

Cluster-robust inference in Stata

Eduardo García Echeverri

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- 1 Clustered errors: Why are they important?
- 2 The cluster-robust variance estimator (CRVE)
- 3 Alternatives when the assumptions of the CRVE fail
 - Adjusted degrees of freedom for `vce(hc2)`
 - Wild cluster bootstrap
- 4 Conclusion

Outline

- 1 Clustered errors: Why are they important?
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Clustered errors in your data

1. Data on firms operating in different sectors:
 - Errors are (probably) correlated within industries.
2. Data on high school students in the USA:
 - Errors are (probably) correlated within schools.
3. Panel data on individuals:
 - For each individual, errors are (probably) serially correlated.
4. Experiments with treatment at an aggregated level.
 - Errors are (probably) correlated within such levels.

Example: States changing minimum wage.

Linear model with clustered errors

Consider the model:

$$y_{ig} = X_{ig}\beta + \varepsilon_{ig}$$

where,

y_{ig} : **outcome** for observation i in cluster g ;

X_{ig} : vector of **covariates** for observation i in cluster g ;

ε_{ig} : **error term** for observation i in cluster g ;

β : **coefficients** of interest;

$g = 1, 2, \dots, \mathbf{G}$.

Clustered errors

Errors **between clusters** are **uncorrelated**:

$$\text{Cor}(\varepsilon_{ig}, \varepsilon_{j\tilde{g}}) = 0$$

Errors **within the same cluster** are (possibly) **correlated**:

$$\text{Cor}(\varepsilon_{ig}, \varepsilon_{jg}) \neq 0$$

Thus, we are relaxing the assumption of **i.i.d. errors**.

Clustered data complicates inference

Problem: CI's assuming i.i.d don't have the right coverage

- Coverage is (typically) less than 95%
- SE are (typically) too small
- May lead to overreject null hypotheses (false positives).

Let's see this in some Monte Carlo simulations.

Linear experimental design

Data generating process:

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

where,

y_{ig} : **outcome** for observation i in cluster g ;

x_{ig}, z_{ig} : **control** variables, $N(0, 3)$ and $\chi^2(7)$ respectively.

Obs = 1000.

g : observations randomly assigned among 100 clusters;

T_g : 33 clusters randomly assigned to treatment ($T_g = 1$)

Clustered errors

Data generating process:

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

where,

ν_g : clustered component of error term, $N(0, 0.5)$;

μ_{ig} : individual component of error term, $N(0, 0.5)$.

\Rightarrow Errors are **correlated** within clusters.

Monte Carlo simulations

Procedure:

1. Simulate the DGP.
2. `regress y x z treat`
3. Store coefficient for `treat`: `[beta]`
4. Check if CI for `treat` contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo simulations

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.008647	.2367399	.1830714	1.775514
contained	1,000	.564	.4961352	0	1

Remarks:

1. Coverage is just 56.4% (vs. 95% nominal size)
2. Estimator is still consistent.
3. CI's are too narrow.

Controlling for cluster dummies is not the solution

Procedure:

1. Simulate the DGP.
2. regress y x z treat i.cvar
3. Store coefficient for treat: [beta]
4. Check if CI for treat contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo controlling for cluster dummies

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.01037	1.482467	-3.426341	5.811284
contained	1,000	.471	.4994081	0	1

Remarks:

1. Coverage fell to 47.1% (vs. 95% nominal size)
2. Inefficient: estimating coefficients of irrelevant variables.

Nonlinear experimental design

Data generating process (Probit model):

$$y_{ig}^* = 1 + x_{ig} + z_{ig} + T_g - \mu_{ig} - \nu_g$$

$$y_{ig} = 1\{y_{ig}^* \geq 0\}$$

where,

y_{ig} : **outcome** for observation i in cluster g ;

x_{ig}, z_{ig} : **control** variables, $N(0, 3)$ and $\chi^2(7)$ respectively.

Obs = 100000

g : observations randomly assigned among 100 clusters;

T_g : 33 clusters randomly assigned to treatment ($T_g = 1$)

Clustered errors

Data generating process:

$$y_{ig}^* = 1 + x_{ig} + z_{ig} + T_g - \mu_{ig} - \nu_g$$

$$y_{ig} = 1\{y_{ig}^* \geq 0\}$$

where,

ν_g : clustered component of error term, $N(0, 0.5)$;

μ_{ig} : individual component of error term, $N(0, 0.5)$.

