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Econometric convergence test and club clustering using Stata

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Abstract. In this article, I introduce a new package with five commands to perform econometric convergence analysis and club clustering as proposed by Phillips and Sul (2007, *Econometrica* 75: 1771–1855). The `logtreg` command performs the log t regression test. The `psecta` command implements the clustering algorithm to identify convergence clubs. The `scheckmerge` command conducts the log t regression test for all pairs of adjacent clubs. The `imergeclub` command tries to iteratively merge adjacent clubs. The `pfilter` command extracts the trend and cyclical components of a time series of each individual in panel data. I provide an example from Phillips and Sul (2009, *Journal of Applied Econometrics* 24: 1153–1185) to illustrate the use of these commands. Additionally, I use Monte Carlo simulations to exemplify the effectiveness of the clustering algorithm.

Keywords: st0503, `logtreg`, `psecta`, `scheckmerge`, `imergeclub`, `pfilter`, convergence, club clustering, log t test

1 Introduction

Convergence in economics refers to the hypothesis that all economies would eventually converge in terms of per-capita output. This issue has played a central role in the empirical growth literature (Pesaran 2007). A large body of literature (for example, Baumol [1986]; Bernard and Durlauf [1995]; Barro and Sala-I-Martin [1997]; Lee, Pesaran, and Smith [1997]; Luginbuhl and Koopman [2004]) has contributed to developing methods for convergence tests and empirically investigating the convergence hypothesis across different countries and regions. In the past years, convergence analysis has also been applied in other topics such as cost of living (Phillips and Sul 2007), carbon dioxide emissions (Panopoulou and Pantelidis 2009), ecoefficiency (Camarero et al. 2013), house prices (Montañés and Olmos 2013), corporate tax (Regis, Cuestas, and Chen 2015), etc.

Phillips and Sul (2007) proposed a novel approach (termed “log t ” regression test)¹ to test the convergence hypothesis based on a nonlinear time-varying factor model. The proposed approach has the following merits: First, it accommodates heterogeneous agent behavior and evolution in that behavior. Second, the proposed test does not impose any particular assumptions concerning trend stationarity or stochastic nonstation-

1. It is also called “log t test” for short.

arity, thereby being robust to the stationarity property of the series. [Phillips and Sul \(2009\)](#) showed that the traditional convergence tests for economic growth have some pitfalls. For instance, estimation of augmented Solow regression under transitional heterogeneity is biased and inconsistent because of the issues of omitted variables and endogeneity. Conventional cointegration tests typically have low power to detect the asymptotic comovement, because the existence of a unit root in the differential of the series does not necessarily lead to the divergence conclusion.

Another common concern with the convergence analysis is the possible existence of convergence clubs. Regarding this issue, traditional studies typically divided all individuals into subgroups based on some prior information (for example, geographical location, institution), then tested the convergence hypothesis for each subgroup respectively. [Phillips and Sul \(2007\)](#) constructed a new algorithm to identify clusters of convergence subgroups. The developed algorithm is a data-driven method that avoids ex-ante sample separation. The relative transition parameter mechanism that [Phillips and Sul \(2007\)](#) proposed to characterize individual variations fits in with some common models.² It can be used as a general panel method to cluster individuals into groups with similar transition paths.

In practice, [Phillips and Sul \(2007, 2009\)](#) provided Gauss codes to perform their empirical studies. Recently, [Schnurbus, Haupt, and Meier \(2017\)](#) provided a set of R functions to replicate the key results of [Phillips and Sul \(2009\)](#). In this article, I introduce a new package, `psecta`, to perform the econometric convergence test and club clustering developed by [Phillips and Sul \(2007\)](#).

The remainder of this article is organized as follows: section 2 briefly describes the methodology of [Phillips and Sul \(2007\)](#); sections 3–7 explain the syntax and options of the new commands; section 8 performs Monte Carlo simulations to examine the effectiveness of the clustering algorithm; and section 9 illustrates the use of the commands with an example from [Phillips and Sul \(2009\)](#).

