

Topic 5: Two-Way Cluster-Robust (TWCR) Standard Errors

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Key points: The validity of Two-Way Cluster-Robust (TWCR) standard errors

Disclaimer: This note is compiled by Sai Zhang.

5.1 One-Way Clustering

First, consider the case of one-way clustering. The linear model with one-way clustering

$$y_{ig} = \mathbf{x}_{ig}\boldsymbol{\beta} + u_{ig}$$

where i denotes the i th of the N individuals in the sample, j denotes the g th of the G clusters, assume that

- $\mathbb{E}[u_{ig} | \mathbf{x}_{ig}] = 0$
- error independence across clusters: for $i \neq j$

$$\mathbb{E}[u_{ig}u_{jg'} | \mathbf{x}_{ig}, \mathbf{x}_{jg'}] = 0 \quad (5.1)$$

unless $g = g'$, that is, errors for individuals within the same cluster may be correlated.

Grouping observations by cluster, get

$$\mathbf{y}_g = \mathbf{X}_g\boldsymbol{\beta} + \mathbf{u}$$

where \mathbf{X}_g has dimension $N_g \times K$ and \mathbf{y}_g has dimension $N_g \times 1$, with N_g observations in cluster g . Stacking over cluster, get the matrix form of the model

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mathbf{u}$$

with \mathbf{y}, \mathbf{u} being $N \times 1$ vectors, \mathbf{X} being an $N \times K$ matrix. OLS estimator gives

$$\hat{\boldsymbol{\beta}} = (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}'\mathbf{y} = \left(\sum_{g=1}^G \mathbf{X}_g' \mathbf{X}_g \right)^{-1} \sum_{g=1}^G \mathbf{X}_g' \mathbf{y}_g \quad (5.2)$$

then, by CLT, we have that $\sqrt{G}(\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}) \xrightarrow{d} \mathcal{N}(0, \boldsymbol{\Sigma})$ where the variance matrix of the limit normal distribution $\boldsymbol{\Sigma}$ is

$$\left(\lim_{G \rightarrow \infty} \frac{1}{G} \sum_{g=1}^G \mathbb{E}[\mathbf{X}_g' \mathbf{X}_g] \right)^{-1} \left(\lim_{G \rightarrow \infty} \frac{1}{G} \sum_{g=1}^G \mathbb{E}[\mathbf{X}_g' \mathbf{u}_g' \mathbf{u}_g \mathbf{X}_g] \right) \times \left(\lim_{G \rightarrow \infty} \frac{1}{G} \sum_{g=1}^G \mathbb{E}[\mathbf{X}_g' \mathbf{X}_g] \right)^{-1} \quad (5.3)$$

If the primary source of clustering is due to group-level common shocks, a useful approximation is that for the j th regressor, the default OLS variance estimate based on $s^2(\mathbf{X}'\mathbf{X})^{-1}$ should be inflated by $\tau_j \approx 1 + \rho_{x_j}\rho_u(\bar{N}_g - 1)$, where

- s is the estimated standard deviation of the error

- ρ_{x_j} is a measure of within-cluster correlation of x_j
- ρ_u is the within-cluster error correlation
- \bar{N}_g is the average cluster size

It's easy to see the τ_j can be large even with small ρ_u (Kloek, 1981; Scott and Holt, 1982; Moulton, 1990). If assume the model for the cluster error variance matrices $\mathbf{\Omega}_g = \mathbb{V}[\mathbf{u}_g | \mathbf{X}_g] = \mathbb{E}[\mathbf{u}_g \mathbf{u}_g' | \mathbf{X}_g]$, and there is a consistent estimate $\hat{\mathbf{\Omega}}_g$ of $\mathbf{\Omega}_g$, we can estimate $\mathbb{E}[\mathbf{X}_g' \mathbf{u}_g \mathbf{u}_g' \mathbf{X}_g] = \mathbb{E}[\mathbf{X}_g' \mathbf{\Omega}_g \mathbf{X}_g]$ via GLS.

