

Example 4-2

September 12, 2020

```
[5]: # install the following packages and library
install.packages("plm")
install.packages("splm")

library("plm")
```

package 'splm' successfully unpacked and MD5 sums checked

The downloaded binary packages are in

C:\Users\cwell\AppData\Local\Temp\RtmpqqV5GV\downloaded_packages

```
[9]: ##-----Block 1-----

#### Example 4-2 ####

## -----
data("RiceFarms", package = "splm")
Rice <- pdata.frame(RiceFarms, index = "id")
rice.w <- plm(log(goutput) ~ log(seed) + log(totlabor) + log(size), Rice)
rice.r <- update(rice.w, model = "random")

# phtest() is the Hausman test.
# arguments are either the within and GLS model or the formula and data
ptest(rice.w, rice.r)
```

Hausman Test

```
data: log(goutput) ~ log(seed) + log(totlabor) + log(size)
chisq = 3.775, df = 3, p-value = 0.2868
alternative hypothesis: one model is inconsistent
```

```
[11]: ##-----Block 2-----

# Mundlak (1978) approach. Takes the difference between the
# within and between estimators
rice.b <- update(rice.w, model = "between")
```

```

cp <- intersect(names(coef(rice.b)), names(coef(rice.w)))
dcoef <- coef(rice.w)[cp] - coef(rice.b)[cp]
V <- vcov(rice.w)[cp, cp] + vcov(rice.b)[cp, cp]
as.numeric(t(dcoef) %*% solve(V) %*% dcoef)

## -----

# correlation coefficient between the individual effects and
# individual means of the explanatory variable
cor(fixef(rice.w), between(log(Rice$goutput)))

```

3.77284055832556

0.448045399935444

```

[12]: ##-----Block 3-----

data("RiceFarms", package = "splm")
pdim(RiceFarms, index = "id")

# piest() is the Chamberlain test. Input argument is formula and the data
piest(log(goutput) ~ log(seed) + log(totlabor) + log(size),
      RiceFarms, index = "id")

```

Balanced Panel: n = 171, T = 6, N = 1026

Chamberlain's pi test

data: log(goutput) ~ log(seed) + log(totlabor) + log(size)
 chisq = 113.72, df = 87, p-value = 0.02882
 alternative hypothesis: within specification does not apply

```

[13]: ##-----Block 4-----

# Angrist and Newey (1991) test. Input argument is formula and the data
aneweytest(log(goutput) ~ log(seed) + log(totlabor) + log(size),
           RiceFarms, index = "id")

```

Angrist and Newey's test of within model

data: log(goutput) ~ log(seed) + log(totlabor) + log(size)
 chisq = 141.89, df = 87, p-value = 0.0001851
 alternative hypothesis: within specification does not apply