

didregress postestimation — Postestimation tools for didregress and xt-didregress

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Postestimation commands

The following postestimation commands are of special interest after `didregress` and `xt-didregress`:

| Command | Description |
|-------------------------------|---|
| <code>estat trendplots</code> | graphical diagnostics for parallel trends |
| <code>estat ptrends</code> | parallel-trends test |
| <code>estat granger</code> | Granger causality test |

The following standard postestimation commands are also available:

| Command | Description |
|------------------------------|---|
| <code>estat summarize</code> | summary statistics for the estimation sample |
| <code>estat vce</code> | variance–covariance matrix of the estimators (VCE) |
| <code>estimates</code> | cataloging estimation results |
| <code>forecast</code> | dynamic forecasts and simulations |
| <code>lincom</code> | point estimates, standard errors, testing, and inference for linear combinations of coefficients |
| <code>nlcom</code> | point estimates, standard errors, testing, and inference for nonlinear combinations of coefficients |
| <code>predict</code> | linear predictions and residuals |
| <code>predictnl</code> | point estimates, standard errors, testing, and inference for generalized predictions |
| <code>test</code> | Wald tests of simple and composite linear hypotheses |
| <code>testnl</code> | Wald tests of nonlinear hypotheses |

predict

Description for predict

`predict` creates a new variable containing predictions such as linear predictors and residuals.

Menu for predict

Statistics > Postestimation

Syntax for predict

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

| <i>statistic</i> | Description |
|------------------|-------------|
|------------------|-------------|

Main

| | |
|------------------------|-------------------------------|
| <code>xb</code> | linear predictor; the default |
| <code>residuals</code> | residuals |

Options for predict

Main

`xb`, the default, calculates the linear predictor. It excludes the effect of the first group or of the panel identifier. All other effects, including the time fixed effects, are included in the linear predictor.

`residuals` calculates the overall residuals. It is the difference of the outcome and the linear predictor, including all group, panel, and time effects. In other words, it is not just the difference of the outcome and the linear predictor.

estat

Description for estat

`estat trendplots` produces two diagnostic plots for assessing the parallel-trends assumption that is required for consistent estimation of the ATET using `didregress` or `xtddidregress`. The first plot consists of two lines showing the mean of the outcome over time for the treatment and the control groups. The second plot augments the DID model to include interactions of time with an indicator of treatment and plots the predicted values of this augmented model for the treatment and control groups. Both plots include a vertical line one period before treatment.

`estat ptrends` performs a test of whether the linear trends in the outcome variable are parallel between control and treatment groups during the pretreatment period.

`estat granger` performs a test of whether treatment effects can be observed in anticipation of the treatment.

Menu for estat

Statistics > Postestimation

Syntax for estat

Graphical diagnostics for parallel trends

```
estat trendplots [ , trend_options plot_options ]
```

Parallel-trends test

```
estat ptrends
```

Granger causality test

```
estat granger
```

| <i>trend_options</i> | Description |
|----------------------|-------------|
|----------------------|-------------|

Plot

| | |
|------------------------------------|--|
| <code>omeans</code> | draw graph showing observed means |
| <code>omeans(plot_options)</code> | draw observed-means graph and affect its rendition |
| <code>ltrends</code> | draw graph showing linear trends |
| <code>ltrends(plot_options)</code> | draw linear-trends graph and affect its rendition |
| <code>notitle</code> | suppress overall title |
| <code>noxline</code> | suppress treatment-time reference line |
| <code>nocommonlegend</code> | display two individual legends |
| <code>legendfrom(#)</code> | specify which legend to use |

| <i>plot_options</i> | Description |
|---|--|
| Plot | |
| <i>cline_options</i> | affect rendition of the plotted trend lines; see [G-3] <i>cline_options</i> |
| <i>line1opts(cline_options)</i> | affect rendition of the line for controls |
| <i>line2opts(cline_options)</i> | affect rendition of the line for treated |
| Y axis, X axis, Titles, Legend, Overall | |
| <i>twoway_options</i> | any options other than <code>by()</code> documented in [G-3] <i>twoway_options</i> |

Options for estat trendplots

Plot

`omeans`, `omeans(plot_options)`, `ltrends`, and `ltrends(plot_options)` specify which graphs are to be included and how they should be individually rendered. The default is `omeans ltrends`, meaning that both graphs are included without any modifications.

