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## Postestimation commands

The following postestimation commands are of special interest after `xtreg`:

Command	Description
<a href="#">xttest0</a>	Breusch and Pagan LM test for random effects
<a href="#">estat mundlak</a>	Mundlak specification test

The following standard postestimation commands are also available:

Command	Description
<a href="#">contrast</a>	contrasts and ANOVA-style joint tests of parameters
* <a href="#">estat ic</a>	Akaike's, consistent Akaike's, corrected Akaike's, and Schwarz's Bayesian information criteria (AIC, CAIC, AICc, and BIC, respectively)
<a href="#">estat summarize</a>	summary statistics for the estimation sample
<a href="#">estat vce</a>	variance–covariance matrix of the estimators (VCE)
<a href="#">estimates</a>	cataloging estimation results
<a href="#">etable</a>	table of estimation results
† <a href="#">forecast</a>	dynamic forecasts and simulations
<a href="#">hausman</a>	Hausman's specification test
<a href="#">lincom</a>	point estimates, standard errors, testing, and inference for linear combinations of parameters
* <a href="#">lrtest</a>	likelihood-ratio test
<a href="#">margins</a>	marginal means, predictive margins, marginal effects, and average marginal effects
<a href="#">marginsplot</a>	graph the results from margins (profile plots, interaction plots, etc.)
<a href="#">nlcom</a>	point estimates, standard errors, testing, and inference for nonlinear combinations of parameters
<a href="#">predict</a>	linear predictions, residuals, error components
<a href="#">predictnl</a>	point estimates, standard errors, testing, and inference for generalized predictions
<a href="#">pwcompare</a>	pairwise comparisons of parameters
<a href="#">test</a>	Wald tests of simple and composite linear hypotheses
<a href="#">testnl</a>	Wald tests of nonlinear hypotheses

\* `estat ic` and `lrtest` are not appropriate after `xtreg` with the `pa`, `re`, or `cre` option.

† `forecast` is not appropriate with `mi` estimation results.

# predict

## Description for predict

`predict` creates a new variable containing predictions such as fitted values, standard errors, predicted values, linear predictions, and the equation-level score.

## Menu for predict

Statistics > Postestimation

## Syntax for predict

*Fixed-effects (FE) model*

```
predict [type] newvar [if] [in] [, FE_statistic]
```

*Between-effects and GLS and ML random-effects (RE) model*

```
predict [type] newvar [if] [in] [, RE/CRE_statistic]
```

*Correlated random-effects (CRE) model*

```
predict [type] newvar [if] [in] [, RE/CRE_statistic]
```

*Population-averaged (PA) model*

```
predict [type] newvar [if] [in] [, PA_statistic nooffset]
```

<i>FE_statistic</i>	Description
Main	
xb	$\alpha + \mathbf{x}_{it}\beta$ , fitted values; the default
stdp	standard error of the fitted values
ue	$u_i + e_{it}$ , the combined residual
* xbu	$\alpha + \mathbf{x}_{it}\beta + u_i$ , prediction including panel effect
* u	$u_i$ , the fixed error component
* xbd	$\alpha + \mathbf{x}_{it}\beta + d_{\text{absorbvars}}$ , prediction including effects of absorbed variables
* d	$d_{\text{absorbvars}}$ , effects of absorbed variables
* e	$e_{it}$ , the overall error component

Unstarred statistics are available both in and out of sample; type `predict ... if e(sample) ...` if wanted only for the estimation sample. Starred statistics are calculated only for the estimation sample, even when `if e(sample)` is not specified.

<i>RE/CRE_statistic</i>	Description
Main	
xb	$\alpha + \mathbf{x}_{it}\beta$ or $\alpha + \mathbf{x}_{it}\beta + \bar{\mathbf{x}}_i\gamma$ for CRE, fitted values; the default
stdp	standard error of the fitted values
ue	$u_i + e_{it}$ , the combined residual
* xbu	$\alpha + \mathbf{x}_{it}\beta + u_i$ or $\alpha + \mathbf{x}_{it}\beta + \bar{\mathbf{x}}_i\gamma + u_i$ for CRE, prediction including panel effect
* u	$u_i$ , the random-error component
* e	$e_{it}$ , the overall error component

Unstarred statistics are available both in and out of sample; type `predict ... if e(sample) ...` if wanted only for the estimation sample. Starred statistics are calculated only for the estimation sample, even when `if e(sample)` is not specified.