Cluster-robust variance matrix estimate consider

$$\hat{\mathbb{V}}[\hat{\beta}] = (\mathbf{X}'\mathbf{X})^{-1} \left(\sum_{g=1}^G \mathbf{X}_g' \hat{\mathbf{u}}_g \hat{\mathbf{u}}_g' \mathbf{X}_g \right) (\mathbf{X}'\mathbf{X})^{-1} \quad (5.4)$$

where $\hat{\mathbf{u}}_g = \mathbf{y}_g - \mathbf{X}_g \hat{\beta}$. This estimate is consistent if

$$G^{-1} \sum_{g=1}^G \mathbf{X}_g' \hat{\mathbf{u}}_g \hat{\mathbf{u}}_g' \mathbf{X}_g - G^{-1} \sum_{g=1}^G \mathbb{E}[\mathbf{X}_g' \mathbf{u}_g \mathbf{u}_g' \mathbf{X}_g] \xrightarrow{P} \mathbf{0}$$

as $G \rightarrow \infty$. An informal presentation of Eq.(5.4) is to rewrite the central matrix as

$$\hat{\mathbf{B}} = \sum_{g=1}^G \mathbf{X}_g' \hat{\mathbf{u}}_g \hat{\mathbf{u}}_g' \mathbf{X}_g = \mathbf{X}' \begin{bmatrix} \hat{\mathbf{u}}_1 \hat{\mathbf{u}}_1' & \mathbf{0} & \cdots & \mathbf{0} \\ \mathbf{0} & \hat{\mathbf{u}}_2 \hat{\mathbf{u}}_2' & & \vdots \\ \vdots & & \ddots & \mathbf{0} \\ \mathbf{0} & \cdots & & \hat{\mathbf{u}}_G \hat{\mathbf{u}}_G' \end{bmatrix} \mathbf{X} = \mathbf{X}' (\hat{\mathbf{u}} \hat{\mathbf{u}}' \otimes \mathbf{S}^G) \mathbf{X} \quad (5.5)$$

where \otimes denotes element-wise multiplication. The (p, q) th element of this matrix is

$$\sum_{i=1}^N \sum_{j=1}^N x_{ia} x_{jb} \hat{u}_i \hat{u}_j \cdot \mathbf{1}(i, j \text{ in the same cluster})$$

with $\hat{u}_i = y_i - \mathbf{x}_i' \hat{\beta}$.

\mathbf{S}^G is an $N \times N$ indicator matrix with $\mathbf{S}_{ij}^G = 1$ only if the i th and j th observation belong to the same cluster: it zeros out a large amount of $\hat{\mathbf{u}} \hat{\mathbf{u}}'$ (asymptotically equivalently, $\mathbf{u} \mathbf{u}'$), specifically, only $\sum_{g=1}^G N_g^2$ out of $N^2 = \left(\sum_{g=1}^G N_g \right)^2$ terms are not zero (sub-matrices on the diagonal). Asymptotically

- for fixed N_g , $\frac{1}{N^2} \sum_{g=1}^G N_g^2 \xrightarrow{G \rightarrow \infty} 0$
- for balanced clusters $N_g = N/G$, $\frac{1}{N^2} \sum_{g=1}^G N_g^2 = \frac{1}{G} \xrightarrow{G \rightarrow \infty} 0$

A strand of literature popularizes this method:

- Liang and Zeger (1986): in a generalized estimatin equations setting
- Arellano (1987): fixed effects estimator in linear panel models
- Hansen (2007): asymptotic theory for panel data where $T \rightarrow \infty$ in addition to $N \rightarrow \infty$ (or $N_g \rightarrow \infty$ in addition to $G \rightarrow \infty$ in the notation above).

5.2 Two-Way Clustering

Now, consider the case of two-way clustering,

$$y_{i,gh} = \mathbf{x}'_{i,gh} \boldsymbol{\beta} + u$$

where each observation may belong to **two** dimension of groups: group $g \in \{1, \dots, G\}$ and $h \in \{1, \dots, H\}$, and for $i \neq j$

$$\mathbb{E} [u_{i,gh} u_{j,g'h'} | \mathbf{x}_{i,gh}, \mathbf{j}, \mathbf{g}'\mathbf{h}'] = 0 \quad (5.6)$$

unless $g = g'$ or $h = h'$, that is, errors for individuals within the same group (along either g or h) may be correlated.