2 Econometric convergence test and club clustering

2.1 Time-varying factor representation

The starting point of the model is decomposing the panel data, X_{it} , as

$$X_{it} = g_{it} + a_{it} \quad (1)$$

where g_{it} represents systematic components such as permanent common components and a_{it} embodies transitory components. To separate common components from idiosyncratic components, we further transform (1) as

2. See, for example, economic growth with heterogeneous technological progress ([Parente and Prescott 1994](#); [Howitt and Mayer-Foulkes 2005](#)), income processes with heterogeneity ([Baker 1997](#); [Moffitt and Gottschalk 2002](#)), and stock price factor models ([Menzly, Santos, and Veronesi 2002](#); [Ludvigson and Ng 2007](#)).

$$X_{it} = \left(\frac{g_{it} + a_{it}}{u_t} \right) u_t = \delta_{it} u_t \quad (2)$$

where δ_{it} is a time-varying idiosyncratic element and u_t is a single common component. Equation (2) is a dynamic-factor model where u_t captures some deterministic or stochastically trending behavior, and the time-varying factor-loading coefficient δ_{it} measures the idiosyncratic distance between X_{it} and the common trend component u_t .

In general, we cannot directly fit the model without imposing some restrictions on δ_{it} and u_t . Thus, [Phillips and Sul \(2007\)](#) proposed removing the common factor as follows:

$$h_{it} = \frac{X_{it}}{\frac{1}{N} \sum_{i=1}^N X_{it}} = \frac{\delta_{it}}{\frac{1}{N} \sum_{i=1}^N \delta_{it}} \quad (3)$$

h_{it} is the relative transition parameter, which measures the loading coefficient relative to the panel average at time t . In other words, h_{it} traces out a transition path of individual i in relation to the panel average. Equation (3) indicates that the cross-sectional mean of h_{it} is unity, and the cross-sectional variance of h_{it} satisfies the following condition:

$$H_{it} = \frac{1}{N} \sum_{i=1}^N (h_{it} - 1)^2 \rightarrow 0 \text{ if } \lim_{t \rightarrow \infty} \delta_{it} = \delta, \text{ for all } i$$

2.2 The log t regression test

The convergence of X_{it} requires the following condition:

$$\lim_{t \rightarrow \infty} \frac{X_{it}}{X_{jt}} = 1, \text{ for all } i \text{ and } j$$

[Phillips and Sul \(2007\)](#) defined this condition as the relative convergence. It is equivalent to the convergence of the time-varying factor-loading coefficient

$$\lim_{t \rightarrow \infty} \delta_{it} = \delta, \text{ for all } i$$

Assume the loading coefficient δ_{it} as

$$\delta_{it} = \delta_i + \sigma_{it} \xi_{it}, \quad \sigma_{it} = \frac{\sigma_i}{L(t)t^\alpha}, t \geq 1, \sigma_i > 0 \text{ for all } i$$

where $L(t)$ is a slowly varying function. Possible choices for $L(t)$ can be $\log(t)$, $\log^2(t)$, or $\log\{\log(t)\}$. The Monte Carlo simulations in [Phillips and Sul \(2007\)](#) indicate that $L(t) = \log(t)$ produces the least size distortion and the best test power. Thus, we set $L(t) = \log(t)$ in our Stata codes.

[Phillips and Sul \(2007\)](#) developed a regression t test for the null hypothesis of convergence,

$$\mathcal{H}_0: \delta_i = \delta \text{ and } \alpha \geq 0$$

against the alternative, $\mathcal{H}_A: \delta_i \neq \delta$ or $\alpha < 0$. Specifically, the hypothesis test can be implemented through the following log t regression model:

$$\log\left(\frac{H_1}{H_t}\right) - 2\log\{\log(t)\} = a + b\log(t) + \varepsilon_t$$

for $t = [rT], [rT] + 1, \dots, T$ with $r > 0$

The selection of the initiating sample fraction r might influence the results of the above regression. The Monte Carlo experiments indicate that $r \in [0.2, 0.3]$ achieves a satisfactory performance. Specifically, it is suggested to set $r = 0.3$ for the small or moderate $T (\leq 50)$ sample and set $r = 0.2$ for the large $T (\geq 100)$ sample.

[Phillips and Sul \(2007\)](#) further showed that $b = 2\alpha$ and \mathcal{H}_0 is conveniently tested through the weak inequality null $\alpha \geq 0$. It implies a one-sided t test. Under some technical assumptions, the limit distribution of the regression t statistic is

$$t_b = \frac{\hat{b} - b}{s_b} \Rightarrow N(0, 1)$$

where

$$s_b^2 = \widehat{lvar}(\hat{\varepsilon}_t) \left\{ \sum_{t=[rT]}^T \left(\log(t) - \frac{1}{T - [rT] + 1} \sum_{t=[rT]}^T \log(t) \right)^2 \right\}^{-1}$$

and $\widehat{lvar}(\hat{\varepsilon}_t)$ is a conventional heteroskedastic and autocorrelated estimate formed from the regression residuals.

