

Linear models and related using Stata

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Introduction

- Linear models have been an integral part of Stata since its inception
- Linear models are the most commonly used tools
 - Teaching
 - Build intuition and explore
 - Punching bag
- We are still developing linear regression extension and refinements
- Most current developments that we have incorporated in Stata
- Along the way I will show some simulation and theoretical results

Stata Menu

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	Choice models		800-782-8272 https://www.stata.com			
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		3978				
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	EMM (finite mixture models)	•		~	Value label	
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Linear models and related

3 - StataNow/MP 19.5			Linear regression			
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	Spatial autoregressive models		Fractional polynomials			
	Longitudinal/panel data	•	Quantile regression			
	Multilevel mixed-effects models	•	Errors-in-variables regression			
	Survival analysis	•	Frontier models		Properties	
	Epidemiology and related	•	Panel data	Linear regression (FE, RE, PA, BE, CRE)	\bullet < >	
	Endogenous covariates	•	Mixed-effects linear regression	Lagrange multiplier test for random effects	Variables	
	Sample-selection models	•	Mixed-effects nonlinear regression	Linear regression with AR(1) disturbance (FE, RE)	Name	
	Causal inference/treatment effects	•	Spatial autoregressive models	Random-coefficients regression by GLS	Label	
	SEM (structural equation modeling)	•	Multiple-equation models	Sample-selection model (RE)	Type	
	LCA (latent class analysis)			Dynamic panel data (DPD)	Format Value label	
	FMM (finite mixture models)		Causal inference/treatment effects	Censored outcomes	Notes	
			FMM (finite mixture models)	Difference in differences (DID)	Data	
	IRT (item response theory)					
	IRT (item response theory) Multivariate analysis	•	Lasso inferential models	Models with endogeneity, selection, and treatment	Frame	
	()	•	Lasso inferential models Bayesian regression	Models with endogeneity, selection, and treatment Contemporaneous correlation	 Frame Filename 	

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Linear models and related

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Linear models and related		
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Fractional outcomes	•	StataCorp 4905 Lakeway Drive
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Game plan

- Estimating β of interest hun JABA
- Inference for $\widehat{\beta}$
- Simulation exercises

Estimating β of interest

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High dimensional "fixed effects"

ssc hot

Top 10	packages at	SSC	
	Jun 2025		
Rank	# hits	Package	Author(s)
1	166246.7	estout	Ben Jann
-	116502.7	asdoc 🚽	Attaullah Shah
3	107891.5	outreg2	Roy Wada
4	100261.0	reghdfe	Sergio Correia
5	92930.7	winsor2 🏹	Yujun Lian
6	59048.5	ftools	Sergio Correia
7	54003.7	sum2docx	Chuntao Li, Yuan Xue
8	45016.8	reg2docx	Chuntao Li, Yuan Xue
9	40756.0	coefplot	Ben Jann
10	39717.2	ivreg210	Mark E Schaffer, Steven Stillman, Christopher F Baum

(Click on package name for description)

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A toy model

$$y_i = \beta_0 + \beta_1 x_i + \sum_{k=1}^{1,502} \gamma_k a_{ik} + \sum_{k=1}^{1,124} \theta_k b_{ik} + \varepsilon_i$$

- We are interested in the marginal effect of x on y
- The categorical variables a and b serve as controls
- The 1,501 parameters of *a* and 1,123 parameters for *b* are not relevant.

High dimensional "fixed effects"



Fixed effects vs. categorical variables

$$\mathbf{y}_{it} = \beta_0 + \beta_1 \mathbf{x}_{it} + \gamma \mathbf{a}_{it} + \theta \mathbf{b}_{it} + \alpha_i + \varepsilon_{it}$$

R AMARIA

- *α_i* is a time invariant unobservable
- *α_i* is a fixed effect
- **a**_{it} and **b**_{it} are categorical variables
- In panel data we absorb α_i

syntax areg

• areg ..., absorb(varlist)

- areg treats data as a cross-section
- Any variable that is absorbed is considered to be a regressor
- In other words there are no incidental parameters or fixed effects
- There is no concept of panel delay

syntax areg

- areg ..., absorb(varlist)
- areg treats data as a cross-section
- Any variable that is absorbed is considered to be a regressor
- In other words there are no incidental parameters or fixed effects
- There is no concept of panel data

syntax xtreg, fe

xtreg ..., fe absorb(varlist)