Cluster-robust variance matrix estimate extending the one-way clustering case, keep elements of $\hat{\mathbf{u}}\hat{\mathbf{u}}'$ where the i th and j th observations share a cluster in **any** dimension, then similar to Eq.(5.5)

$$\hat{\mathbf{B}} = \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^{GH} \right) \mathbf{X} \quad (5.7)$$

here \mathbf{S}^{GH} is an $N \times N$ indicator matrix with $S_{ij}^{GH} = 1$ only if the i th and j th observation share any cluster, the (p, q) th element of this matrix is

$$\sum_{i=1}^N \sum_{j=1}^N x_{ia} x_{jb} \hat{u}_i \hat{u}_j \cdot \mathbf{1}(i, j \text{ share any cluster})$$

$\hat{\mathbf{B}}$ can also be presented in one-way cluster-robust fashion:

$$\hat{\mathbf{B}} = \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^{GH} \right) \mathbf{X} = \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^G \right) \mathbf{X} + \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^H \right) \mathbf{X} - \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^{G \cap H} \right) \mathbf{X} \quad (5.8)$$

where $\mathbf{G}^{GH} = \mathbf{G}^G + \mathbf{G}^H - \mathbf{G}^{G \cap H}$, with

- \mathbf{G}^G : $G_{ij}^G = 1$ only if the i th and j th observation belong to the same cluster $g \in \{1, 2, \dots, G\}$
- \mathbf{G}^H : $G_{ij}^H = 1$ only if the i th and j th observation belong to the same cluster $h \in \{1, 2, \dots, H\}$
- $\mathbf{G}^{G \cap H}$: $G_{ij}^{G \cap H} = 1$ only if the i th and j th observation belong to **both** the same cluster $g \in \{1, 2, \dots, G\}$ and the same cluster $h \in \{1, 2, \dots, H\}$

then, similar to one-way clustering case,

$$\begin{aligned} \hat{\mathbf{V}}[\hat{\boldsymbol{\beta}}] &= (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^G \right) \mathbf{X} (\mathbf{X}'\mathbf{X})^{-1} \\ &\quad + (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^H \right) \mathbf{X} (\mathbf{X}'\mathbf{X})^{-1} \\ &\quad - (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}' \left(\hat{\mathbf{u}}\hat{\mathbf{u}}' \otimes \mathbf{S}^{G \cap H} \right) \mathbf{X} (\mathbf{X}'\mathbf{X})^{-1} \end{aligned} \quad (5.9)$$

that is,

$$\hat{\mathbf{V}}[\hat{\boldsymbol{\beta}}] = \hat{\mathbf{V}}^G[\hat{\boldsymbol{\beta}}] + \hat{\mathbf{V}}^H[\hat{\boldsymbol{\beta}}] - \hat{\mathbf{V}}^{G \cap H}[\hat{\boldsymbol{\beta}}] \quad (5.10)$$

each of Eq.(5.10) can be separately computed by OLS of \mathbf{y} on \mathbf{X} , with variance matrix estimates $\hat{\mathbf{V}}$ based on

- clustering on $g \in \{1, 2, \dots, G\}$
- clustering on $h \in \{1, 2, \dots, H\}$
- clustering on $(g, h) \in \{(1, 1), \dots, (G, H)\}$

Practical considerations It is required to know what *ways* will be potentially important for clustering, which can be tested via checking the dimension of correlations in the errors. There are several ways to test

- estimate sample covariances of $\mathbf{X}'\hat{\mathbf{u}}$ within dimensions, test the null that the **average** of such covariances is 0: rejecting this null is sufficient (not necessary) to reject the null of no clustering (White, 1980)
- for **small samples**, Eq. (5.4) is biased downwards. This is corrected (in Stata) by replacing $\hat{\mathbf{u}}_g$ with $\sqrt{c}\hat{\mathbf{u}}_g$, where $c = \frac{G}{G-1} \frac{N-1}{N-K} \simeq \frac{G}{G-1}$. For two-way clustering (Eq. 5.8), there are 2 ways of correction:
 - choose correction terms for each of the 3 components:

$$c_1 = \frac{G}{G-1} \frac{N-1}{N-K}, c_2 = \frac{H}{H-1} \frac{N-1}{N-K}, c_3 = \frac{I}{I-1} \frac{N-1}{N-K}$$

with I being the number of unique clusters determined by $G \cap H$

- choose a constant terms for all components:

$$c = \frac{J}{J-1} \frac{N-1}{N-K}$$

with $J = \min(G, H)$

- **Var-cov matrix not positive-semidefinite**: $\hat{\mathbf{V}}[\hat{\boldsymbol{\beta}}]$ might have negative elements on the diagonal (Eq. 5.10), informly, this is more likely to arise when clustering is done over the same groups as the fixed effects. One way to address this issue is using *eigendecomposition* technique:

$$\hat{\mathbf{V}}[\hat{\boldsymbol{\beta}}] = \mathbf{U}\mathbf{\Lambda}\mathbf{U}'$$

where

- \mathbf{U} containing the eigenvectors of $\hat{\mathbf{V}}$
- $\mathbf{\Lambda} = \text{diag}[\lambda_1, \dots, \lambda_d]$ contains the eigenvalues of $\hat{\mathbf{V}}$

then create $\mathbf{\Lambda}^+ = \text{diag}[\lambda_1^+, \dots, \lambda_d^+]$ with $\lambda_j^+ = \max(0, \lambda_j)$ and use $\hat{\mathbf{V}}^+[\hat{\boldsymbol{\beta}}] = \mathbf{U}\mathbf{\Lambda}^+\mathbf{U}'$ as the estimate

5.3 Multiway Clustering

Cameron et al. (2011) extended the framework¹ to allow clustering in D dimensions, then we can do the following reframing

- G_d : the number of clusters in dimension $d \in \{1, 2, \dots, D\}$
- D -vector $\boldsymbol{\delta}_i = \boldsymbol{\delta}(i)$, with function $\boldsymbol{\delta} : \{1, 2, \dots, N\} \rightarrow \times_{d=1}^D \{1, 2, \dots, G_d\}$ lists the cluster membership in each dimension of each observation

then we have

$$\mathbf{1}[i, j \text{ shares a cluster}] = 1 \Leftrightarrow \delta_{id} = \delta_{jd}$$

for some $d \in \{1, 2, \dots, D\}$, where δ_{id} denotes the d th element of $\boldsymbol{\delta}_i$. Also

- D -vector \mathbf{r} : define the set

$$R \equiv \{\mathbf{r} : r_d \in \{0, 1\}, d = 1, 2, \dots, D, \mathbf{r} \neq \mathbf{0}\}$$

elements of the set R can be used to index all cases where 2 observations share a cluster in at least one dimension. Define the function

$$\mathbf{I}_r(i, j) \equiv \mathbf{1}[r_d \delta_{id} = r_d \delta_{jd}, \forall d]$$

¹Also proposed by Thompson (2011).

which indicates whether observations i and j have identical cluster membership for **all** dimensions d s.t. $r_d = 1$. Then we have a *aggregate* identifier

$$\mathbf{I}(i, j) = 1 \Leftrightarrow \mathbf{I}_r(i, j) = 1 \text{ for some } \mathbf{r} \in R$$

i.e., 2 observations share **at least** one dimension.

The define the $2^D - 1$ matrices

$$\tilde{\mathbf{B}}_r \equiv \sum_{i=1}^N \sum_{j=1}^N \mathbf{x}_i \mathbf{x}_j' \hat{u}_i \hat{u}_j \mathbf{I}_r(i, j) \quad (5.11)$$

with $\mathbf{r} \in R$.

Var-cov matrix estimator consider, similarly, an estimator

$$\hat{\mathbb{V}}[\hat{\beta}] = (\mathbf{X}'\mathbf{X})^{-1} \tilde{\mathbf{B}} (\mathbf{X}'\mathbf{X})^{-1} \equiv (\mathbf{X}'\mathbf{X})^{-1} \left(\sum_{\|\mathbf{r}\|=k, \mathbf{r} \in R} (-1)^{k+1} \tilde{\mathbf{B}}_r \right) (\mathbf{X}'\mathbf{X})^{-1} \quad (5.12)$$

where cases of clustering on an odd number of dimensions are added, those of clustering on an even number of dimensions are subtracted. Consider the case of $D = 3$,

$$(\tilde{\mathbf{B}}_{(1,0,0)} + \tilde{\mathbf{B}}_{(0,1,0)} + \tilde{\mathbf{B}}_{(0,0,1)}) - (\tilde{\mathbf{B}}_{(1,1,0)} + \tilde{\mathbf{B}}_{(1,0,1)} + \tilde{\mathbf{B}}_{(0,1,1)}) + \tilde{\mathbf{B}}_{(1,1,1)}$$

