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Description

estat wcorrelation is for use after estimation with menl and mixed.

estat wcorrelation displays the overall correlation matrix for a given cluster calculated on the basis of the design of the random effects and their assumed covariance and the correlation structure of the residuals. This allows for a comparison of different multilevel models in terms of the ultimate within-cluster correlation matrix that each model implies.

Menu for estat

Statistics > Postestimation

Syntax

estat wcorrelation [, *options*]

<i>options</i>	Description
at(<i>at_spec</i>)	specify the cluster for which you want the correlation matrix; default is the first two-level cluster encountered in the data
all	display correlation matrix for all the data
covariance	display the covariance matrix instead of the correlation matrix
list	list the data corresponding to the correlation matrix
nosort	list the rows and columns of the correlation matrix in the order they were originally present in the data
<u>iterate</u> (#)	maximum number of iterations to compute random effects; default is <code>iterate(50)</code> ; only for use after <code>menl</code>
<u>tolerance</u> (#)	convergence tolerance when computing random effects; default is <code>tolerance(1e-6)</code> ; only for use after <code>menl</code>
<u>nrtolerance</u> (#)	scaled gradient tolerance when computing random effects; default is <code>nrtolerance(1e-5)</code> ; only for use after <code>menl</code>
<u>nonrtolerance</u>	ignore the <code>nrtolerance()</code> option; only for use after <code>menl</code>
<u>format</u> (% <i>fmt</i>)	set the display format; default is <code>format(%6.3f)</code>
<u>matlist_options</u>	style and formatting options that control how matrices are displayed

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

Options

`at(at_spec)` specifies the cluster of observations for which you want the within-cluster correlation matrix. `at_spec` is

```
relevel_var = value [, relevel_var = value ...]
```

For example, if you specify

```
. estat wcorrelation, at(school = 33)
```

you get the within-cluster correlation matrix for those observations in school 33. If you specify

```
. estat wcorrelation, at(school = 33 classroom = 4)
```

you get the correlation matrix for classroom 4 in school 33.

If `at()` is not specified, then you get the correlations for the first level-two cluster encountered in the data. This is usually what you want.

`all` specifies that you want the correlation matrix for all the data. This is not recommended unless you have a relatively small dataset or you enjoy seeing large $n \times n$ matrices. However, this can prove useful in some cases.

`covariance` specifies that the within-cluster covariance matrix be displayed instead of the default correlations and standard deviations.

`list` lists the model data for those observations depicted in the displayed correlation matrix. With linear mixed-effects models, this option is also useful if you have many random-effects design variables and you wish to see the represented values of these design variables.

`nosort` lists the rows and columns of the correlation matrix in the order that they were originally present in the data. Normally, `estat wcorrelation` will first sort the data according to level variables, by-group variables, and time variables to produce correlation matrices whose rows and columns follow a natural ordering. `nosort` suppresses this.

`iterate(#)` specifies the maximum number of iterations when computing estimates of the random effects. The default is `iterate(50)`. This option is only for use after `menl`.

`tolerance(#)` specifies a convergence tolerance when computing estimates of the random effects. The default is `tolerance(1e-6)`. This option is only for use after `menl`.

`nrtolerance(#)` and `nonrtolerance` control the tolerance for the scaled gradient when computing estimates of the random effects. These options are only for use after `menl`.

`nrtolerance(#)` specifies the tolerance for the scaled gradient. Convergence is declared when $g(-H^{-1})g'$ is less than `nrtolerance(#)`, where g is the gradient row vector and H is the approximated Hessian matrix from the current iteration. The default is `nrtolerance(1e-5)`.

`nonrtolerance` specifies that the default `nrtolerance()` criterion be turned off.

`format(%fmt)` sets the display format for the standard deviation vector and correlation matrix. The default is `format(%6.3f)`.

`matlist_options` are style and formatting options that control how the matrix (or matrices) is displayed; see [P] [matlist](#) for a list of options that are available.

Remarks and examples

► Example 1: Displaying within-cluster correlations for different clusters

Here we fit a model where different clusters have different within-cluster correlations, and we show how to display them for different clusters. We use the Asian children weight data from [example 6](#) of [\[ME\] mixed](#).

```

use https://www.stata-press.com/data/r19/childweight
(Weight data on Asian children)

. mixed weight age || id: age, covariance(unstructured)

Performing EM optimization ...

Performing gradient-based optimization:
Iteration 0:  Log likelihood = -344.37065
Iteration 1:  Log likelihood = -342.83814
Iteration 2:  Log likelihood = -342.71861
Iteration 3:  Log likelihood = -342.71777
Iteration 4:  Log likelihood = -342.71777

Computing standard errors ...

Mixed-effects ML regression                                Number of obs   =   198
Group variable: id                                       Number of groups =    68
                                                         Obs per group:
                                                         min   =    1
                                                         avg   =   2.9
                                                         max   =    5
                                                         Wald chi2(1)    =  755.27
                                                         Prob > chi2     =  0.0000

Log likelihood = -342.71777

```

weight	Coefficient	Std. err.	z	P> z	[95% conf. interval]	
age	3.459671	.1258878	27.48	0.000	3.212936	3.706407
_cons	5.110496	.149478	34.19	0.000	4.817524	5.403468

Random-effects parameters	Estimate	Std. err.	[95% conf. interval]	
id: Unstructured				
var(age)	.2023928	.12429	.0607393	.6744041
var(_cons)	.0970259	.1108024	.0103473	.9098067
cov(age,_cons)	.1401334	.0566912	.0290206	.2512461
var(Residual)	1.357922	.1650507	1.070075	1.723199

LR test vs. linear model: $\chi^2(3) = 27.38$ Prob > $\chi^2 = 0.0000$

Note: LR test is conservative and provided only for reference.