- Any variable that is absorbed in absorb() is considered to be a regressor
- The time invariant heterogeneity (fixed effect) is not estimable
- xtreg treats data as panel data
 - When errors are not assumed to be i.i.d the "fixed-effects" are not accounted in d.f.a
 - Weights are constant within panel
 - Panels should be nested within cluster

Generate a panel data model

```
clear
set seed 111
set obs 10000
qen id = n
generate ai = rnormal(0, 4)
expand 10
bysort id: generate time = _n + 2015
generate a = runiformint(1,1000)
generate b = runiformint(1, 1000)
generate x = rnormal()
generate e = rnormal(0, 6)
generate y = 10 - 10 \times x + a/3000 - b/2000 + e + ai
```

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xtreg vs. areg I

. xtset id time
Panel variable: id (strongly balanced) Time variable: time, 2016 to 2025 Delta: 1 unit
. quietly xtreg y x, absorb(a b) fe . estimates store fe_iid guietly areg y y absorb(a b id)
. quietly areg y x, absorb(a b id) . estimates store areg idd
. etable, estimates (fe iid areg idd) column(command) ///
<pre>> cstat(_r_b, nformat(%9.4g)) cstat(_r_se, nformat(%9.4g))</pre>
xtreg areg
x -10.02 -10.02
(.02024) (.02024)
Intercept 9.898 9.898
(.01893) (.01893)
Number of observations 100000 100000

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xtreg vs. areg II

$$\mathbf{y}_{it} = \boldsymbol{\beta}_0 + \boldsymbol{\beta}_1 \mathbf{x}_{it} + \gamma \mathbf{a}_{it} + \theta \mathbf{b}_{it} + \alpha_i + \varepsilon_{it}$$

- Error term is $\alpha_i + \varepsilon_{it}$
- There is intra panel correlation to be accounted for
- Absorbing α_i does not eliminate all correlation

xtreg vs areg II

- . quietly xtreg y x, absorb(a b) fe vce(robust)
- . estimates store fe_robust
- . quietly areg y x, absorb(a b id) vce(robust)
- . estimates store areg_robust
- etable, estimates(fe_robust areg_robust) column(command)
- > cstat(_r_b,nformat(%9.4g)) cstat(_r_se,nformat(%9.4g))

	xtreg	areg
x	-10.02	-10.02
	(.01999)	(.02012)
Intercept	9.898	9.898
-	(5.05e-06)	(.01893)
Number of observations	100000	100000
		53
		10
		~ ^

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More linear models

- didregress
- xtdidregress
- xthdidregress twfe
- xtreg, cre
- All of this are using _regress except for xtreg, cre

A .

More linear models

- didregress
- xtdidregress
- xthdidregress twfe
- xtreg, cre
- All of this are using _regress except for xtreg, cre

didregress

didregress (y xvars) (treatment), group(group) time(t)

- areg y xvars i.t treatment, absorb(group) /// vce(cluster group)
- treatment indicator for being in a treated group in the period after intervention
- treatment is an interaction of a treated group and post period
- All individuals in a group should have the same treatment status at the same time
- There should be a control group (never issated)
- Standard errors are cluster robust by defaults

- E - N

didregress

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- Standard errors are cluster robust by default

xthdidregress twfe

xthdidregress twfe (y xvars) (d), group(group) time(t)

- regress y on cohort, time, xvars and interactions and their panel level means
- a regression with panel level means of regressors is a Mundlak regression (CRE)
- cohort variable is created (defines first time a group is treated)
- compute contrast of treatment over cohorts at a given time (margins)
- variance covariance matrix is vce (unconditional), takes into account variation in covariates

AKC data

. use akc

(Fictional dog breed and AKC registration data)