<i>PA_statistic</i>	Description
Main	
mu	predicted value of <i>depvar</i> ; considers the <code>offset()</code>
rate	predicted value of <i>depvar</i>
xb	linear prediction
stdp	standard error of the linear prediction
score	first derivative of the log likelihood with respect to $\mathbf{x}_{it}\beta$

These statistics are available both in and out of sample; type `predict ... if e(sample) ...` if wanted only for the estimation sample.

## Options for predict

### Main

xb calculates the linear prediction. This is the default for all except the population-averaged model. The linear prediction equals  $\alpha + \mathbf{x}_{it}\beta$  for the fixed-, between-, and random-effects models; and equals  $\alpha + \mathbf{x}_{it}\beta + \bar{\mathbf{x}}_i\gamma$  for the correlated random-effects model. Panel means are recalculated when predicting out of sample in the correlated random-effects model.

stdp calculates the standard error of the linear prediction. For the fixed-effects model, this excludes the variance due to uncertainty about the estimate of  $u_i$ .

`mu` and `rate` both calculate the predicted value of *depvar*. `mu` takes into account the `offset()`, and `rate` ignores those adjustments. `mu` and `rate` are equivalent if you did not specify `offset()`. `mu` is the default for the population-averaged model.

`ue` calculates the prediction of  $u_i + e_{it}$ .

`xbu` calculates the linear prediction including the fixed or random component. This prediction equals  $\alpha + \mathbf{x}_{it}\beta + u_i$  for the fixed-, between-, and random-effects models; and equals  $\alpha + \mathbf{x}_{it}\beta + \bar{\mathbf{x}}_i\gamma + u_i$  for the correlated random-effects model.

`u` calculates the prediction of  $u_i$ , the estimated fixed or random effect.

`xbd` calculates the prediction of  $\alpha + \mathbf{x}_{it}\beta + d_{\text{absorbvars}}$ , the prediction including the absorbed variables' effects.

`d` calculates the prediction of  $d_{\text{absorbvars}}$ , the absorbed variables' effects.

`e` calculates the prediction of  $e_{it}$ .

`score` calculates the equation-level score,  $u_{it} = \partial \ln L(\mathbf{x}_{it}\beta) / \partial (\mathbf{x}_{it}\beta)$ .

`nooffset` is relevant only if you specified `offset(varname)` for `xtreg, pa`. It modifies the calculations made by `predict` so that they ignore the offset variable; the linear prediction is treated as  $\mathbf{x}_{it}\beta$  rather than  $\mathbf{x}_{it}\beta + \text{offset}_{it}$ .

# margins

## Description for margins

margins estimates margins of response for fitted values, probabilities, and linear predictions.

## Menu for margins

Statistics > Postestimation

## Syntax for margins

```
margins [marginlist] [ , options ]
margins [marginlist] , predict(statistic ...) [predict(statistic ...) ...] [options]
```

### Fixed-effects (FE) model

FE_statistic	Description
xb	$\alpha + \mathbf{x}_{it}\beta$ , fitted values; the default
stdp	not allowed with margins
ue	not allowed with margins
xbu	not allowed with margins
u	not allowed with margins
xbd	not allowed with margins
d	not allowed with margins
e	not allowed with margins

### Between-effects, GLS and ML random-effects (RE), and correlated random-effects (CRE) model

RE_statistic	Description
xb	$\alpha + \mathbf{x}_{it}\beta$ or $\alpha + \mathbf{x}_{it}\beta + \bar{\mathbf{x}}_i\gamma$ for CRE, fitted values; the default
stdp	not allowed with margins
ue	not allowed with margins
xbu	not allowed with margins
u	not allowed with margins
e	not allowed with margins

### Population-averaged (PA) model

PA_statistic	Description
mu	probability of <i>depvar</i> ; considers the offset()
rate	probability of <i>depvar</i>
xb	linear prediction
stdp	not allowed with margins
<u>score</u>	not allowed with margins

Statistics not allowed with margins are functions of stochastic quantities other than e(b).

For the full syntax, see [\[R\] margins](#).

# xttest0

## Description for xttest0

xttest0, for use after xtreg, re, presents the [Breusch and Pagan \(1980\)](#) Lagrange multiplier test for random effects, a test that  $\text{Var}(\nu_i) = 0$ .