2.3 Club convergence test and clustering

Rejection of the null hypothesis of convergence for the whole panel cannot rule out the existence of convergence in subgroups of the panel. To investigate the possibility of convergence clusters, [Phillips and Sul \(2007\)](#) developed a data-driven algorithm. [Schnurbus, Haupt, and Meier \(2017\)](#) advocated making some minor adjustments to the original algorithm. We sketch their ideas as follows:

1. Cross-section sorting

Sort individuals in the panel decreasingly according to their observations in the last period. If there is substantial time-series volatility in the data, the sorting can be implemented based on the time-series average of the last fraction (for example, $1/2, 1/3$) of the sample. Index individuals with their orders $\{1, \dots, N\}$.

2. Core group formation

- 2.1 Find the first k such that the test statistic of the log t regression $t_k > -1.65$ for the subgroup with individuals $\{k, k + 1\}$. If there is no k satisfying $t_k > -1.65$, exit the algorithm, and conclude that there are no convergence subgroups in the panel.
- 2.2 Start with the k identified in step 2.1, perform log t regression for the subgroups with individuals $\{k, k + 1, \dots, k + j\}, j \in \{1, \dots, N - k\}$. Choose j^* such that the subgroup with individuals $\{k, k + 1, \dots, k + j^*\}$ yields the highest value of the test statistic. Individuals $\{k, k + 1, \dots, k + j^*\}$ form a core group.

3. Sieve individuals for club membership

- 3.1 Form a complementary group $G_{j^*}^c$ with all the remaining individuals not included in the core group. Add one individual from $G_{j^*}^c$ at each time to the core group and run the log t test. Include the individual in the club candidate group if the test statistic is greater than the critical value c^* .³
- 3.2 Run the log t test for the club candidate group identified in step 3.1. If the test statistic \hat{t}_b is greater than -1.65 , the initial convergence club is obtained. If not, [Phillips and Sul \(2007\)](#) advocated raising the critical value c^* and repeating steps 3.1 and 3.2 until $\hat{t}_b > -1.65$. [Schnurbus, Haupt, and Meier \(2017\)](#) proposed adjusting this step as follows: If the convergence hypothesis does not hold for the club candidate group, sort the club candidates w.r.t. decreasing \hat{t}_b obtained in step 3.1. If there are some $\hat{t}_b > -1.65$, add the individual with the highest value of \hat{t}_b to form an extended core group. Add one individual from the remaining candidates at a time, run the log t test, and denote the test statistic \hat{t}_b . If the highest value of \hat{t}_b is not greater than -1.65 , stop the procedure; the extended core group will form an initial convergence club. Otherwise, repeat the above procedure to add the individual with the highest \hat{t}_b .

4. Recursion and stopping rule

Form a subgroup of the remaining individuals that are not sieved by step 3. Perform the log t test for this subgroup. If the test statistic is greater than -1.65 , the subgroup forms another convergence club. Otherwise, repeat steps 1–3 on this subgroup.

5. Club merging

Perform the log t test for all pairs of the subsequent initial clubs. Merge those clubs fulfilling the convergence hypothesis jointly. [Schnurbus, Haupt, and Meier \(2017\)](#) advocated conducting club merging iteratively as follows: run the log t test for the initial clubs 1 and 2; if they fulfill the convergence hypothesis jointly,

3. When T is small, the sieve criterion c^* can be set to 0 to ensure that it is highly conservative, whereas for large T , c^* can be set to the asymptotic 5% critical value -1.65 .

merge them to form the new club 1, then run the $\log t$ test for the new club 1 and the initial club 3 jointly; if not, run the $\log t$ test for initial clubs 2 and 3, etc. The new club classifications would be obtained by the above procedure. After that, one can also repeat the procedure on the newly obtained club classifications until no clubs can be merged, which leads to the classifications with the smallest number of convergence clubs.

3 The **logtreg** command

logtreg performs the $\log t$ test using linear regression with heteroskedasticity- and autocorrelation-consistent standard errors.

3.1 Syntax

logtreg *varname* [*if*] [*in*] [, **kq(#)** **nomata**]

3.2 Options

kq(#) specifies the first **kq()** proportion of the data to be discarded before regression. The default is **kq(0.3)**.

nomata bypasses the use of community-contributed Mata functions; by default, community-contributed Mata functions are used.

