

## Postestimation commands

The following standard postestimation commands are available after `etpoisson`:

Command	Description
<code>contrast</code>	contrasts and ANOVA-style joint tests of parameters
<code>estat ic</code>	Akaike's, consistent Akaike's, corrected Akaike's, and Schwarz's Bayesian information criteria (AIC, CAIC, AICc, and BIC, respectively)
<code>estat summarize</code>	summary statistics for the estimation sample
<code>estat vce</code>	variance–covariance matrix of the estimators (VCE)
<code>estat (svy)</code>	postestimation statistics for survey data
<code>estimates</code>	cataloging estimation results
<code>etable</code>	table of estimation results
* <code>hausman</code>	Hausman's specification test
<code>lincom</code>	point estimates, standard errors, testing, and inference for linear combinations of parameters
* <code>lrtest</code>	likelihood-ratio test
<code>margins</code>	marginal means, predictive margins, marginal effects, and average marginal effects
<code>marginsplot</code>	graph the results from margins (profile plots, interaction plots, etc.)
<code>nlcom</code>	point estimates, standard errors, testing, and inference for nonlinear combinations of parameters
<code>predict</code>	potential-outcome means, observed-outcome means, conditional treatment effects, etc.
<code>predictnl</code>	point estimates, standard errors, testing, and inference for generalized predictions
<code>pwcompare</code>	pairwise comparisons of parameters
<code>suest</code>	seemingly unrelated estimation
<code>test</code>	Wald tests of simple and composite linear hypotheses
<code>testnl</code>	Wald tests of nonlinear hypotheses

\*`hausman` and `lrtest` are not appropriate with `svy` estimation results.

# predict

## Description for predict

predict creates a new variable containing predictions such as counts, conditional treatment effects, probabilities, and linear predictions.

## Menu for predict

Statistics > Postestimation

## Syntax for predict

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

```
predict [type] stub* [if] [in] , scores
```

statistic	Description
Main	
<u>p</u> mean	potential-outcome mean (the predicted count); the default
<u>o</u> mean	observed-outcome mean (the predicted count)
cte	conditional treatment effect at treatment level
pr( <i>n</i> )	probability $\Pr(y_j = n)$
pr( <i>a</i> , <i>b</i> )	probability $\Pr(a \leq y_j \leq b)$
xb	linear prediction
<u>x</u> btreat	linear prediction for treatment equation

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

`pomean`, the default, calculates the potential-outcome mean.

`omean` calculates the observed-outcome mean.

`cte` calculates the treatment effect, the difference of potential-outcome means, conditioned on treatment level.

`pr( $n$ )` calculates the probability  $\Pr(y_j = n)$ , where  $n$  is a nonnegative integer that may be specified as a number or a variable.

`pr( $a, b$ )` calculates the probability  $\Pr(a \leq y_j \leq b)$ , where  $a$  and  $b$  are nonnegative integers that may be specified as numbers or variables;

$b$  missing ( $b \geq .$ ) means  $+\infty$ ;

`pr(20, .)` calculates  $\Pr(y_j \geq 20)$ ;

`pr(20,  $b$ )` calculates  $\Pr(y_j \geq 20)$  in observations for which  $b \geq .$  and calculates

$\Pr(20 \leq y_j \leq b)$  elsewhere.

`pr(. ,  $b$ )` produces a syntax error. A missing value in an observation of the variable  $a$  causes a missing value in that observation for `pr( $a, b$ )`.

`xb` calculates the linear prediction for the dependent count variable, which is  $\mathbf{x}_j\boldsymbol{\beta}$  if neither `offset()` nor `exposure()` was specified;  $\mathbf{x}_j\boldsymbol{\beta} + \text{offset}_j^\beta$  if `offset()` was specified; or  $\mathbf{x}_j\boldsymbol{\beta} + \ln(\text{exposure}_j)$  if `exposure()` was specified.

`xbtreat` calculates the linear prediction for the endogenous treatment equation, which is  $\mathbf{w}_j\boldsymbol{\gamma}$  if `offset()` was not specified in `treat()` and  $\mathbf{w}_j\boldsymbol{\gamma} + \text{offset}_j^\alpha$  if `offset()` was specified in `treat()`.

`nooffset` is relevant only if you specified `offset()` or `exposure()` when you fit the model. It modifies the calculations made by `predict` so that they ignore the offset or exposure variable. `nooffset` removes the offset from calculations involving both the `treat()` equation and the dependent count variable.

`scores` calculates equation-level score variables.

The first new variable will contain  $\partial \ln L / \partial (\mathbf{x}_j\boldsymbol{\beta})$ .

The second new variable will contain  $\partial \ln L / \partial (\mathbf{w}_j\boldsymbol{\gamma})$ .

The third new variable will contain  $\partial \ln L / \partial \text{atanh } \rho$ .

The fourth new variable will contain  $\partial \ln L / \partial \ln \sigma$ .

# margins

## Description for margins

margins estimates margins of response for counts, conditional treatment effects, probabilities, and linear predictions.

## Menu for margins

Statistics > Postestimation

## Syntax for margins

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

statistic	Description
<u>p</u> omean	potential-outcome mean (the predicted count); the default
<u>o</u> mean	observed-outcome mean (the predicted count)
cte	conditional treatment effect at treatment level
pr( <i>n</i> )	probability $\Pr(y_j = n)$
pr( <i>a</i> , <i>b</i> )	probability $\Pr(a \leq y_j \leq b)$
xb	linear prediction
<u>x</u> b <u>t</u> reat	linear prediction for treatment equation

Statistics not allowed with margins are functions of stochastic quantities other than e(b).  
For the full syntax, see [R] margins.

## Remarks and examples

The average treatment effect (ATE) and the average treatment effect on the treated (ATET) are the parameters most frequently estimated by postestimation techniques after etpoisson.

You can use the margins command (see [R] margins) after etpoisson to estimate the ATE or ATET. See example 2 of [CAUSAL] etpoisson for an example of ATE estimation. See example 3 of [CAUSAL] etpoisson for an example of ATET estimation.

See example 1 of [CAUSAL] etpoisson for an example using lincom after etpoisson.

## Methods and formulas

See *Methods and formulas* of [CAUSAL] etpoisson for details.

## Also see

[CAUSAL] [etpoisson](#) — Poisson regression with endogenous treatment effects

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

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