

pset 3 631

```
#####  
# ==== IO Pset 3 ====  
#####  
  
#####  
# ==== load packages/data ====  
#####  
  
rm(list = ls(pos = ".GlobalEnv"), pos = ".GlobalEnv")  
options(scipen = 999)  
cat("\f")  
  
library(package)  
library(data.table)  
library(xtable)  
library(boot)  
library(broom)  
library(rmarkdown)  
  
# set option for who is running this  
opt_nate <- TRUE  
  
# load data and set directories  
if(opt_nate){  
  # load data  
  gmdt <- fread("c:/Users/Nmath_000/Documents/MI_school/Third Year/Econ 631/ps3/GMdata.csv")  
  f_out <- "c:/Users/Nmath_000/Documents/Code/Econ_631/ps3/"  
  
  # if running on tyler's computer  
}else{  
  # load data from tylers locaiton  
  gmdt <- fread("C:/Users/tyler/Box/coursework/Econ_631/ps3/GMdata.csv")  
  f_out <- "C:/Users/tyler/Box/coursework/Econ_631/ps3/"  
  
}  
  
#####  
# ==== summary stats ====  
#####  
  
#####  
# ==== Make balanced panel ====
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#####

# make the balanced panel version of data
# first get number of years by firm
gmdt[, num_years := .N, index]

# check how many are in each group
gmdt[, .N, num_years]

# make balanced panel
gmdt_b <- gmdt[num_years == max(gmdt$num_years)]

# drop variable
gmdt[, num_years := NULL]
gmdt_b[, num_years := NULL]

#####
# ==== do comparison ====
#####

# make list of variables
vars <- grep("l", colnames(gmdt_b), value = TRUE)

# get each column of summart stats
sum_stats_li <- list()
sum_stats_li[[1]] <- data.table(Variable = vars)
# means
sum_stats_li[[2]] <- gmdt[, list("Full Mean" = lapply(.SD, mean)), .SDcols = vars]
sum_stats_li[[3]] <- gmdt_b[, list("Bal. Mean" = lapply(.SD, mean)), .SDcols = vars]

# t.test for mean equlity
# function to do it
tstat_fun <- function(var_i){
  res <- t.test(gmdt[, get(var_i)], gmdt_b[, get(var_i)])

  return(res$statistic)
}

# apply over variables
sum_stats_li[[4]] <- data.table( "Tstat" = unlist(lapply(vars, tstat_fun)))

# varianvce
sum_stats_li[[5]] <- gmdt[, list("Full VAR" = lapply(.SD, var)), .SDcols = vars]
sum_stats_li[[6]] <- gmdt_b[, list("Bal. VAR" = lapply(.SD, var)), .SDcols = vars]

# fstat for equalit of variance
ftest_fun <- function(var_i){
  res <- var.test(gmdt[, get(var_i)], gmdt_b[, get(var_i)])

  return(res$statistic)
}

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}

#apply over variables
sum_stats_li[[7]] <- data.table( "Fstat" = unlist(lapply(vars, ftest_fun)))

#min
sum_stats_li[[8]] <- gmdt[, list("Full Min" = lapply(.SD, min)), .SDcols = vars]
sum_stats_li[[9]] <- gmdt_b[, list("Bal. Min" = lapply(.SD, min)), .SDcols = vars]
#max
sum_stats_li[[10]] <- gmdt[, list("Full max" = lapply(.SD, max)), .SDcols = vars]
sum_stats_li[[11]] <- gmdt_b[, list("Bal. max" = lapply(.SD, max)), .SDcols = vars]

# get variance
sum_stats <- do.call(cbind, sum_stats_li)
#check it out!