$\tilde{\mathbf{B}}$ is identical to $\hat{\mathbf{B}}$ defined analogically as in Eq.(5.8), since

- no observation pair with $\mathbf{I}(i, j) = 0$: this is immediate, since $\mathbf{I}(i, j) = 0 \Leftrightarrow \mathbf{I}_r(i, j) = 0, \forall \mathbf{r}$
- the covariance term corresponding to each observation pair with $\mathbf{I}(i, j) = 1$ is included **exactly once** in $\tilde{\mathbf{B}}$: by inclusion-exclusion principle for set cardinality

$$\mathbf{I}(i, j) \Rightarrow \sum_{\|\mathbf{r}\|=k, \mathbf{r} \in R} (-1)^{k+1} \mathbf{I}_r(i, j) = 1$$

Curse of dimensionality this could arise in a setting with **many dimensions** of clustering, and in which one or more dimensions have **few** clusters². **Cameron et al. (2011)** suggested an ad-hoc rule of thumb for approximating sufficient numbers of clusters.

5.3.1 Non-linear Estimators

m -Estimators Consider an m -estimator that solves

$$\sum_{i=1}^N \mathbf{h}_i(\hat{\theta}) = \mathbf{0}$$

under standard assumptions, $\hat{\theta}$ is asymptotically normal with estimated variance matrix

$$\hat{\mathbb{V}}[\hat{\theta}] = \hat{\mathbf{A}}^{-1} \hat{\mathbf{B}} \hat{\mathbf{A}}'^{-1} \quad (5.13)$$

where $\hat{\mathbf{A}} = \sum_i \frac{\partial \mathbf{h}_i}{\partial \theta'} \Big|_{\hat{\theta}}$ and $\hat{\mathbf{B}}$ is an estimate of $\mathbb{V}[\sum_i \mathbf{h}_i]$.

²The square design (each dimension has the same number of clusters) with orthogonal dimensions has the **least** independence of observations.

- **one-way clustering** $\hat{\mathbf{B}} = \sum_{g=1}^G \hat{\mathbf{h}}_g \hat{\mathbf{h}}_g'$ where $\hat{\mathbf{h}}_g = \sum_{i=1}^{N_g} \hat{\mathbf{h}}_{ig}$, clustering may not lead to parameter inconsistency, depending on whether $\mathbb{E}[\mathbf{h}_i(\boldsymbol{\theta})] = \mathbf{0}$ with clustering
 - **population-averaged approach**: assume $\mathbb{E}[y_{ig} | \mathbf{x}_{ig}] = \Phi(\mathbf{x}_{ig}'\boldsymbol{\beta})$
 - **random effects approach**: let $y_{ig} = 1$ if $y_{ig}^* > 0$ where $y_{ig}^* = \mathbf{x}_{ig}'\boldsymbol{\beta} + \epsilon_g + \epsilon_{ig}$, where
 - * idiosyncratic error $\epsilon_{ig} \sim \mathcal{N}(0, 1)$
 - * cluster-specific error $\epsilon_g \sim \mathcal{N}(0, \sigma_g^2)$
 then we have the alternative moment condition

$$\mathbb{E}[y_{ig} | \mathbf{x}_{ig}] = \Phi\left(\frac{\mathbf{x}_{ig}'\boldsymbol{\beta}}{\sqrt{1 + \sigma_g^2}}\right)$$

- **multiway clustering** replacing $\hat{u}_i \mathbf{x}_i$ in Eq.(5.11) with $\hat{\mathbf{h}}_i$, then we have

$$\hat{\mathbb{V}}[\hat{\boldsymbol{\theta}}] = \hat{\mathbf{A}}^{-1} \hat{\mathbf{B}} \hat{\mathbf{A}}^{-1}$$

where

$$\hat{\mathbf{A}} = \sum_i \frac{\partial \mathbf{h}_i}{\partial \boldsymbol{\theta}'} \bigg|_{\hat{\boldsymbol{\theta}}} \quad \hat{\mathbf{B}} = \sum_{\|\mathbf{r}\|=k, \mathbf{r} \in R} (-1)^{k+1} \tilde{\mathbf{B}}_r \quad \tilde{\mathbf{B}}_r \equiv \sum_{i=1}^N \sum_{j=1}^N \hat{\mathbf{h}}_i \hat{\mathbf{h}}_j' \mathbb{I}_r(i, j)$$

with $\mathbf{r} \in R^3$.