Monte Carlo simulations

Procedure:

1. Simulate the DGP.
2. `probit y x z treat`
3. Store coefficient for `treat`: `[beta]`
4. Check if CI for `treat` contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo nonlinear experiment

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.022974	.1905835	.3237958	1.550756
contained	1,000	.558	.4968731	0	1

Remarks:

1. Coverage is just 55.8% (vs. 95% nominal size)
2. Estimator is still consistent.
3. CI's are too narrow.

Solutions

1. Cluster-robust variance estimator:

- Option `vce(cluster cvarlist)`
Liang and Zeger (1986)
- Adjust degrees of freedom: `vce(hc2 cvar, dfadjust)` **[Stata 18]**
Bell and McCaffrey (2002)

2. Wild cluster bootstrap **[Stata 18]**

Cameron, Gelbach, and Miller (2008)

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Cluster-robust variance estimator (CRVE)

CI's can be **corrected** using the **CRVE**:

$$\hat{V} = \frac{G(N-1)}{(G-1)(N-k)} (XX')^{-1} \left(\sum_{g=1}^G X'_g \hat{\epsilon}_g \hat{\epsilon}'_g X_g \right) (XX')^{-1}$$

The 95% **corrected CI** for β_k : $\left[\hat{\beta}_k - 1.96 \sqrt{\hat{V}_{k,k}}, \hat{\beta}_k + 1.96 \sqrt{\hat{V}_{k,k}} \right]$

Implementation – option `vce(cluster)`

Example:

```
estimation_command ..., vce(cluster cvarlist)
```

Check availability:

```
help estimation_command
```

Monte Carlo simulations – Linear experimental design

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

Procedure:

1. Simulate the DGP.
2. regress y x z treat, `vce(cluster cvar)`
3. Store coefficient for treatment: [beta]
4. Check if CI for treatment contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo CRVE

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	.9974977	.2236554	.1211772	1.817533
contained	1,000	.959	.1983894	0	1

Remarks:

1. Coverage is 95.9% (vs. 95% nominal size)

Monte Carlo simulations – Probit experimental design

$$y_{ig}^* = 1 + x_{ig} + z_{ig} + T_g - \mu_{ig} - \nu_g$$
$$y_{ig} = 1\{y_{ig}^* \geq 0\}$$

Procedure:

1. Simulate the DGP.
2. `probit y x z treat, vce(cluster cvar)`
3. Store coefficient for treat: `[beta]`
4. Check if CI for treat contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo Probit CRVE

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.022447	.1861081	.3533935	1.591309
contained	1,000	.929	.2569534	0	1

Remarks:

1. Coverage is 92.9% (vs. 95% nominal size)

CRVE typically increases SE, improving CI coverage

Example: Linear regression with and without CRVE
(wage_work.dta)

	wage_CRVE	wage_iid
Job tenure	0.609 (0.020)	0.609 (0.016)
Labor-market condition	-0.049 (0.029)	-0.049 (0.030)
Age in years	0.198 (0.007)	0.198 (0.005)
Intercept	13.506 (0.284)	13.506 (0.212)
Number of observations	1928	1928

Limitations of CRVE

The **CRVE can work well**, but the asymptotics depend on G .

The **CRVE can perform poorly** when:

1. The number of clusters **G is small**.
2. Cluster have very **different sizes**.

Again, let see it in a Monte Carlo simulation.

Linear experimental design with few clusters

Data generating process:

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

where,

y_{ig} : **outcome** for observation i in cluster g ;

x_{ig}, z_{ig} : **control** variables, $N(0, 3)$ and $\chi^2(7)$ respectively.

Obs = 1000.

g : observations randomly assigned among **21 clusters**;

T_g : **7 clusters** randomly assigned to treatment ($T_g = 1$)

Results – Monte Carlo few clusters

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.021723	.8693036	-1.44612	3.697521
contained	1,000	.879	.3262905	0	1

Remarks:

1. Coverage of 87.9% (vs. 95% nominal size)

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Solution 1: Adjusted degrees of freedom (DoF)

Bell and McCaffrey (2002): adjust DoF based on `cvar`

- Improves CI coverage when clusters are very few.

`hc2`: requests more conservative estimator for residuals

Implementation:

```
estimation_command ..., vce(hc2 cvar, dfadjust)
```

Check availability:

```
help estimation_command
```

Let's see it in action in a Monte Carlo simulation.