. list in 1/15, sepby(breed) noobs abbreviate(20)

registered	best	movie	breed	year
1653	0	0	Affenpinscher	2031
1340	0	0 💊	Affenpinscher	2032
1180	0	0	Affenpinscher	2033
1602	0	0	Affenpinscher	2034
934	0	0	Affenpinscher	2035
497	0	0	Affenpinscher	2036
1395	201	0	Affenpinscher	2037
1656	0	0	Affenpinscher	2038
1663	0	0	Affenpinscher	2039
1166	0	0	Affenpinscher	2040
1341	0	0	Afghan Hound	2031
1398	0	0	Afghan Hound	2032
1544	0	0	Afghan Hound	2033
791	0	0	Afghan Hound	2034
	0	0	Afghan Hound	2035

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xtreg, cre (etable)

. quietly use akc, clear
. quietly xtset breed year
. bysort breed: egen mmovie = mean(movie)
. quietly xtreg registered movie, fe vce(robust)
. estimates store fe
. quietly regress registered movie mmovie, vce(cluster breed)
. estimates store mundlak
. quietly xtreg registered movie, cre vce(robust)
. estimates store cre
. etable, estimates(fe mundlak cre)
▼

	registered register	ed registered
Was a movie protagonist	2185.141 2185.1	
mmovie	(69.375) (69.40 -230.6 (70.93	53
Intercept	1011.490 1028.3	39
Was a movie protagonist	(5.068) (13.16	2185.141 (69.400)
Intercept		1028.339
Was a movie protagonist		(13.168) -230.653 (70.930)
Number of observations	1410 14	10 1410

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xtreg, cre

. quietly use akc, clear
. quietly xtset breed year
. bysort breed: egen mmovie = mean(movie)
. quietly xtreg registered movie, fe vce (robust)
. estimates store fe
. quietly regress registered movie mmovie, vce(cluster breed)
. estimates store mundlak
. quietly xtreg registered movie, cre vce(robust)
. estimates store cre
. etable, estimates(fe mundlak cre) column(estimates) ///
<pre>> cstat(_r_b,nformat(%9.4g)) cstat(_r_se,nformat(%9.4g)) ///</pre>
<pre>> eqrecode(registered = xb registered = xb xit_vars =xb)</pre>

	fe	mundlak	cre
Was a movie protagonist	2185	2185 (69.4)	2185
mmovie	(09.30)	-230.7	(09.4)
Intercept	1011	(70.93)	1028
Was a movie protagonist	(5.068)	. , .	13.17) -230.7
Number of observations	1410	(1410	70.93) 1410

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Inference for $\widehat{\beta}$

What is new in inference

- Wild cluster bootstrap
- HC2 and HC3 for clusters
- HC2 and HC3 for clusters with degrees of freedom adjustment
- HC3 with Hansen degrees of freedom adjustment and rescaling

C.I.C.H.S.H

Multiway clustering

Why do we care

- Bertrand, Duflo, and Mulainathan (2004)
- Cameron and Miller (2015)
- Imbens and Kolesar (2016)
- McKinnon, Nielsen, and Webb (2023)
- Hansen (2024)

Wild cluster bootstrap

wildbootstrap estimator y xvars [if] [in] [weights], opts

- estimator: areg, regress, xtreg
- Computes confidence intervals and p-values
- Does not compute standard errors

Wild cluster bootstrap

- regress y on x₁..., x_k obtain t-statistic
- 2) impose constraint $\beta_i = 0$
- ${f 0}\,$ run regression imposing constraint and obtain ${f ildeeta}$ and ${f ildearepsilon}$

Rech SH

- **(**) generate $\tilde{y} = x\tilde{\beta} + \tilde{\varepsilon}\nu$
- **3** run regression using \tilde{y} and original covariates

wilbootstrap example

```
. wildbootstrap xtreg registered movie, rseed(111)
Panel variable: breed (strongly balanced)
Time variable: year, 2031 to 2040
        Delta: 1 unit
Performing 1,000 replications for p-value for movie = 0 ...
Computing confidence interval for movie
 Lower bound: ......10......20... done (22)
 Wild cluster bootstrap
                                               Number of obs
                                                                 = 1,410
Fixed-effects linear regression
                                               Number of clusters =
                                                                    141
                                               Cluster size:
Cluster variable: breed
                                                              min =
                                                                      10
Error weight: Rademacher
                                                              avg =
                                                                    10.0
                                                              max =
                                                                      10
                                           p-value
             registered
                           Estimate
                                                      [95% conf. interval]
constraint
                                     31.50
                                            0.000
                                                                 2322.348
              movie = 0
                           2185.141
                                                     2029.609
```