GMM estimation Consider an example of over-identified models: linear two stage least squares with more instruments than endogenous regressors, we have

$$\hat{\boldsymbol{\theta}} = \arg \min_{\boldsymbol{\theta}} Q(\boldsymbol{\theta}) = \arg \min_{\boldsymbol{\theta}} \left(\sum_{i=1}^N \mathbf{h}_i(\boldsymbol{\theta}) \right)' \mathbf{W} \left(\sum_{i=1}^N \mathbf{h}_i(\boldsymbol{\theta}) \right)$$

where \mathbf{W} is a symmetric positive definite weighting matrix. Under standard regularity conditions, $\hat{\boldsymbol{\theta}}$ is asymptotically normal, with estimated variance matrix

$$\hat{\mathbb{V}}[\hat{\boldsymbol{\theta}}] = (\hat{\mathbf{A}}' \mathbf{W} \hat{\mathbf{A}})^{-1} \hat{\mathbf{A}}' \mathbf{W} \hat{\mathbf{B}} \mathbf{W} \hat{\mathbf{A}} (\hat{\mathbf{A}}' \mathbf{W} \hat{\mathbf{A}})^{-1}$$

again, $\hat{\mathbf{A}} = \sum_i \frac{\partial \mathbf{h}_i}{\partial \boldsymbol{\theta}'} \bigg|_{\hat{\boldsymbol{\theta}}}$, and $\hat{\mathbf{B}}$ is an estimate of $\mathbb{V}[\sum_i \mathbf{h}_i]$.

5.4 Menzel (2021): Asymptotic Gaussianity

One key of TWCR inference is the asymptotic Gaussianity, [Menzel \(2021\)](#) pointed out the potential non-Gaussianity of the limit distribution. Still, consider a random array (Y_{it}) indexed by two dimensions by $i = 1, \dots, N$ and $t = 1, \dots, T$. Clusters are sampled independently at random from an infinite population, but otherwise **unrestricted** in dependence within each row $\mathbf{Y}_i := (Y_{i1} \dots, Y_{iT})$ and within each column $\mathbf{Y}_t := (Y_{1t}, \dots, Y_{Nt})$.

³This multiway clustering can be implemented using several one-way clustered bootstraps. Each of the one-way cluster robust matrices is estimated by a pairs cluster bootstrap that resamples with replacement from the appropriate cluster dimension. They are then combined as if they had been estimated analytically ([Cameron et al., 2011](#)).

5.4.1 Distribution of Sample Average

First, consider

$$\bar{Y}_{NT} := \frac{1}{NT} \sum_{i=1}^N \sum_{t=1}^T Y_{it}$$

and approximate the asymptotic distribution regardless of whether, or what type of, cluster-dependence is present.

3 scenarios of the array (Y_{it})

- **no cluster-dependence**: (Y_{it}) are mutually independent, CLT at a rate of $(NT)^{-1/2}$ applies (under regularity conditions)
- **correlation within clusters**: the convergence rate of (Y_{it}) is determined by the number of relevant clusters
- **non-separable models of heterogeneity (dependence with clusters, even uncorrelated)**⁴: The asymptotic behavior is non-standard

Consider 2 examples:

- **Additive factor model**

$$Y_{it} = \mu + \alpha_i + \gamma_t + \epsilon_{it}$$

where μ is a constant, and $\alpha_i, \gamma_t, \epsilon_{it}$ are zero-mean i.i.d. random variables for $i = 1, \dots, N$ and $t = 1, \dots, T$ with bounded second moments, and $N = T$. Based on a standard central limit theory, we have

- in the non-degenerate case with $\text{Var}(\alpha_i) > 0$ or $\gamma_t > 0$, the sample distribution

$$\sqrt{N} \left(\bar{Y}_{NT} - \mathbb{E}[Y_{it}] \right) \xrightarrow{d} \mathcal{N}(0, \text{Var}(\alpha_i) + \text{Var}(\gamma_t))$$