## Menu for xttest0

Statistics > Longitudinal/panel data > Linear models > Lagrange multiplier test for random effects

## Syntax for xttest0

```
xttest0

collect is allowed; see [U] 11.1.10 Prefix commands.
```

# estat mundlak

## Description for estat mundlak

estat mundlak performs a Mundlak specification test, a test that  $\nu_i$  is uncorrelated with the regressors in  $\mathbf{x}_{it}$ . estat mundlak is for use after xtreg with the re, cre, and fe options but not with the absorb() option or the vce(hc2) option.

## Menu for estat

Statistics > Postestimation

## Syntax for estat mundlak

estat mundlak [ , options ]	
options	Description
* reps(#)	perform # bootstrap replications; default is reps(50)
* rseed(#)	set random-number seed to #
*reps() and rseed() allowed only after xtreg, vce(bootstrap).	
collect is allowed; see [U] 11.1.10 Prefix commands.	

## Options for estat mundlak

- reps(#) specifies the number of bootstrap replications to perform when computing the *p*-value for the Mundlak test. This option is allowed only after xtreg, vce(bootstrap).
- rseed(#) sets the random-number seed. This option is allowed only after xtreg, vce(bootstrap).

## Remarks and examples

### ▷ Example 1: Lagrange multiplier test for random effects

Continuing with our random-effects model from [example 4](#) in xtreg, we can see that xttest0 will report a test of  $\nu_i = 0$ . In case we have any doubts, we could type

```
. use https://www.stata-press.com/data/r19/nlswork
(National Longitudinal Survey of Young Women, 14-24 years old in 1968)
. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp
> tenure c.tenure#c.tenure 2.race not_smsa south, re theta
(output omitted)
. xttest0
Breusch and Pagan Lagrangian multiplier test for random effects
      ln_wage[idcode,t] = Xb + u[idcode] + e[idcode,t]
Estimated results:
      _____+-----+-----+-----+-----+-----+-----+-----+
               Var      SD = sqrt(Var)
      _____+-----+-----+-----+-----+-----+-----+-----+
      ln_wage      .2283326      .4778416
      e             .0845002      .2906892
      u             .0665151      .2579053
Test: Var(u) = 0
               chibar2(01) = 14779.98
      Prob > chibar2 = 0.0000
```

◀

### ▷ Example 2: Hausman specification test

More importantly, after xtreg, re estimation, hausman will perform the Hausman specification test. If our model is correctly specified, and if  $\nu_i$  is uncorrelated with  $\mathbf{x}_{it}$ , the (subset of) coefficients that are estimated by the fixed-effects estimator and the same coefficients that are estimated here should not statistically differ:

```
. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp
> tenure c.tenure#c.tenure 2.race not_smsa south, re
(output omitted)
. estimates store random_effects
. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp
> tenure c.tenure#c.tenure 2.race not_smsa south, fe
(output omitted)
```

```
. hausman . random_effects
```

	Coefficients		(b-B) Difference	sqrt(diag(V_b-V_B)) Std. err.
	(b)	(B) random_eff~s		
	.			
age	.0359987	.0368059	-.0008073	.0013177
c.age#c.age	-.000723	-.0007133	-9.68e-06	.0000184
t1l_exp	.0334668	.0290208	.0044459	.001711
c.t1l_exp#				
c.t1l_exp	.0002163	.0003049	-.0000886	.000053
tenure	.0357539	.0392519	-.003498	.0005797
c.tenure#				
c.tenure	-.0019701	-.0020035	.0000334	.0000373
not_smsa	-.0890108	-.1308252	.0418144	.0062745
south	-.0606309	-.0868922	.0262613	.0081345

b = Consistent under H0 and Ha; obtained from **xtreg**.

B = Inconsistent under Ha, efficient under H0; obtained from **xtreg**.

Test of H0: Difference in coefficients not systematic

$$\begin{aligned}\text{chi2}(8) &= (b-B)'[(V_b-V_B)^{-1}](b-B) \\ &= 149.43\end{aligned}$$

Prob > chi2 = 0.0000

We can reject the hypothesis that the coefficients are the same. Before turning to what this means, note that `hausman` listed the coefficients estimated by the two models. It did not, however, list `grade` and `2.race`. `hausman` did not make a mistake; in the Hausman test, we compare only the coefficients estimated by both techniques.

This means that we have an unpleasant choice: we can admit that our model is misspecified—that we have not parameterized it correctly—or we can hold that our specification is correct, in which case the observed differences must be due to the zero correlation of  $\nu_i$  and the  $\mathbf{x}_{it}$  assumption.