`omeans` specifies that the observed-means graph be included. Specifying `omeans` suppresses the linear-trends model graph unless `ltrends` or `ltrends(plot_options)` is also specified.

`omeans(plot_options)` specifies that the observed-means graph be included and affects its rendition. Specifying `omeans(plot_options)` suppresses the linear-trends model graph unless `ltrends` or `ltrends(plot_options)` is also specified.

`ltrends` specifies that the linear-trends model graph be included. Specifying `ltrends` suppresses the observed-means graph unless `omeans` or `omeans(plot_options)` is also specified.

`ltrends(plot_options)` specifies that the linear-trends model graph be included and affects its rendition. Specifying `ltrends(plot_options)` suppresses the observed-means graph unless `omeans` or `omeans(plot_options)` is also specified.

`notitle` suppresses the overall title of the rendered graph.

`noxline` suppresses rendering of the treatment-time reference line.

`nocommonlegend` suppresses the display of one common legend and renders two individual legends.

`legendfrom(#)` specifies which legend to use; the default is `legendfrom(1)`, which refers to the legend of the first plot (observed means). `legendfrom(#)` is not allowed with the `nocommonlegend` option.

cline_options affect the rendition of the plotted trend lines, including their style, size, and color; see [G-3] *cline_options*.

line1opts(cline_options) affect the rendition of the plotted trend lines for the group of controls, including their style, size, and color; see [G-3] *cline_options*.

line2opts(cline_options) affect the rendition of the plotted trend lines for the group of treated, including their style, size, and color; see [G-3] *cline_options*.

Y axis, X axis, Titles, Legend, Overall

twoway_options are any of the options documented in [G-3] *twoway_options*, excluding `by()`. These include options for titling the graph (see [G-3] *title_options*) and for saving the graph to disk (see [G-3] *saving_option*).

Stored results for estat

`estat ptrends` stores the following results for the test of linear trends in `r()`:

Scalars

| | |
|----------------------|--|
| <code>r(N)</code> | number of observations |
| <code>r(F)</code> | test statistic |
| <code>r(df_r)</code> | number of degrees of freedom of the residuals for the F distribution under H_0 |
| <code>r(p)</code> | p -value |
| <code>r(df_m)</code> | number of degrees of freedom of the test for the F distribution under H_0 |

`estat granger` stores the following results for the test of treatment anticipation in `r()`:

Scalars

| | |
|----------------------|--|
| <code>r(N)</code> | number of observations |
| <code>r(F)</code> | test statistic |
| <code>r(df_r)</code> | number of degrees of freedom of the residuals for the F distribution under H_0 |
| <code>r(p)</code> | p -value |
| <code>r(df_m)</code> | number of degrees of freedom of the test for the F distribution under H_0 |

Methods and formulas

The tests performed with `estat ptrends` and `estat granger` are based on augmented difference-in-differences (DID) models. With `estat ptrends`, we augment the DID model with terms that capture the differences in slopes between treated and controls. With `estat granger`, we augment the model by interacting the dummy variable that marks treated observations with dummy variables for time periods prior to the treatment to capture any potential anticipatory treatment effects.

Let's consider the case of panel data for individuals over time, in which individuals belong to a group s . Groups could be states, occupational categories, districts, etc. Let y_{its} be the outcome of individual i , who belongs to group s , at time t , where $i = 1, \dots, N$, $t = 1, \dots, T$, and $s = 1, \dots, S$.