- in the degenerate case of no clustering with $\text{Var}(\alpha_i) = \text{Var}(\gamma_t) = 0$, the sample distribution

$$\sqrt{NT} \left(\bar{Y}_{NT} - \mathbb{E}[Y_{it}] \right) \xrightarrow{d} \mathcal{N}(0, \text{Var}(\epsilon_{it}))$$

if marginal distributions of $\alpha_i, \gamma_t, \epsilon_{it}$ are known, we can simulate from the joint distribution of (Y_{it}) by sampling the individual components at random, a bootstrap procedure would be consistent. If **unknown**, consider estimators

$$\begin{aligned} \hat{\alpha}_i &:= \frac{1}{T} \sum_{t=1}^T (Y_{it} - \bar{Y}_{NT}) = \alpha_i + \frac{1}{T} \sum_{t=1}^T (\epsilon_{it} - \bar{\epsilon}_{NT}) \\ \hat{\gamma}_t &:= \frac{1}{N} \sum_{i=1}^N (Y_{it} - \bar{Y}_{NT}) = \gamma_t + \frac{1}{N} \sum_{i=1}^N (\epsilon_{it} - \bar{\epsilon}_{NT}) \\ \hat{\epsilon}_{it} &:= Y_{it} - \bar{Y}_{NT} - \hat{\alpha}_i - \hat{\gamma}_t \end{aligned}$$

then use these empirical distributions for estimation and form a bootstrap sample

$$Y_{it}^* := \bar{Y}_{NT} + \alpha_i^* + \gamma_t^* + \epsilon_{it}^*$$

⁴This is specific to clustering in 2 or more dimensions.

by drawing from these estimators and obtain $\bar{Y}_{NT}^* := \frac{1}{NT} \sum_{i=1}^N \sum_{t=1}^T Y_{it}^*$, and verify the conditional variances of the bootstrap distribution given the sample:

$$\begin{aligned} \frac{1}{N} \sum_{i=1}^N \left(\hat{\alpha}_i - \frac{1}{N} \sum_{j=1}^N \hat{\alpha}_j \right)^2 - \left[\text{Var}(\alpha_i) + \frac{\text{Var}(\epsilon_{it})}{T} \right] &\xrightarrow{p} 0 \\ \frac{1}{T} \sum_{t=1}^T \left(\hat{\gamma}_t - \frac{1}{T} \sum_{s=1}^T \hat{\gamma}_s \right)^2 - \left[\text{Var}(\gamma_t) + \frac{\text{Var}(\epsilon_{it})}{N} \right] &\xrightarrow{p} 0 \end{aligned}$$

then the bootstrap distribution is

– in the non-degenerate case,

$$\sqrt{N} \left(\bar{Y}_{NT}^* - \bar{Y}_{NT} \right) \xrightarrow{d} \mathcal{N} \left(0, \text{Var}(\alpha_i) + \text{Var}(\gamma_t) \right)$$

the estimation error $\hat{\alpha}_i$ does **NOT** affect the asymptotic variance.

– in the degenerate case,

$$\sqrt{NT} \left(\bar{Y}_{NT}^* - \bar{Y}_{NT} \right) \xrightarrow{d} \mathcal{N} \left(0, 3\text{Var}(\epsilon_{it}) \right)$$

asymptotically overestimates the variance of the sampling distribution, leading to inconsistency of this naive bootstrapping procedure.

- **Non-Gaussian limit distribution**

$$Y_{it} = \alpha_i \gamma_t + \epsilon_{it}$$

where $\alpha_i, \gamma_t, \epsilon_{it}$ are independently distributed with $\mathbb{E}[\epsilon_{it}] = 0$, $\text{Var}(\alpha_i) = \sigma_\alpha^2$, $\text{Var}(\gamma_t) = \sigma_\gamma^2$, $\text{Var}(\epsilon_{it}) = \sigma_\epsilon^2$.