## □ Technical note

We can also mechanically explore the underpinnings of the test's dissatisfaction. In the comparison table from `hausman`, it is the coefficients on `not_smsa` and `south` that exhibit the largest differences. In [equation \(1'\)](#) of [XT] **xtreg**, we showed how to decompose a model into within and between effects. Let's do that with these two variables, assuming that changes in the average have one effect, whereas transitional changes have another:



```
. egen avgnsmsa = mean(not_smsa), by(id)
. generate devnsma = not_smsa -avgnsmsa
(8 missing values generated)
. egen avgsouth = mean(south), by(id)
. generate devsouth = south - avgsouth
(8 missing values generated)

. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp tenure c.tenure#
> c.tenure 2.race avgnsm devnsm avgsou devsou
Random-effects GLS regression
Group variable: idcode
R-squared:
    Within = 0.1723
    Between = 0.4809
    Overall = 0.3737
Number of obs      =      28,091
Number of groups   =       4,697
Obs per group:
    min =          1
    avg =         6.0
    max =         15
Wald chi2(12)      =    9319.56
Prob > chi2        =     0.0000
corr(u_i, X) = 0 (assumed)
```

ln_wage	Coefficient	Std. err.	z	P> z	[95% conf. interval]	
grade	.0631716	.0017903	35.29	0.000	.0596627	.0666805
age	.0375196	.0031186	12.03	0.000	.0314072	.043632
c.age#c.age	-.0007248	.00005	-14.50	0.000	-.0008228	-.0006269
ttl_exp	.0286543	.0024207	11.84	0.000	.0239098	.0333989
c.ttl_exp#						
c.ttl_exp	.0003222	.0001162	2.77	0.006	.0000945	.0005499
tenure	.0394423	.001754	22.49	0.000	.0360044	.0428801
c.tenure#						
c.tenure	-.0020081	.0001192	-16.85	0.000	-.0022417	-.0017746
race						
Black	-.0545936	.0102101	-5.35	0.000	-.074605	-.0345821
avgnsmsa	-.1833237	.0109339	-16.77	0.000	-.2047537	-.1618937
devnsma	-.0887596	.0095071	-9.34	0.000	-.1073931	-.070126
avgsouth	-.1011235	.0098789	-10.24	0.000	-.1204858	-.0817611
devsouth	-.0598538	.0109054	-5.49	0.000	-.081228	-.0384797
_cons	.2682987	.0495778	5.41	0.000	.171128	.3654694
sigma_u	.2579182					
sigma_e	.29068923					
rho	.44047745	(fraction of variance due to u_i)				

We will leave the reinterpretation of this model to you, except that if we were really going to sell this model, we would have to explain why the between and within effects are different. Focusing on residence in a non-SMSA, we might tell a story about rural people being paid less and continuing to get paid less when they move to the SMSA. Given our panel data, we could create variables to measure this (an indicator for moved from non-SMSA to SMSA) and to measure the effects. In our assessment of this model, we should think about women in the cities moving to the country and their relative productivity in a bucolic setting.

In any case, the Hausman test now is

```
. estimates store new_random_effects
. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp
> tenure c.tenure#c.tenure 2.race avgnsma devnsma avgsou devsou, fe
(output omitted)
. hausman . new_random_effects
```

	Coefficients		(b-B) Difference	sqrt(diag(V_b-V_B)) Std. err.
	(b) .	(B) new_random-s		
age	.0359987	.0375196	-.0015209	.0013198
c.age#c.age	-.000723	-.0007248	1.84e-06	.0000184
ttl_exp	.0334668	.0286543	.0048124	.0017127
c.ttl_exp#c.ttl_exp	.0002163	.0003222	-.0001059	.0000531
tenure	.0357539	.0394423	-.0036884	.0005839
c.tenure#c.tenure	-.0019701	-.0020081	.000038	.0000377
devnsma	-.0890108	-.0887596	-.0002512	.000683
devsouth	-.0606309	-.0598538	-.0007771	.0007618

b = Consistent under H0 and Ha; obtained from **xtreg**.

B = Inconsistent under Ha, efficient under H0; obtained from **xtreg**.

Test of H0: Difference in coefficients not systematic

```
chi2(8) = (b-B)'[(V_b-V_B)^(-1)](b-B)
= 92.52
```

```
Prob > chi2 = 0.0000
```

We have mechanically succeeded in greatly reducing the  $\chi^2$ , but not by enough. The major differences now are in the age, experience, and tenure effects. We already knew this problem existed because of the ever-increasing effect of experience. More careful parameterization work rather than simply including squares needs to be done.