We use estat wcorrelation to display the within-cluster correlations for the first cluster.

```
. estat wcorrelation, list
Standard deviations and correlations for id = 45:
Standard deviations:
      obs |      1      2      3      4      5
-----|-----
      sd |  1.224  1.314  1.448  1.506  1.771
Correlations:
      obs |      1      2      3      4      5
-----|-----
       1 |  1.000
       2 |  0.141  1.000
       3 |  0.181  0.274  1.000
       4 |  0.193  0.293  0.376  1.000
       5 |  0.230  0.348  0.447  0.477  1.000
Data:
```

	id	weight	age
1.	45	5.171	.136893
2.	45	10.86	.657084
3.	45	13.15	1.21834
4.	45	13.2	1.42916
5.	45	15.88	2.27242

We specified the list option to display the data associated with the cluster. The next cluster in the dataset has ID 258. To display the within-cluster correlations for this cluster, we specify the at () option.

```
. estat wcorrelation, at(id=258) list
Standard deviations and correlations for id = 258:
Standard deviations:
      obs |      1      2      3      4
-----|-----
      sd |  1.231  1.320  1.424  1.782
Correlations:
      obs |      1      2      3      4
-----|-----
       1 |  1.000
       2 |  0.152  1.000
       3 |  0.186  0.270  1.000
       4 |  0.244  0.356  0.435  1.000
Data:
```

	id	weight	age
1.	258	5.3	.19165
2.	258	9.74	.687201
3.	258	9.98	1.12799
4.	258	11.34	2.30527

The within-cluster correlations for this model depend on age. The values for age in the two clusters are different, as are the corresponding within-cluster correlations.



See [example 1](#) of [\[ME\] mixed postestimation](#) for a model fit where each cluster had the same model-implied within-cluster correlations.

Stored results

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

Matrices

<code>r(sd)</code>	standard deviations
<code>r(Corr)</code>	within-cluster correlation matrix
<code>r(Cov)</code>	within-cluster variance–covariance matrix
<code>r(G)</code>	variance–covariance matrix of random effects
<code>r(Z)</code>	model-based design matrix
<code>r(R)</code>	variance–covariance matrix of level-one errors
<code>r(path)</code>	path identifying cluster for which correlation is reported

Results `r(G)`, `r(Z)`, and `r(R)` are available only after `mixed`. Result `r(path)` is available only after `menl`.

Methods and formulas

Methods and formulas are presented under the following headings:

[Linear mixed-effects model](#)

[Nonlinear mixed-effects model](#)

Linear mixed-effects model

A two-level linear mixed model of the form

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

implies the marginal model

$$\mathbf{y}_j = \mathbf{X}_j\boldsymbol{\beta} + \boldsymbol{\epsilon}_j^*$$

where $\boldsymbol{\epsilon}_j^* \sim N(\mathbf{0}, \mathbf{V}_j)$, $\mathbf{V}_j = \mathbf{Z}_j\mathbf{G}\mathbf{Z}_j' + \mathbf{R}$. In a marginal model, the random part is described in terms of the marginal or total residuals $\boldsymbol{\epsilon}_j^*$, and \mathbf{V}_j is the covariance structure of these residuals.

`estat wcorrelation` calculates the marginal covariance matrix $\tilde{\mathbf{V}}_j$ for cluster j and by default displays the results in terms of standard deviations and correlations. This allows for a comparison of different multilevel models in terms of the ultimate within-cluster correlation matrix that each model implies.

Calculation of the marginal covariance matrix extends naturally to higher-level models; see, for example, chapter 4.8 in [West, Welch, and Galecki \(2022\)](#).

Nonlinear mixed-effects model

For nonlinear mixed-effects models, there is no closed-form expression for the marginal covariance matrix $\text{Cov}(\mathbf{y}_j)$. This is because it is expressed in terms of a q -dimensional integral (q is the number of random effects in the model), which, in general, is analytically intractable. Under the linear mixed-effects approximation, the marginal covariance matrix is estimated by $\hat{\mathbf{V}}_j = \hat{\mathbf{Z}}_j\hat{\boldsymbol{\Sigma}}\hat{\mathbf{Z}}_j' + \hat{\sigma}^2\hat{\boldsymbol{\Lambda}}_j$, where $\hat{\mathbf{Z}}_j$, $\hat{\boldsymbol{\Sigma}}$, and $\hat{\boldsymbol{\Lambda}}_j$ are defined in [Methods and formulas](#) of [\[ME\] menl](#).

`estat wcorrelation` calculates the estimated marginal covariance matrix $\widehat{\mathbf{V}}_j$ for cluster j and by default displays the results in terms of standard deviations and correlations.

Under the linear mixed-effects approximation, estimation of the marginal covariance matrix extends naturally to higher-level models; see, for example, chapter 4.8 in [West, Welch, and Galecki \(2022\)](#).

Reference

West, B. T., K. B. Welch, and A. T. Galecki. 2022. *Linear Mixed Models: A Practical Guide Using Statistical Software*. 3rd ed. Boca Raton, FL: CRC Press.

Also see

[ME] [menl](#) — Nonlinear mixed-effects regression

[ME] [mixed](#) — Multilevel mixed-effects linear regression

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

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