Back to the linear experimental design (7 clusters)

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

Procedure:

1. Simulate the DGP.
2. regress y x z treat, `vce(hc2 cvar, dfadjust)`
3. Store coefficient for treat: [beta]
4. Check if CI for treat contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Monte Carlo with few clusters, adjusted DoF

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	1.040855	.835802	-1.290347	4.12301
contained	1,000	.978	.1467567	0	1

Remarks:

1. Coverage of 97.8% (vs. 95% nominal size)

Solution 2: The wildbootstrap command

Syntax:

wildbootstrap *estimator depvar [indepvars] [if] [in] [weight] [, options]*

estimator:

- regress
- areg
- xtreg (**fixed-effects** model only; no need to specify fe option)

Quick Start

Estimate the WCB p-value and CI for the coefficient on x_1 in a linear regression of y on x_1 with clusters identified in `cvar`

```
wildbootstrap regress y x1, cluster(cvar)
```

Wild cluster bootstrap algorithm

Consider the model:

$$y = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_G \end{bmatrix} = X\beta + \epsilon = \begin{bmatrix} X_1 \\ X_2 \\ \vdots \\ X_G \end{bmatrix} \begin{bmatrix} \beta_1 \\ \beta_2 \\ \vdots \\ \beta_k \end{bmatrix} + \begin{bmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_G \end{bmatrix}$$

where,

y_g : vector of **outcomes** for cluster g ,

X_g : vector of **covariates** for cluster g ,

ϵ_g : vector of **errors** for cluster g ,

β : **coefficients** of interest.

The wild cluster restricted bootstrap (WCRB)

The **WCRB algorithm** consists of 4 steps:

Suppose, for example, we want to test $\mathbb{H}_0 : \beta_k = 0$.

1. **Re-estimate** the linear model **with the restriction** $\beta_k = 0$.

$\tilde{\beta}$: restricted **estimated coefficients**

$\tilde{\epsilon}$: restricted **estimated residuals**

WCRB algorithm

2. Create a **bootstrap replication b** (repeat B times):

2.1 Generate **random variable** ν_g^b for each cluster g :

- Distribution specified in `errorweight()`

E.g. `rademacher`: -1 or $+1$ with equal probability

2.2 Generate a **new dependent variable** y_{ig}^b :

$$y_{ig}^b = X_{ig}\tilde{\beta} + \tilde{\epsilon}_{ig}\nu_g^b.$$

2.3 **Re-estimate the model** using variable y_{ig}^b on the LHS.

2.4 Calculate the **t-statistic** $t_k^b = \frac{\hat{\beta}_k^b}{\sqrt{\hat{V}_{k,k}^b}}$

$\hat{\beta}_k^b$: **estimated coefficient** in bootstrap replication.

$\hat{V}_{k,k}^b$: **CRVE** for k -th coefficient in the bootstrap replication.

WCRB algorithm

Suppose $\mathbb{H}_A : \beta_k \neq 0$.

3. **Calculate p-values** depending on the the t -statistics' distribution:

- **Symmetric** around 0: $p_S = \frac{1}{B} \sum_{b=1}^B I(|t_k^b| > |t_k|)$

t_k : original sample t-statistic

- **Otherwise:** $p_e = 2 \min(p_1, p_2)$

p_1, p_2 : Bootstrap p-values for **one-sided alternative hypotheses**.

4. Obtain CI's by **inverting the test**. Find t -statistics such that p -value is 0.05.

Back to the linear experimental design (7 clusters)

$$y_{ig} = 1 + x_{ig} + z_{ig} + T_g + \mu_{ig} + \nu_g$$

Procedure:

1. Simulate the DGP.
2. `wildbootstrap reg y x z treat, cluster(cvar)`
3. Store coefficient for treatment: `[beta]`
4. Check if CI for treatment contains 1
5. Repeat 1000 times steps 1-4.
6. Count number of times 1 was contained in CI.

Results – Wild cluster bootstrap

Variable	Obs	Mean	Std. dev.	Min	Max
beta	1,000	.9961465	.4699228	-1.036362	2.653632
contained	1,000	.957	.2029586	0	1

Remarks:

1. Coverage of 95.7% (vs. 95% nominal size)

Example 1: Simple linear regression

```
. use https://www.stata-press.com/data/r18/wagework, replace
(Wages for 20 to 77 year olds, 2013-2016)
```

```
. wildbootstrap regress wage tenure, cluster(personid) rseed(12345)
```

Performing 1,000 replications for p-value for tenure = 0 ...