If $\mathbb{E}[\alpha_i] = \mathbb{E}[\gamma_t] = 0$, then CLT and Continuous Mapping Theorem (CMT) imply

$$\begin{aligned} \sqrt{NT} \cdot \bar{Y}_{NT} &= \frac{1}{\sqrt{NT}} \sum_{i=1}^N \sum_{t=1}^T (\alpha_i \gamma_t + \epsilon_{it}) \\ &= \left(\frac{1}{\sqrt{N}} \sum_{i=1}^N \alpha_i \right) \left(\frac{1}{\sqrt{T}} \sum_{t=1}^T \gamma_t \right) + \frac{1}{\sqrt{NT}} \sum_{i=1}^N \sum_{t=1}^T \epsilon_{it} \\ &\xrightarrow{d} \sigma_\alpha \sigma_\gamma Z_1 Z_2 + \sigma_\epsilon Z_3 \end{aligned}$$

then even without correlation within clusters, non-separable heterogeneity can still generate dependence in 2nd or higher moments in the limiting distribution⁵.

5.4.2 Menzel (2021)'s Bootstrap procedure

5.4.2.1 Notation

For the array $(Y_{it})_{i,t}$, denote

- \mathbb{P} : joint distribution of $(Y_{it})_{i,t}$

⁵2 major issues arise:

- The limiting distribution needs **not** be Gaussian: plug-in asymptotic inference based on the normal distribution is invalid
- It only comes from two-or-more-dimension cluster dependence, not single-dimension cluster dependence.

- \mathbb{P}_{NT} : drifting DGP indexed by N, T
- \mathbb{P}_{NT}^* : bootstrap distribution for (Y_{it}^*) given the realizations $(Y_{it} : i = 1, \dots, N; t = 1, \dots, T)$
- respective distributions $\mathbb{E}, \mathbb{E}_{NT}, \mathbb{E}_{NT}^*$

5.4.2.2 Inference: Sample Mean

First, consider the assumption of *separate exchangeability*

Assumption 5.4.1: Separate Exchangeability

- A **separately exchangeable** array is an infinite array $(Y_{it})_{i,t}$ such that for any integers \tilde{N}, \tilde{T} and permutations $\pi_1 : \{1, \dots, \tilde{N}\} \rightarrow \{1, \dots, \tilde{N}\}$ and $\pi_2 : \{1, \dots, \tilde{T}\} \rightarrow \{1, \dots, \tilde{T}\}$, we have

$$(Y_{\pi_1(i), \pi_2(t)})_{i,t} \stackrel{d}{=} (Y_{it})_{i,t}$$

such an array is called **dissociated** if for any $N_0, T_0 \geq 1$, $(Y_{it})_{i=1, t=1}^{i=N_0, t=T_0}$ is independent of $(Y_{it})_{i>N_0, t>T_0}$.

- For dyadic data, consider the alternative assumption **jointly exchangeable** arrays $(Y_{ij})_{i,j}$ satisfying

$$(Y_{\pi(i), \pi(j)})_{i,j} \stackrel{d}{=} (Y_{ij})_{i,j}$$

for any permutation π on $\{1, \dots, \tilde{N}\}$, in addition, $(Y_{ij})_{i,j=1}^{N_0}$ is independent of $(Y_{ij})_{i,j>N_0}$

This assumption can be interpreted as rows (and columns) corresponding to units that are drawn independently from a common population, where we then observe the joint outcome for every row-column pair, consider the requirements in the following applications

- **DiD/matched data**: the units corresponding to either dimension of the sample to represent independent draws from a common, infinite population
- **non-exhaustively matched data**: only observe joint outcomes for a possibly self-selected subset of unit pairs, sample selection should be (jointly or separately) exchangeable
- **U-/V-statistics**: the kernel $Y_{i_1, \dots, i_D} := h(X_{i_1}, \dots, X_{i_D})$ evaluated at i.i.d. observations X_1, \dots, X_N forms a dissociated, jointly exchangeable array
- **Network**: unlabeled⁶ data implies finite exchangeability, the sampled graph has joint (*infinite*) exchangeability if it is a subgraph of an infinite graph

Directly from Assumption 5.4.1, any dissociated separately exchangeable array can be represented as

$$Y_{it} = f(\alpha_i, \gamma_t, \epsilon_{it})$$

for some function $f(\cdot)$ where $\alpha_1, \dots, \alpha_N, \gamma_1, \dots, \gamma_T, \epsilon_{11}, \dots, \epsilon_{NT}$ are mutually independent, uniformly distributed random variables.

A Theoretical

Chiang and Sasaki (2023)

⁶Unlabeled: model identifiers do not carry any significance for the statistical model.

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