□

### ► Example 3: Mundlak specification test

Suppose now that the errors in our random-effects model are serially correlated within `idcode`. For most panel-data structures, this is probably a more realistic assumption than assuming i.i.d. errors. We can no longer rely on the Hausman specification test when the errors are correlated. However, we can still conduct a cluster-robust specification test by using `estat mundlak`.

We first refit the random-effects model with cluster-robust standard errors and then use `estat mundlak` to perform the specification test.

```
. xtreg ln_w grade age c.age#c.age ttl_exp c.ttl_exp#c.ttl_exp tenure
> > c.tenure#c.tenure 2.race not_smsa south, re vce(cluster idcode)
(output omitted)
. estat mundlak
Mundlak specification test
H0: Covariates are uncorrelated with unobserved panel-level effects
      chi2(8) = 123.10
Prob > chi2 = 0.0000
Notes: Fixed effects and correlated random effects are
       consistent under H0 and Ha.
       Random effects are efficient under H0.
```

With a Wald test statistic of 123.10 and a  $p$ -value of 0.0000, we reject the null hypothesis that  $\nu_i$  is uncorrelated with the regressors in  $\mathbf{x}_{it}$ . This result implies that the random-effects model as stated is misspecified and that either the model needs to be reparameterized or estimation by correlated random effects or fixed effects should be considered.

◀

## Stored results

`xttest0` stores the following in `r()`:

Scalars

<code>r(lm)</code>	Lagrange multiplier statistic
<code>r(df)</code>	degrees of freedom
<code>r(p)</code>	$p$ -value

`estat mundlak` stores the following in `r()`:

Scalars

<code>r(chi2_mundlak)</code>	Mundlak test $\chi^2$ statistic
<code>r(df_mundlak)</code>	Mundlak test degrees of freedom
<code>r(p_mundlak)</code>	Mundlak test $p$ -value
<code>r(N_reps)</code>	number of complete replications
<code>r(N_clust)</code>	number of clusters

Macros

<code>r(rngstate)</code>	random-number state
<code>r(clustvar)</code>	name of cluster variable

## Methods and formulas

Methods and formulas are presented under the following headings:

*Predictions for fixed-effects model with absorbed variables*  
`xttest0`  
`estat mundlak`

## Predictions for fixed-effects model with absorbed variables

The following uses the notation introduced in [Methods and formulas](#) in [XT] `xtreg`.

Suppose we fit the model

$$y_{it} = \alpha + \mathbf{x}_{it}\beta + \eta_{it} + \epsilon_{it}$$

where  $\eta_{it} = \mathbf{d}_{1(it)}\gamma_1 + \mathbf{d}_{2(it)}\gamma_2 + \cdots + \mathbf{d}_{h(it)}\gamma_h$  and  $\mathbf{d}_{k(it)}$  is an indicator vector for absorbed variable  $k$  in panel  $i$  at time  $t$ . By convention, we define the first absorb variable ( $k = 1$ ) as the panel-ID variable and write  $\eta_{it} = \nu_i + \sum_{k=2}^h \mathbf{d}_{k(it)}\gamma_k = \nu_i + \Delta_{it}$ . Let  $\boldsymbol{\eta}$  be the  $1 \times N$  vector with the values of  $\eta_{it}$  in the sample. Define vectors  $\Delta$  and  $\boldsymbol{\nu}$  similarly.

We are interested in making predictions for the panel effects  $\boldsymbol{\nu}$  (the `u` option), the absorbed variables' effects  $\Delta$  (the `d` option), and the residuals  $\epsilon$  (the `e` option). These postestimation predictions use alternating projection methods to avoid the computational burden of estimating the  $\gamma$ 's. Two alternating projection method algorithms are available: Halperin and Cimmino. For a description of these two algorithms, see [Methods and formulas](#) in [XT] `xtreg`.