Computing confidence interval for tenure

Lower bound:10.....20. done (21)

Upper bound:10..... done (17)

Wild cluster bootstrap

Linear regression

Number of obs = 1,928

Number of clusters = 589

Cluster size:

min = 1

avg = 3.3

max = 4

Cluster variable: personid

Error weight: Rademacher

	wage	Estimate	t	p-value	[95% conf. interval]	
constraint						
	tenure = 0	.7807403	27.19	0.000	.7209754	.8368386

Example 1: Using CRVE instead

```
. regress wage tenure, vce(cluster personid)
```

Linear regression

```
Number of obs      =      1,928
F(1, 588)           =      739.36
Prob > F            =      0.0000
R-squared           =      0.4212
Root MSE           =      3.5097
```

(Std. err. adjusted for 589 clusters in personid)

wage	Coefficient	Robust std. err.	t	P> t	[95% conf. interval]	
tenure	.7807403	.028713	27.19	0.000	.7243477	.8371328
_cons	20.89884	.2135686	97.86	0.000	20.47939	21.31829

Remark: Large G and clusters have similar size, $CI_{CRVE} \approx CI_{WCRB}$

Example 2: Few clusters of heterogeneous size

```
. use https://www.stata-press.com/data/r18/nlsw88, replace
(NLSW, 1988 extract)
```

```
. wildbootstrap regress wage tenure, cluster(industry) rseed(12345)
```

Performing 1,000 replications for p-value for tenure = 0 ...

Computing confidence interval for tenure

Lower bound:10.....20..... done (26)

Upper bound:10.....20.... done (24)

Wild cluster bootstrap

Linear regression

Number of obs = 2,217

Number of clusters = 12

Cluster size:

min = 4

avg = 184.8

max = 817

Cluster variable: industry

Error weight: Rademacher

	wage	Estimate	t	p-value	[95% conf. interval]	
constraint						
	tenure = 0	.1830716	6.95	0.000	.1274023	.3258156

Example 2: Using CRVE instead

```
. regress wage tenure, vce(cluster industry)
```

Linear regression

```
Number of obs   =      2,217
F(1, 11)        =      48.30
Prob > F        =      0.0000
R-squared       =      0.0305
Root MSE       =      5.6853
```

(Std. err. adjusted for 12 clusters in industry)

wage	Coefficient	Robust std. err.	t	P> t	[95% conf. interval]	
tenure	.1830716	.026341	6.95	0.000	.1250953	.2410478
_cons	6.710915	.4147967	16.18	0.000	5.797954	7.623877

Remark: Small G and clusters of dissimilar size, $CI_{CRVE} \neq CI_{WCRB}$

Example 3: Two regressors

```
. wildbootstrap regress wage tenure age, cluster(industry) rseed(12345)

Performing 1,000 replications for p-value for tenure = 0 ...
Computing confidence interval for tenure
  Lower bound: .....10.....20.. done (22)
  Upper bound: .....10.....20.... done (24)

Performing 1,000 replications for p-value for age = 0 ...
Computing confidence interval for age
  Lower bound: .....10.....20.... done (24)
  Upper bound: .....10.....20..... done (27)

Wild cluster bootstrap
Linear regression
Cluster variable: industry
Error weight: Rademacher

Number of obs      = 2,217
Number of clusters = 12
Cluster size:
min = 4
avg = 184.8
max = 817
```

	wage	Estimate	t	p-value	[95% conf. interval]	
constraints						
	tenure = 0	.1869715	7.18	0.000	.13478	.3280472
	age = 0	-.0946592	-2.55	0.006	-.2232396	-.0279091

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Conclusion

1. It's crucial to adjust standard errors when dealing with clustered data.
 - CI's can be very misleading otherwise
 - Specially in experimental designs with treatment by clusters.
2. When clusters are many and homogeneous:
 - CRVE: `vce(cluster cvar)`
3. When clusters are few and heterogeneous:
 - Adjust degrees of freedom: `vce(hc2 cvar, dfadjust)`
 - Wild cluster bootstrap: `wildbootstrap`

Learning more...

1. `help` command
 - Access to all our documentation.
2. www.stata.com
 - Access to all our documentation;
 - Frequently asked questions.
3. www.youtube.com/@statacorp/featured
4. tech-support@stata.com
 - Specific questions about our software.

References

1. When should you adjust standard errors for clustering?
 - Abadie, Athey, Imbens, and Wooldridge (NBER, Working paper)
2. How much should we trust differences-in-differences estimates?
 - Esther Duflo (QJE)
3. Bootstrap-based improvements for inference with clustered errors
 - Cameron, Gelbach, and Miller (ReStat)
4. Bias reduction in standard errors for linear regression with multi-stage samples
 - Bell and McCaffrey (Survey Methodology 2002)

Thank you!