3.3 Stored results

logtreg stores the following in **e()**:

| | |
|-------------------|--|
| Scalars | |
| e(N) | number of individuals |
| e(T) | number of time periods |
| e(nreg) | number of observations used for the regression |
| e(beta) | $\log t$ coefficient |
| e(tstat) | t statistic for $\log t$ |
| Macros | |
| e(cmd) | logtreg |
| e(cmdline) | command as typed |
| e(varlist) | name of the variable for $\log t$ test |
| Matrices | |
| e(res) | table of estimation results |

4 The psecta command

`psecta` implements club convergence and clustering analysis using the algorithm proposed by [Phillips and Sul \(2007\)](#).

4.1 Syntax

```
psecta varname [ , name(panelvar) kq(#) gen(newvar) cr(#) incr(#)
    maxcr(#) adjust fr(#) nomata noprtlogtreg ]
```

4.2 Options

`name(panelvar)` specifies a panel variable to be displayed for the clustering results; by default, the panel variable specified by `xtset` is used.

`kq(#)` specifies the first `kq()` proportion of the data to be discarded before regression. The default is `kq(0.3)`.

`gen(newvar)` creates a new variable to store club classifications. For the individuals that are not classified into any convergence club, missing values are generated.

`cr(#)` specifies the critical value for club clustering. The default is `cr(0)`.

`incr(#)` specifies the increment of `cr()` when the initial `cr()` value fails to sieve individuals for clusters. The default is `incr(0.05)`.

`maxcr(#)` specifies the maximum of `cr()` value. The default is `maxcr(50)`.

`adjust` specifies using the adjusted method proposed by [Schnurbus, Haupt, and Meier \(2017\)](#) instead of raising `cr()` when the initial `cr()` value fails to sieve individuals for clusters. See [Schnurbus, Haupt, and Meier \(2017\)](#) for more details.

`fr(#)` specifies sorting individuals by the time-series average of the last `fr()` proportion periods. The default is `fr(0)`, sorting individuals according to the last period.

`nomata` bypasses the use of community-contributed Mata functions; by default, community-contributed Mata functions are used.

`noprtlogtreg` suppresses the estimation results of the `logtreg`.

4.3 Stored results

`psecta` stores the following in `e()`:

| | |
|-------------------------|--|
| Scalar | |
| <code>e(nclub)</code> | number of convergent clubs |
| Macros | |
| <code>e(cmd)</code> | <code>psecta</code> |
| <code>e(cmdline)</code> | command as typed |
| <code>e(varlist)</code> | name of the variable for log <i>t</i> test |
| Matrices | |
| <code>e(bm)</code> | log <i>t</i> coefficients |
| <code>e(tm)</code> | <i>t</i> statistics |
| <code>e(club)</code> | club classifications |

4.4 Dependency of `psecta`

`psecta` depends on the Mata function `mm_which()`. If not already installed, you can install it by typing `ssc install moremata`.

5 The `scheckmerge` command

`scheckmerge` performs the log *t* test for all pairs of adjacent clubs.

5.1 Syntax

`scheckmerge varname, club(varname) kq(#) [mdiv nomata]`

5.2 Options

`club(varname)` specifies the initial club classifications. `club()` is required.

`kq(#)` specifies the first `kq()` proportion of the data to be discarded before regression. `kq()` is required.

`mdiv` specifies including the divergence group for the log *t* test; by default, the divergence group is excluded.

`nomata` bypasses the use of community-contributed Mata functions; by default, community-contributed Mata functions are used.

5.3 Stored results

`scheckmerge` stores the following in `e()`:

| | |
|-------------------------|--|
| Macros | |
| <code>e(cmd)</code> | <code>scheckmerge</code> |
| <code>e(cmdline)</code> | command as typed |
| <code>e(varlist)</code> | name of the variable for log <i>t</i> test |
| Matrices | |
| <code>e(bm)</code> | log <i>t</i> coefficients |
| <code>e(tm)</code> | <i>t</i> statistics |

6 The `imergeclub` command

`imergeclub` iteratively conducts merging adjacent clubs.

6.1 Syntax

```
imergeclub varname, club(varname) kq(#) [name(panelvar) gen(newvar)  
imore mdiv nomata noprtlogtreg]
```

6.2 Options

`club(varname)` specifies the initial club classifications. `club()` is required.

`kq(#)` specifies the first `kq()` proportion of the data to be discarded before regression. `kq()` is required.

`name(panelvar)` specifies a panel variable to be displayed for the clustering results; by default, the panel variable specified by `xtset` is used.

`gen(newvar)` creates a new variable to store the new club classifications. For the individuals that are not classified into any convergence club, missing values are generated.