$$\widehat{\mathbf{V}}_{hc2cluster} = \left(\mathbf{X}'\mathbf{X}\right)^{-1} \left(\sum_{g=1}^{G} \mathbf{X}'_{g} \mathbf{M}_{g}^{+1/2} \widehat{\epsilon}_{g} \widehat{\epsilon}_{g} \mathbf{M}_{g}^{+1/2} \mathbf{X}_{g}\right) \left(\mathbf{X}'\mathbf{X}\right)^{-1}$$

- $\mathbf{M}_{g}^{+1/2} = \left(\mathbf{I}_{n_{g}} \mathbf{X}_{g} (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}_{g}'\right)^{+1/2}$ is an eliminator matrix and we are using a Moore-Penrose inverse
- Imbens and Kolesar found good properties for cluster case
- It is a generalization of HC2 with a degrees of freedom adjustment of Bell and McCaffrey

HC2 syntax

• estimator ..., vce(hc2 [clustervar], [dfadjust]

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- ▶ areg
- xtreg
- regress

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HC2

. xtdidregress (y x1 x2 i.a i.b) (digt), group(G) time(t) vce(hc2) Computing degrees of freedom ... Treatment and time information Time variable: t Control: digt = 0digt = 1Treatment: Control Treatment Group 46 G Time Minimum Maximum Difference-in-differences regression Number of obs = 30,000 No. of clusters = 50 Data type: Longitudinal (Std. err. adjusted for 50 clusters in id) Robust HC2 Coefficient std. err. P>(t) [95% conf. interval] V ATET diat (1 vs 0) -.0653023 .1189582 -0.550.610 -.385554 .2549495 Note: ATET estimate adjusted for covariates, panel effects, and time effects. . estimates store hc2 э.

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$$\widehat{\mathbf{V}}_{hc2cluster} = (\mathbf{X}'\mathbf{X})^{-1} \left(\sum_{g=1}^{G} \mathbf{X}'_{g} \mathbf{M}_{g}^{+} \widehat{\epsilon}_{g} \widehat{\epsilon}_{g} \mathbf{M}_{g}^{+} \mathbf{X}_{g} \right) (\mathbf{X}'\mathbf{X})^{-1}$$

- $\mathbf{M}_{g}^{+} = \left(\mathbf{I}_{n_{g}} \mathbf{X}_{g} \left(\mathbf{X}'\mathbf{X}\right)^{-1} \mathbf{X}'_{g}\right)^{+}$ is an eliminator matrix and we are using a Moore-Penrose inverse
- Suggested as a possilibity by Cameron and Miller (2015)
- It is a generalization of HC2 with a degrees of freedom adjustment of Bell and McCaffrey

HC3 syntax

- estimator ..., vce(hc3 [clustervar], [dfadjust hansen] hun JAIR AL
 - areq
 - xtreq
 - regress

(I) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1)) < ((1))

HC3

. xtdidregress (y x1 x2 i.a i.b) (digt), group(G) time(t) vce(hc3) Computing degrees of freedom ... Treatment and time information Time variable: t Control: digt = 0digt = 1Treatment: Control Treatment Group 46 G Time Minimum Maximum Difference-in-differences regression Number of obs = 30,000 No. of clusters = 50 Data type: Longitudinal (Std. err. adjusted for 50 clusters in id) Robust HC3 Coefficient std. err. P>(t) [95% conf. interval] V ATET diat (1 vs 0) -.0653023 .1344837 -0.490.653 -.4400695 .309465 Note: ATET estimate adjusted for covariates, panel effects, and time effects. . estimates store hc3 э.