sum_stats

#=====#
# ==== save table to tex ====
#=====#

# split this into two tables
sum_stat_1 <- sum_stats[, 1:7, with=FALSE]
sum_stat_2 <- sum_stats[, c(1, 8:11), with = FALSE]

# save summary stats
print(xtable(sum_stat_1, type = "latex"),
      file = paste0(f_out, "sum_stats_1.tex"),
      include.rownames = FALSE,
      floating = FALSE)

print(xtable(sum_stat_2, type = "latex"),
      file = paste0(f_out, "sum_stats_2.tex"),
      include.rownames = FALSE,
      floating = FALSE)

#=====#
# ==== Estimate Production Function ==== #
#=====#

# Step 1: Get measurement error from second-order polynomial
# (including interactions) of emp, dnpt, drst and investment
setnames(gmdt, c("ldsal", "ldnpt", "ldrst"), c("lsales", "lcap", "lrdcap"))

gmdt[, dummy:=1]

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# do it for balanced too just for fun
setnames(gmdt_b, c("ldsai", "ldnpt", "ldrst"), c("lsales", "lcap", "lrdcap"))

gmdt_b[,dummy:=1]

#####
# === internal functions ===
#####

# in_data <- gmdt_b
get_resid_fun <- function(in_data){

  dt_copy <- copy(in_data)

  X <- as.matrix(dt_copy[,.(dummy, lemp,lcap,lrdcap,ldinv, lemp*lemp,lemp*lcap, lemp*lrdcap,
    lemp*ldinv,lcap*lcap,lcap*lrdcap,lcap*ldinv,lrdcap*lrdcap,lrdcap*ldinv,ld

  Y <- as.matrix(dt_copy[,.(lsales)])

  # Get the coefficients of all these interactions
  beta1 <- solve(t(X)%*%X)%*%t(X)%*%Y

  # Get the fitted values
  theta <- X%*%beta1

  # Add the fitted values to the datatable
  dt_copy[,theta:= theta]

  # get lags of emp, cap, rdcap and theta, remove observations without lag values
  cols = c("lemp","lcap","lrdcap", "theta")
  anscols = paste("lag", cols, sep="_")
  dt_copy[, (anscols) := shift(.SD, 1, NA, "lag"), .SDcols=cols, index]

  dt_copy <- dt_copy[!is.na(lag_lemp),]

  return(dt_copy)

}

# # ttest the funciton
# gmdt2 <- get_resid_fun(gmdt)
#

# function to run 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 sget a bit wild with all these
gmm_obj_f <- function(parm_vector.in,
  Y.in = Y,
  X.in = X,
  lX.in = lX,
  ltheta.in = ltheta,

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        Z.in = Z,
        W.in = W){

beta <- as.matrix(parm_vector.in[1:4])
rho <- parm_vector.in[5]

current.resid <- Y.in - X.in**beta

lag.w <- as.matrix(ltheta.in) - lX.in**beta

m <- current.resid - rho*lag.w

m <- as.matrix(m)

distance <- t(Z.in)**m

result <- t(distance/length(m))** W.in ** (distance/length(m))

# get function value
return(result)
}

# TEST
# # parms to test the function
# parm_vector <- c(1, .6, .2, .2, .8)
#
# # Get X's and lag X's and Zs for GMM estimation
# X <- as.matrix(gmdt2[,.(dummy, lemp, lcap, lrdcap)])
# lX <- as.matrix(gmdt2[,.(dummy, lag_lemp, lag_lcap, lag_lrdcap)])
# Z <- as.matrix(gmdt2[,.(dummy, lag_lemp, lcap, lrdcap)])
# Y <- as.matrix(gmdt2[,.(lsales)])
# ltheta <- as.matrix(gmdt2[,.(lag_theta)])
# W <- diag(1, 4, 4)
#
# # test it out
# f <- gmm_obj_f(parm_vector.in = parm_vector,
#               Y.in           = Y,
#               X.in           = X,
#               lX.in          = lX,
#               ltheta.in      = ltheta,
#               Z.in           = Z,
#               W.in           = W)
#
# # Run the initial GMM using the identity matrix as the weighting matrix
# Results.step1 <- optim(par      = parm_vector,
#                       fn        = gmm_obj_f,
#                       Y.in      = Y,
#                       X.in      = X,
#                       lX.in     = lX,
#                       ltheta.in = ltheta,
#                       Z.in      = Z,
#                       W.in      = W)

# Function to calculate optimal weighting matrix

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find.optimal.W <- function(results.in,
                           Y.in = Y,
                           X.in = X,
                           lX.in = lX,
                           ltheta.in = ltheta,
                           Z.in = Z){
  beta <- as.matrix(results.in$par[1:4])
  rho <- results.in$par[5]

  current.resid <- Y.in - X.in%%beta

  lag.w <- as.matrix(ltheta.in) - lX.in%%beta

  m <- current.resid - rho*lag.w

  m <- as.matrix(m)

  distance <- t(Z.in*cbind(m, m, m, m))%(Z.in*cbind(m, m, m, m))
  W <- distance/length(m)

  W.inv <- solve(W)
  return(W.inv)
}

# test
# # Get optimal weighting matrix
# W.opt <- find.optimal.W(results.in = Results.step1,
#                           Y.in = Y,
#                           X.in = X,
#                           lX.in = lX,
#                           ltheta.in = ltheta,
#                           Z.in = Z)
#
# # Run again with optimal weighting matrix
# Results.final <- optim(par      = parm_vector,
#                         fn      = gmm_obj_f,
#                         Y.in = Y,
#                         X.in = X,
#                         lX.in = lX,
#                         ltheta.in = ltheta,
#                         Z.in = Z,
#                         W.in = W.opt)

#####
# ==== boot function ====
#####

# in_data <- gmdt
# sample <- 1:nrow(in_data)
to_boot_fun <- function(in_data, sample){