`imore` specifies merging clubs iteratively until no clubs can be merged. By default, the procedure is conducted as follows: First, run the log *t* test for the individuals belonging to the initial clubs 1 and 2 (obtained from club clustering). Second, if they fulfill the convergence hypothesis jointly, merge them to be the new club 1, and then run the log *t* test for the new club 1 and the initial club 3; if not, run the log *t* test for initial clubs 2 and 3, etc. If `imore` is chosen, the above procedure is repeated until no clubs can be merged.

`mdiv` specifies including the divergence group during club merging; by default, the divergence group is excluded.

`nomata` bypasses the use of community-contributed Mata functions; by default, community-contributed Mata functions are used.

`noprtlogtreg` suppresses the estimation results of the `logtreg`.

6.3 Stored results

`imergeclub` stores the following in `e()`:

| | |
|-------------------------|--|
| Scalars | |
| <code>e(nclub)</code> | number of convergent clubs |
| Macros | |
| <code>e(cmd)</code> | <code>imergeclub</code> |
| <code>e(cmdline)</code> | command as typed |
| <code>e(varlist)</code> | name of the variable for log <i>t</i> test |
| Matrices | |
| <code>e(bm)</code> | log <i>t</i> coefficients |
| <code>e(tm)</code> | <i>t</i> statistics |

7 The pfilter command

`pfilter` applies the `tsfilter` (see [TS] `tsfilter`) command into a panel-data context. It can be used to extract trend and cyclical components for each individual in the panel, respectively.

7.1 Syntax

`pfilter varname, method(string) [trend(newvar) cyc(newvar) options]`

7.2 Options

`method(string)` specifies the filter method. `string` should be chosen from `bk`, `bw`, `cf`, or `hp`. `method()` is required.

`trend(newvar)` creates a new variable for the trend component.

`cyc(newvar)` creates a new variable for the cyclical component.

`options` are any options available for `tsfilter` (see [TS] `tsfilter`).

7.3 Stored results

`pfilter` stores the following in `r()`:

| | |
|-------------------------|----------------------|
| Macros | |
| <code>r(cmd)</code> | <code>pfilter</code> |
| <code>r(cmdline)</code> | command as typed |
| <code>r(varlist)</code> | name of the variable |

8 Monte Carlo simulation

Phillips and Sul (2007) performed extensive simulations for the power and size of the “log t ” test. They also showed how the clustering procedure works through Monte Carlo experiments. In this section, we further do simulation exercises to exemplify the effectiveness of the clustering algorithm in the finite sample.

Referring to Phillips and Sul (2007), we consider the following data-generating process,

$$\begin{aligned} X_{it} &= \theta_{it}\mu_t, \quad \theta_{it} = \theta_i + \theta_{it}^0 \\ \theta_{it}^0 &= \rho_i\theta_{it-1}^0 + \varepsilon_{it} \end{aligned}$$

where $t = 1, \dots, T$; $\varepsilon_{it} \sim iid N\{0, \sigma_i^2 \log(t+1)^{-2} t^{-2\alpha_i}\}$, $\sigma_i \sim U[0.02, 0.28]$, $\alpha_i \sim U[0.2, 0.9]$; $\rho_i \sim U[0, 0.4]$. Note that the common component μ_t cancels out in the test procedure. It is not needed to generate the realizations of μ_t .

For simulations, we set $T = 20, 40, 60, 100$ and $N = 40, 80, 120$. The number of replications is 1,000. We first consider the following two cases:

Case 1: One convergence club and one divergence subgroup. We consider two equal-sized groups in the panel with numbers $N_1 = N_2 = N/2$. We set $\theta_i = 1$ and $\theta_i \sim U[1.5, 5]$ for the first and second groups, respectively. This implies that the first group forms a convergence club and the second group is divergent.

Case 2: Two convergence clubs. Two groups are set much like case 1, except that $\theta_i = 1$ and $\theta_i = 2$ for the first and second groups, respectively.

Tables 1 and 2 report the false discovery rate (the ratio of the replications that fail to identify the club memberships) for case 1 and case 2, respectively. Generally speaking, the false discovery rate is acceptable. In case 1, the false discovery rate is lower than 10% when $T = 20$, and it becomes lower than 5% when $T \geq 40$. In case 2, the false discovery rate is lower than 5% for all combinations of N and T except for $(N = 120, T = 40)$.

