- cbind(P,P,P,P)</pre>
dmda <- Z*temp
#----#
# ==== now get dmds ====
#======#
  #======#
  # ==== Get first one ====
  #----#
    # set phi
    phi <- .001
    # first get delta with + phi
    # get new MU matrix
    mu <- find_mu(X, sd_sug.in = sd_final[[1]] + phi, sd_mush.in = sd_final[[2]], v)</pre>
    # update delta
    squarem.output <- squarem(par = delta.initial,</pre>
                             fixptfn = blp_inner,
                             mu.in = mu,
                              s.jm.in = s.jm,
                             mkt.id.in = mkt.id,
                              control
                                      = list(trace = TRUE))
    delta_phi_p0 <- squarem.output$par</pre>
    temp <- delta_phi_p0 - XP%*%theta_final</pre>
    temp <- cbind(temp, temp, temp, temp)</pre>
    m_p0 \leftarrow Z*(temp)
    # do it again
    mu <- find_mu(X, sd_sug.in = sd_final[[1]] - phi, sd_mush.in = sd_final[[2]], v)</pre>
    # update delta
    squarem.output <- squarem(par = delta.initial,</pre>
                             fixptfn = blp_inner,
                             mu.in = mu,
                              s.jm.in = s.jm,
                             mkt.id.in = mkt.id,
                              control = list(trace = TRUE))
    delta_phi_m0 <- squarem.output$par</pre>
    temp <- delta_phi_m0 - XP%*%theta_final</pre>
    temp <- cbind(temp, temp, temp, temp)</pre>
    m_m0 \leftarrow Z*(temp)
    # now get first dmds
    dmds1 <- (m_p0 - m_m0)/(2*phi)
```

```
#----#
# ==== get second one ====
#----#
  mu <- find_mu(X, sd_sug.in = sd_final[[1]], sd_mush.in = sd_final[[2]] + phi, v)</pre>
  # update delta
  squarem.output <- squarem(par
                                    = delta.initial,
                           fixptfn = blp_inner,
                           mu.in
                                    = mu,
                            s.jm.in = s.jm,
                           mkt.id.in = mkt.id,
                            control = list(trace = TRUE))
  delta_phi_Op <- squarem.output$par</pre>
  temp <- delta_phi_Op - XP%*%theta_final</pre>
  temp <- cbind(temp, temp, temp, temp)</pre>
  m_0p \leftarrow Z*(temp)
  mu <- find_mu(X, sd_sug.in = sd_final[[1]], sd_mush.in = sd_final[[2]] - phi, v)</pre>
  # update delta
  squarem.output <- squarem(par = delta.initial,</pre>
                           fixptfn = blp_inner,
                           mu.in = mu,
                            s.jm.in = s.jm,
                           mkt.id.in = mkt.id,
                            control = list(trace = TRUE))
  delta_phi_0m <- squarem.output$par</pre>
  temp <- delta_phi_Om - XP%*%theta_final</pre>
  temp <- cbind(temp, temp, temp, temp)</pre>
  m_0m \leftarrow Z*(temp)
  # Now finish getting dmds
  dmds2 <- (m_0p - m_0m)/(2*phi)
#=======#
# ==== get standard errors ====
#======#
  isSymmetric(S_final)
  S_final
  m_final
  dmdb1
  dmdb2
  dmda
  dmds1
```

```
dmds2
      Rho <- t(rbind(colMeans(dmdb1),</pre>
                   colMeans(dmdb2),
                   colMeans(dmda),
                   colMeans(dmds1),
                   colMeans(dmds2)))
   SE_mat_final <- (1/JM)*(solve(t(Rho) %*% Rho, tol = 10^(-25)) %*% (t(Rho) %*% S_final %*% Rho) %*%
   SE_final <- sqrt(diag(SE_mat_final))</pre>
   \# put estiamates and SE into a table
   q3_p5 <- data.table(Variable = c("Sigma Sugar", "Sigma Mushy", "Beta Sugar", "Deta Mushy", "Alpha")
#======#
# ==== save output to latex ====
#----#
 if(opt_save){
   print(xtable(probit_res, type = "latex"),
         file = pasteO(f_out, "q1_p1.tex"),
         include.rownames = FALSE,
         floating = FALSE)
   print(xtable(logit_res, type = "latex"),
         file = paste0(f_out, "q1_p3.tex"),
         include.rownames = FALSE,
         floating = FALSE)
   print(xtable(p6_res_tab, type = "latex"),
         file = paste0(f_out, "q1_p6.tex"),
         include.rownames = FALSE,
         floating = FALSE)
   print(xtable(q2_p3_ols, type = "latex"),
         file = pasteO(f_out, "q2_p3a.tex"),
         include.rownames = FALSE,
         floating = FALSE)
   print(xtable(q2_p3_iv1, type = "latex"),
         file = pasteO(f_out, "q2_p3b.tex"),
         include.rownames = FALSE,
         floating = FALSE)
   print(xtable(q2_p3_iv2, type = "latex"),
         file = pasteO(f_out, "q2_p3c.tex"),
```