  boot_dt <- in_data[sample]

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# get resids
boot_dt <- get_resid_fun(boot_dt)

parm_vector <- c(1, .6, .2, .2, .8)

# Get X's and lag X's and Zs for GMM estimation
X <- as.matrix(boot_dt[,.(dummy, lemp, lcap, lrdcap)])
lX <- as.matrix(boot_dt[,.(dummy, lag_lemp, lag_lcap, lag_lrdcap)])
Z <- as.matrix(boot_dt[,.(dummy, lag_lemp, lcap, lrdcap)])
Y <- as.matrix(boot_dt[,.(lsales)])
ltheta <- as.matrix(boot_dt[,.(lag_theta)])
W <- diag(1, 4, 4)

# test it out
f <- gmm_obj_f(parm_vector.in = parm_vector,
               Y.in           = Y,
               X.in           = X,
               lX.in          = lX,
               ltheta.in      = ltheta,
               Z.in           = Z,
               W.in           = W)

# Run the initial GMM using the identity matrix as the weighting matrix
Results.step1 <- optim(par      = parm_vector,
                      fn       = gmm_obj_f,
                      Y.in     = Y,
                      X.in     = X,
                      lX.in     = lX,
                      ltheta.in = ltheta,
                      Z.in     = Z,
                      W.in     = W)

# Get optimal weighting matrix
W.opt <- find.optimal.W(results.in = Results.step1,
                        Y.in = Y,
                        X.in = X,
                        lX.in = lX,
                        ltheta.in = ltheta,
                        Z.in = Z)

# Run again with optimal weighting matrix
Results.final <- optim(par      = parm_vector,
                      fn       = gmm_obj_f,
                      Y.in     = Y,
                      X.in     = X,
                      lX.in     = lX,
                      ltheta.in = ltheta,
                      Z.in     = Z,
                      W.in     = W.opt)

return(Results.final$par)
}

```

```

# to_boot_fun(gmdt, 1:(nrow(gmdt)))

#####
# ==== Do bootstrapping ====
#####

#####
# ==== write out own boot ====
#####

n_sim <- 1000

# block boot function to iterate
block_boot <- function(iteration, in_data){

  # if iteration is 1 use full sample
  if(iteration == 1){

    sample_i <- 1:1400
  }else{
    sample_i <- sample(1:1400, 1400, replace = TRUE)
  }

  # grab a random sample
  dt_i <- in_data[index %in% sample_i]

  # run function on subsample
  res_i <- to_boot_fun(dt_i, 1:nrow(dt_i))

  return(data.table(var = parms , res = res_i))
}

# apply function n_sim times
re_list_b <- lapply(1:n_sim, block_boot, in_data = gmdt_b)

# do it again on unbalanced sample
re_list_u <- lapply(1:n_sim, block_boot, in_data = gmdt)

# bind the lists together
re_dt_b <- rbindlist(re_list_b)
re_dt_u <- rbindlist(re_list_u)

# Now get standard errors for balanced
re_dt_b[, mean := mean(res), var]
re_dt_b[, to_sum := (res-mean)^2 ]
out_dt_b <- re_dt_b[, list(std.error = sqrt(1/(n_sim - 1) * sum(to_sum))), var]

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# Now get standard errors fo unbalanced
re_dt_u[, mean := mean(res), var]
re_dt_u[, to_sum := (res-mean)^2 ]
out_dt_u <- re_dt_u[, list(std.error = sqrt(1/(n_sim - 1) * sum(to_sum))), var]

# Now add in estiamtes
est_b <- re_list_b[[1]]
est_u <- re_list_u[[1]]
out_dt_b <- merge(est_b, out_dt_b, "var")
out_dt_u <- merge(est_u, out_dt_u, "var")

# change the variable names
setnames(out_dt_b, colnames(out_dt_b), c("Parm", "Statistic", "Std.Error"))
setnames(out_dt_u, colnames(out_dt_u), c("Parm", "Statistic", "Std.Error"))

# save table
print(xtable(out_dt_b, type = "latex"),
      file = paste0(f_out, "boot_res.tex"),
      include.rownames = FALSE,
      floating = FALSE)

print(xtable(out_dt_u, type = "latex"),
      file = paste0(f_out, "boot_res_unbalanced.tex"),
      include.rownames = FALSE,
      floating = FALSE)

#=====#
# ==== run r markdown for tex file ====
#=====#

# load data and set directories
if(opt_nate){
  rmarkdown::render(input = "C:/Users/Nmath_000/Documents/Code/Econ_631/ps3/ps3_r_markdown.Rmd",
                    output_format = "pdf_document",
                    output_file = paste0(f_out, "assignment_3_r_code_pdf.pdf"))

  # if running on tyler's computer
}else{
  rmarkdown::render(input = "C:/Users/tyler/Box/coursework/Econ_631/ps3/ps3_r_markdown.Rmd",
                    output_format = "pdf_document",
                    output_file = paste0(f_out, "assignment_3_r_code_pdf.pdf"))
}

```