We can write the DID model for such setups as follows:

$$y_{ist} = \gamma_i + \gamma_t + \mathbf{x}_{ist}\boldsymbol{\beta} + D_{st}\delta + \epsilon_{ist}$$

Here γ_i are individual fixed effects, γ_t are time fixed effects, \mathbf{x}_{ist} are covariates, D_{st} is a variable that is 1 if an individual belongs to a group s that is treated at time t and is 0 otherwise, and ϵ_{ist} is an error term. The coefficient δ represents the average treatment effect on the treated (ATET).

To simplify the exposition below, we rewrite the model as follows:

$$\begin{aligned} y_{ist} &= \gamma_i + \gamma_t + \mathbf{x}_{ist}\boldsymbol{\beta} + D_{st}\delta + \epsilon_{ist} \\ y_{ist} &= \text{DID}_{ist} + \epsilon_{ist} \end{aligned} \tag{1}$$

The linear-trends model that is used for the parallel-trends test with `estat ptrends` augments the above model with two more terms. Let $d_{t,0} = 1(d_t = 0)$ be a variable indicating pretreatment time periods, and let $d_{t,1} = 1(d_t = 1)$ be a variable indicating posttreatment time periods. Also, let w_i be a variable that is 1 if the individual belongs to a treated group and is 0 otherwise. The augmentation terms then consist of two 3-way interactions between $d_{t,0}$, w_i , and t , and $d_{t,1}$, w_i , and t :

$$y_{ist} = \text{DID}_{ist} + w_i d_{t,0} t \zeta_1 + w_i d_{t,1} t \zeta_2 + \epsilon_{ist} \tag{2}$$

Under this specification, the coefficient ζ_1 captures the differences in slopes between treatment group and control group in pretreatment periods, while ζ_2 captures the differences in slopes in posttreatment periods. If ζ_1 is 0, the linear trends in the outcome are parallel during pretreatment periods. The same is true for ζ_2 with respect to the posttreatment period; however, posttreatment differences in trends are not relevant for assessing the parallel-trends assumption. `estat ptrends` uses a Wald test of ζ_1 against 0 to assess whether the linear trends are parallel prior to treatment. Thus, the null hypothesis of this test is that the linear trends are parallel.

`estat granger` performs a Granger-type causality test to assess whether treatment effects are observed prior to the treatment. To illustrate this, suppose the treatment took place at time $t = j$. We could express D_{st} as $D_{st} = 1(t \geq j)w_i$. The Granger-type test augments the model with counterfactual treatment-time indicators. For example, if the treatment occurred at time $j - 1$, then we could construct a new treatment as $1(t_{it} \geq j - 1)w_i$, and if we have sufficient time points, we could construct another counterfactual treatment as $1(t_{it} \geq j - 2)w_i$, and so on. These terms are referred to as leads in the DID literature. The model used by `estat granger` uses the model in (1) and augments it with all leads leaving out one for identification purposes. Let J index the time at which the treatment occurs.

$$y_{ist} = \text{DID}_{ist} + \sum_{j=2}^{J-1} 1(t_{it} \geq j)w_i\lambda_j + \nu_{ist}$$

The test result is then obtained by performing a joint Wald test on the coefficients λ_j . Thus, the null hypothesis for this test is that the coefficients in λ_j are jointly 0, which is to say there are no anticipatory effects.

The `estat trendplots` command produces two plots. The first plot is simply plotting the observed means for each treatment group at each point in time. The second plot is based on the model in (2), which is the model used for the parallel-trends test, but this model centers the continuous time variable around its minimum value:

$$y_{ist} = \text{DID}_{ist} + w_i d_{t,0} \{t - \min(t)\} \zeta_1 + w_i d_{t,1} \{t - \min(t)\} \zeta_2 + \mu_{ist}$$

Centering around the minimum time value provides a common reference point at the first observed time point such that deviations from parallelism are easily detectable. The graph then shows the predicted values from this model, evaluated at all observed time points for each of the treatment groups and at the means of the covariates.

While the formulas above are shown for the case of panel data, these methods work the same way for data that consist of repeated cross-sections.

Also see

[TE] **didregress** — Difference-in-differences estimation

[TE] **DID intro** — Introduction to difference-in-differences estimation

[U] **20 Estimation and postestimation commands**