Table 1. Simulation result of case 1

| $T \backslash N$ | 40 | 80 | 120 |
|------------------|-------|-------|-------|
| 20 | 0.097 | 0.070 | 0.082 |
| 40 | 0.048 | 0.035 | 0.047 |
| 60 | 0.023 | 0.032 | 0.038 |
| 100 | 0.027 | 0.033 | 0.039 |

Table 2. Simulation result of case 2

| $T \backslash N$ | 40 | 80 | 120 |
|------------------|-------|-------|-------|
| 20 | 0.026 | 0.023 | 0.046 |
| 40 | 0.014 | 0.031 | 0.056 |
| 60 | 0.012 | 0.022 | 0.040 |
| 100 | 0.015 | 0.015 | 0.042 |

For the experiments of case 1 and case 2, the values of θ_i are given. Here we provide another experiment in which the values of θ_i are unknown, and the data are generated by copying actual data with noises. The experiment is described as follows:

Case 3: We collect per-capita gross domestic product of the United States and Democratic Republic of the Congo and denote them as X_t^U and X_t^C , respectively.⁴ The simulation data are generated by $N/2$ copies of X_t^U and X_t^C with noises as follows:

$$X_{it}^j = X_t^j + \theta_{it}^0 X_t^j, \quad j = \{U, C\}$$

θ_{it}^0 is set as described above except that $\alpha_i = (0.1, 0.3, 0.6, 0.8)$.

The result is given in table 3, which shows that the false discovery rate is lower than 5% for all combinations of N and α_i except for $(N = 80, \alpha_i = 0.1)$. Taking the results presented in tables 1, 2, and 3 together, we can conclude that the clustering algorithm generally achieves a satisfactory performance.

Table 3. Simulation result of case 3

| $\alpha_i \backslash N$ | 40 | 80 | 120 |
|-------------------------|-------|-------|-------|
| 0.1 | 0.034 | 0.054 | 0.044 |
| 0.3 | 0.013 | 0.030 | 0.044 |
| 0.6 | 0.014 | 0.030 | 0.041 |
| 0.8 | 0.010 | 0.031 | 0.040 |

9 Example

The example provided here is a replication of the key results of Phillips and Sul (2009). They collected panel data covering 152 economies during the period of 1970–2003. They first examined whether the convergence hypothesis holds for the whole sample. Then, they investigated the possibility of club convergence using their proposed clustering algorithm. The replication is conducted as follows:

4. The period is 1950–2014, namely, $T = 65$.

```

. use ps2009
(PWT152, from Phillips and Sul (2009) in Journal of Applied Econometrics)
. egen id=group(country)
. xtset id year
    panel variable: id (strongly balanced)
    time variable: year, 1970 to 2003
    delta: 1 unit
. generate lnpgd2=ln(pgdp)
. pfilter lnpgd2, method(hp) trend(lnpgd2) smooth(400)

```

First, the `pfilter` command is used to wipe out the cyclical component. A new variable, `lnpgd2`, is generated to store the trend component. We then run the $\log t$ regression for the convergence test. The output reports the coefficient, standard error, and t statistic for $\log(t)$. Because the value of the t statistic (calculated as -159.55) is less than -1.65 , the null hypothesis of convergence is rejected at the 5% level.

```

. logtreg lnpgd2, kq(0.333)
log t test:
-----  

Variable | Coeff SE T-stat  

-----  

log(t) | -0.8748 0.0055 -159.5544
-----  

The number of individuals is 152.  

The number of time periods is 34.  

The first 11 periods are discarded before regression.

```

Furthermore, identifying convergence clubs is conducted by the `psecta` command. The output presents the club classifications. The `noprtlogtreg` option suppresses the estimation details. If not, the results of the $\log t$ regression would be displayed following each club. We put all the estimation results together in the matrix `result1` and display it. Additionally, a new variable `club` is generated to store the club classifications.

```

. psecta lnpgd2, name(country) kq(0.333) gen(club) noprt
xxxxxxxxxxxxxxxxxxxxxxxxxxxx Club classifications xxxxxxxxxxxxxxxxxxxxxxxxx
----- Club 1 :(50)-----
| Antigua | Australia | Austria | Belgium | Bermuda | Botswana |
| Brunei | Canada | Cape Verde | Chile | China | Cyprus | Denmark |
| Dominica | Equatorial Guinea | Finland | France | Germany |
| Hong Kong | Iceland | Ireland | Israel | Italy | Japan |
| Korea, Republic of | Kuwait | Luxembourg | Macao | Malaysia |
| Maldives | Malta | Mauritius | Netherlands | New Zealand |
| Norway | Oman | Portugal | Puerto Rico | Qatar | Singapore |
| Spain | St. Kitts & Nevis | St.Vincent & Grenadines | Sweden |
| Switzerland | Taiwan | Thailand | United Arab Emirates |
| United Kingdom | United States |
-----  