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HC3 with Hansen D.F. Adjustment Example

. xtdidregress (y x1 x2 i.a i.b) (digt), group(G) time(t) vce(hc3, hansen) Computing degrees of freedom ... Treatment and time information Time variable: t Control: digt = 0digt = 1Treatment: Control Treatment Group G 46 Time Minimum Maximum Difference-in-differences regression Number of obs = 30.000 No. of clusters = 50 Data type: Longitudinal (Std. err. adjusted for 50 clusters in id) Robust HC3 P>(t) Coefficient std. err. [95% conf. interval] V ATET digt (1 vs 0) -.0653023 .1344837 -0.490.611 -.3955562 .2649517 Note: ATET estimate adjusted for covariates, panel effects, and time effects. Note: p-values and confidence intervals computed using Hansen adjustment. . estimates store hchansen

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Simulation exercises ALLON WWW HONE THE REAL AND A CHINE AND A

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DID DGP

$$\begin{array}{rcl} y_{igt}\left(0\right) &=& \epsilon_{igt} + u_g + h_{ig}\nu_g \\ \epsilon_{igt} &\sim& N\left(0,1\right) \\ u_g &\sim& N\left(0,1\right) \\ \nu_g &\sim& N\left(0,1\right) \\ h_{ig} &=& 1-2\left(\mathbbm{1}\left\{U(0,1)>.5\right\}\right) \\ y_{igt}\left(1\right) &=& y_{igt}\left(0\right) + \varepsilon_{ig} \\ \varepsilon_{ig} &\sim& N\left(0,1\right) \\ y_{igt} &=& d_{igt}\left(y_{igt}\left(1\right) - y_{igt}\left(0\right)\right) + \mathbf{x}\beta + \alpha_i + \epsilon_{igt} \end{array}$$

Simulation parameters

- Number of clusters: 10 (small) or 50 (large)
- Cluster sizes: homogeneous (uniform distribution) or heterogeneous (truncated beta)
- Panel of 3000 individuals and 10 time periods (fixed across designs)
- 4 treated clusters in all designs
- Treatment happens for all in period 2

Many homogeneous clusters

HC1 HC2 DF HC3 DF HC3 Hansen	.821 .943 .966
HC3 DF	
	966
HC3 Hancon	.300
105 hansen	.948
Wildboot R	.883
Wildboot W	.898

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Few homogeneous clusters

HC1 .886 HC2 DF .923 HC3 DF .954 HC3 Hansen .927 Wildboot R .909 Wildboot W .900	Estimator	Rejection rate
HC3 DF .954 HC3 Hansen .927 Wildboot R .909	HC1	.886
HC3 Hansen .927 Wildboot R .909	HC2 DF	.923
Wildboot R .909	HC3 DF	.954
	HC3 Hansen	.927
Wildboot W .900	Wildboot R	.909
	Wildboot W	.900

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Many heterogeneous clusters

HC1 .871 HC2 DF .934 HC3 DF .956 HC3 Hansen .938 Wildboot R .919	Estimator	Rejection rate
HC3 DF .956 HC3 Hansen .938 Wildboot R .919	HC1	.871
HC3 Hansen .938 Wildboot R .919	HC2 DF 🍌	.934
Wildboot R .919	HC3 DF	.956
	HC3 Hansen	.938
	Wildboot R	.919
Wildboot W .913	Wildboot W	.913

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Few heterogeneous clusters

HC1 .890 HC2 DF .913 HC3 DF .938 HC3 Hansen .918 Wildboot B .906	Estimator	Rejection rate
HC3 DF .938 HC3 Hansen .918	HC1	.890
HC3 Hansen .918	HC2 DF 🍌	.913
	HC3 DF	.938
Wildboot B 906	HC3 Hansen	.918
1000	Wildboot R	.906
Wildboot W .900	Wildboot W	.900

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Takeaways

- Default cluster robust standard errors tend to under-reject
- HC2 and HC3 standard errors with degrees of freedom correction perform well
- HC3 with Hansed D.F. correction has a similar behavior to other D.F. adjustment methods

Conclusion

- Linear models and related have new and important features in the last two releases
- We are still working and improving to add more features both in estimation and inference
- We have added the absorb () option to areg and xtreg, fe with important interpretation differences
- HC2 and HC3 estimator with degrees of freedom seem to be a good alternative when there are few and beterogeneous clusters

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- We are still working and improving to add more features both in estimation and inference
- We have added the absorb() option to areg and xtreg, fe with important interpretation differences
- HC2 and HC3 estimator with degrees of freedom seem to be a good alternative when there are few and heterogeneous clusters