Estimates for  $\boldsymbol{\nu}$ ,  $\Delta$ , and the residuals can be extracted from  $\hat{\mathbf{v}} = \mathbf{y} - \mathbf{X}\hat{\beta}$ , where  $\hat{\beta}$  is the estimated regression coefficient vector. Here is the procedure. First, separate the absorbed variables into two sets:  $\iota_1 = \{1\}$ , the panel effect, and  $\iota_2 = \{2, \dots, h\}$ , the remaining absorbed variables. Let  $\text{project}(\hat{\mathbf{v}}, \iota_i)$  denote the projection of vector  $\hat{\mathbf{v}}$  over the variables in set  $\iota_i$ , using either the Halperin or Cimmino algorithm, and let  $\text{p\_error}(\hat{\mathbf{v}}, \iota_i)$  denote the corresponding projection error. Second, initialize the values of the algorithm to  $\tilde{\epsilon}^{(1)} = \hat{\mathbf{v}}$ ,  $\tilde{\boldsymbol{\nu}}^{(1)} = \mathbf{0}$ , and  $\tilde{\Delta}^{(1)} = \mathbf{0}$ . Third and finally, iterate until convergence using the following formulas:

$$\tilde{\boldsymbol{\nu}}^{(j+1)} = \tilde{\boldsymbol{\nu}}^{(j)} + \text{project}(\tilde{\epsilon}^{(j)}, \iota_1)$$

$$\tilde{\epsilon}^{(j+1)} = \text{p\_error}\{\text{p\_error}(\tilde{\epsilon}^{(j)}, \iota_1), \iota_2\}$$

$$\tilde{\Delta}^{(j+1)} = \tilde{\Delta}^{(j)} + \text{project}\{\text{p\_error}(\tilde{\epsilon}^{(j)}, \iota_1), \iota_2\}$$

Convergence is declared at  $j = j$  when the elements of  $\text{project}(\tilde{\epsilon}^{(j)}, \iota_1)$  and  $\text{project}(\tilde{\epsilon}^{(j)}, \iota_2)$  are negligible. At convergence,  $\tilde{\epsilon}^{(j)}$  contains the estimated residuals,  $\tilde{\boldsymbol{\nu}}^{(j)}$  the estimated panel effects, and  $\tilde{\Delta}^{(j)}$  the estimated absorbed variables' effects.

## xttest0

`xttest0` reports the Lagrange multiplier test for random effects developed by [Breusch and Pagan \(1980\)](#) and as modified by [Baltagi and Li \(1990\)](#). The model

$$y_{it} = \alpha + \mathbf{x}_{it}\beta + \nu_i$$

is fit via OLS, and then the quantity

$$\lambda_{\text{LM}} = \frac{(n\bar{T})^2}{2} \left\{ \frac{A_1^2}{(\sum_i T_i^2) - n\bar{T}} \right\}$$

is calculated, where

$$A_1 = 1 - \frac{\sum_{i=1}^n (\sum_{t=1}^{T_i} v_{it})^2}{\sum_i \sum_t v_{it}^2}$$

The Baltagi and Li modification allows for unbalanced data and reduces to the standard formula

$$\lambda_{\text{LM}} = \begin{cases} \frac{nT}{2(T-1)} \left\{ \frac{\sum_i (\sum_t v_{it})^2}{\sum_i \sum_t v_{it}^2} - 1 \right\}^2, & \hat{\sigma}_u^2 \geq 0 \\ 0, & \hat{\sigma}_u^2 < 0 \end{cases}$$

when  $T_i = T$  (balanced data). Under the null hypothesis,  $\lambda_{\text{LM}}$  is distributed as a 50:50 mixture of a point mass at zero and  $\chi^2(1)$ .

## estat mundlak

`estat mundlak` performs a Mundlak specification test to help decide between estimation by random effects or estimation by correlated random effects or fixed effects. The test consists of first fitting a correlated random-effects model, using the same covariates and the same sample as the previously fitted `xtreg` model, and then testing whether the coefficients of the panel means are all jointly equal to 0. See [Mundlak \(1978\)](#), [Wooldridge \(2019\)](#), and `xtreg, cre` in `[XT] xtreg` for more details on correlated random-effects models and the Mundlak specification test. `estat mundlak` is available after `xtreg, re`; `xtreg, fe`; and `xtreg, cre`.

`estat mundlak` uses the same VCE specified in the estimation step to fit the underlying correlated random-effects model. So, for example, if the `vce(cluster clustvar)` option was specified in the estimation step, `estat mundlak` will fit the correlated random-effects model using the same level of clustering for the standard errors. If the `vce(bootstrap)` option was specified in the estimation step, `estat mundlak` computes a bootstrapped  $p$ -value for the Mundlak test following the procedure for Wald-type bootstrap tests in [Hansen \(2022, 294\)](#). You can set the number of bootstrap replications used in this computation with the `reps()` option.

## References

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## Also see

[XT] [xtreg](#) — Linear models for panel data

[U] [20 Estimation and postestimation commands](#)

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