```

```

----- Club 2 :(30)-----
| Argentina | Bahamas | Bahrain | Barbados | Belize | Brazil |
| Colombia | Costa Rica | Dominican Republic | Egypt | Gabon |
| Greece | Grenada | Hungary | India | Indonesia | Mexico |
| Netherlands Antilles | Panama | Poland | Saudi Arabia |
| South Africa | Sri Lanka | St. Lucia | Swaziland | Tonga |
| Trinidad &Tobago | Tunisia | Turkey | Uruguay |

----- Club 3 :(21)-----
| Algeria | Bhutan | Cuba | Ecuador | El Salvador | Fiji |
| Guatemala | Iran | Jamaica | Lesotho | Micronesia, Fed. Sts. |
| Morocco | Namibia | Pakistan | Papua New Guinea | Paraguay |
| Peru | Philippines | Romania | Suriname | Venezuela |

----- Club 4 :(24)-----
| Benin | Bolivia | Burkina Faso | Cameroon | Cote d Ivoire |
| Ethiopia | Ghana | Guinea | Honduras | Jordan | Korea, Dem. Rep. |
| Laos | Mali | Mauritania | Mozambique | Nepal | Nicaragua | Samoa |
| Solomon Islands | Syria | Tanzania | Uganda | Vanuatu | Zimbabwe |

----- Club 5 :(14)-----
| Cambodia | Chad | Comoros | Congo, Republic of | Gambia, The |
| Iraq | Kenya | Kiribati | Malawi | Mongolia | Nigeria |
| Sao Tome and Principe | Senegal | Sudan |

----- Club 6 :(11)-----
| Afghanistan | Burundi | Central African Republic |
| Guinea-Bissau | Madagascar | Niger | Rwanda | Sierra Leone |
| Somalia | Togo | Zambia |

----- Club 7 :(2)-----
| Congo, Dem. Rep. | Liberia |

xxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxx

. matrix b=e(bm)
. matrix t=e(tm)
. matrix result1=(b \ t)
. matlist result1, border(rows) rowtitle("log(t)") format(%9.3f) left(4)



| log(t) | Club1 | Club2 | Club3 | Club4 | Club5 |
|--------|-------|-------|-------|-------|-------|
| Coeff  | 0.382 | 0.240 | 0.110 | 0.131 | 0.190 |
| T-stat | 9.282 | 6.904 | 3.402 | 2.055 | 1.701 |



| log(t) | Club6 | Club7  |
|--------|-------|--------|
| Coeff  | 1.003 | -0.470 |
| T-stat | 6.024 | -0.559 |


```

Finally, we use the `scheckmerge` and `imergeclub` commands to perform possible club merging. We see that the initial clubs 4 and 5 can be merged to form a larger convergent club. The results obtained here are the same as those in Phillips and Sul (2009) and Schnurbus, Haupt, and Meier (2017).

```
. scheckmerge lnpgd2, kq(0.333) club(club) mdiv
```

The log t test for Club 1+2

log t test:

| Variable | Coeff | SE | T-stat |
|----------|---------|--------|---------|
| log(t) | -0.0507 | 0.0232 | -2.1909 |

The number of individuals is 80.

The number of time periods is 34.

The first 11 periods are discarded before regression.

The log t test for Club 2+3

log t test:

| Variable | Coeff | SE | T-stat |
|----------|---------|--------|---------|
| log(t) | -0.1041 | 0.0159 | -6.5339 |

The number of individuals is 51.

The number of time periods is 34.

The first 11 periods are discarded before regression.

The log t test for Club 3+4

log t test:

| Variable | Coeff | SE | T-stat |
|----------|---------|--------|---------|
| log(t) | -0.1920 | 0.0379 | -5.0684 |

The number of individuals is 45.

The number of time periods is 34.

The first 11 periods are discarded before regression.

The log t test for Club 4+5

log t test:

| Variable | Coeff | SE | T-stat |
|----------|---------|--------|---------|
| log(t) | -0.0443 | 0.0696 | -0.6360 |

The number of individuals is 38.

The number of time periods is 34.

The first 11 periods are discarded before regression.

```

The log t test for Club 5+6
log t test:
-----  


| Variable | Coeff   | SE     | T-stat  |
|----------|---------|--------|---------|
| log(t)   | -0.2397 | 0.0612 | -3.9178 |


The number of individuals is 25.
The number of time periods is 34.
The first 11 periods are discarded before regression.
-----  

The log t test for Club 6+7
log t test:
-----  


| Variable | Coeff   | SE     | T-stat   |
|----------|---------|--------|----------|
| log(t)   | -1.1163 | 0.0602 | -18.5440 |


The number of individuals is 13.
The number of time periods is 34.
The first 11 periods are discarded before regression.
-----  

. matrix b=e(bm)
. matrix t=e(tm)
. matrix result2=(b \ t)
. matlist result2, border(rows) rowtitle("log(t)") format(%9.3f) left(4)
-----  


| log(t) | Club1+2 | Club2+3 | Club3+4 | Club4+5 | Club5+6 |
|--------|---------|---------|---------|---------|---------|
| Coeff  | -0.051  | -0.104  | -0.192  | -0.044  | -0.240  |
| T-stat | -2.191  | -6.534  | -5.068  | -0.636  | -3.918  |


-----  


| log(t) | Club6+7 |
|--------|---------|
| Coeff  | -1.116  |
| T-stat | -18.544 |


-----  

. imergeclub lnpgd2, name(country) kq(0.333) club(club) gen(finalclub) nopr
xxxxxxxxxxxxxxxxxxxxxxxxxxxx Final Club classifications xxxxxxxxxxxxxxxxxxxxxxxx
----- Club 1 :(50)-----
| Antigua | Australia | Austria | Belgium | Bermuda | Botswana |
| Brunei | Canada | Cape Verde | Chile | China | Cyprus | Denmark |
| Dominica | Equatorial Guinea | Finland | France | Germany |
| Hong Kong | Iceland | Ireland | Israel | Italy | Japan |
| Korea, Republic of | Kuwait | Luxembourg | Macao | Malaysia |
| Maldives | Malta | Mauritius | Netherlands | New Zealand |
| Norway | Oman | Portugal | Puerto Rico | Qatar | Singapore |
| Spain | St. Kitts & Nevis | St.Vincent & Grenadines | Sweden |
| Switzerland | Taiwan | Thailand | United Arab Emirates |
| United Kingdom | United States |
-----
```

```

----- Club 2 :(30)-----
| Argentina | Bahamas | Bahrain | Barbados | Belize | Brazil |
| Colombia | Costa Rica | Dominican Republic | Egypt | Gabon |
| Greece | Grenada | Hungary | India | Indonesia | Mexico |
| Netherlands Antilles | Panama | Poland | Saudi Arabia |
| South Africa | Sri Lanka | St. Lucia | Swaziland | Tonga |
| Trinidad &Tobago | Tunisia | Turkey | Uruguay |

----- Club 3 :(21)-----
| Algeria | Bhutan | Cuba | Ecuador | El Salvador | Fiji |
| Guatemala | Iran | Jamaica | Lesotho | Micronesia, Fed. Sts. |
| Morocco | Namibia | Pakistan | Papua New Guinea | Paraguay |
| Peru | Philippines | Romania | Suriname | Venezuela |

----- Club 4 :(38)-----
| Benin | Bolivia | Burkina Faso | Cambodia | Cameroon | Chad |
| Comoros | Congo, Republic of | Cote d Ivoire | Ethiopia |
| Gambia, The | Ghana | Guinea | Honduras | Iraq | Jordan | Kenya |
| Kiribati | Korea, Dem. Rep. | Laos | Malawi | Mali |
| Mauritania | Mongolia | Mozambique | Nepal | Nicaragua |
| Nigeria | Samoa | Sao Tome and Principe | Senegal |
| Solomon Islands | Sudan | Syria | Tanzania | Uganda | Vanuatu |
| Zimbabwe |

----- Club 5 :(11)-----
| Afghanistan | Burundi | Central African Republic |
| Guinea-Bissau | Madagascar | Niger | Rwanda | Sierra Leone |
| Somalia | Togo | Zambia |

----- Club 6 :(2)-----
| Congo, Dem. Rep. | Liberia |

xxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxxx

. matrix b=e(bm)
. matrix t=e(tm)
. matrix result3=(b \ t)
. matlist result3, border(rows) rowtitle("log(t)") format(%9.3f) left(4)



| log(t) | Club1 | Club2 | Club3 | Club4  | Club5 |
|--------|-------|-------|-------|--------|-------|
| Coeff  | 0.382 | 0.240 | 0.110 | -0.044 | 1.003 |
| T-stat | 9.282 | 6.904 | 3.402 | -0.636 | 6.024 |



| log(t) | Club6  |
|--------|--------|
| Coeff  | -0.470 |
| T-stat | -0.559 |


```

10 Acknowledgments

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