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<b>poisson postestimation</b> — Postestimation to	ls for	poisson
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**Postestimation commands** 

# The following postestimation commands are of special interest after poisson:

Command	Description
estat gof	goodness-of-fit test
lassogof	calculate goodness-of-fit predictions

estat gof is not appropriate with svy estimation results.

The following standard postestimation commands are also available:

Command	Description	
contrast	contrasts and ANOVA-style joint tests of estimates	
estat ic	Akaike's, consistent Akaike's, corrected Akaike's, and Schwarz's Bayesian in formation criteria (AIC, CAIC, AICc, and BIC)	
estat summarize	summary statistics for the estimation sample	
estat vce	variance-covariance matrix of the estimators (VCE)	
estat (svy)	postestimation statistics for survey data	
estimates	cataloging estimation results	
etable	table of estimation results	
*forecast	dynamic forecasts and simulations	
*hausman	Hausman's specification test	
lincom	point estimates, standard errors, testing, and inference for linear combinations of coefficients	
linktest	link test for model specification	
*lrtest	likelihood-ratio test	
margins	marginal means, predictive margins, marginal effects, and average marginal effects	
marginsplot	graph the results from margins (profile plots, interaction plots, etc.)	
nlcom	point estimates, standard errors, testing, and inference for nonlinear combinations of coefficients	
predict	number of events, incidence rates, probabilities, etc.	
predictnl	point estimates, standard errors, testing, and inference for generalized predictions	
pwcompare	pairwise comparisons of estimates	
suest	seemingly unrelated estimation	
test	Wald tests of simple and composite linear hypotheses	
testnl	Wald tests of nonlinear hypotheses	

<sup>\*</sup>forecast, hausman, and lrtest are not appropriate with svy estimation results. forecast is also not appropriate with mi estimation results.

## predict

## **Description for predict**

predict creates a new variable containing predictions such as numbers of events, incidence rates, probabilities, linear predictions, standard errors, and the equation-level score.

## Menu for predict

Statistics > Postestimation

## Syntax for predict

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

statistic	Description
Main	
n	number of events; the default
ir	incidence rate
pr(n)	probability $Pr(y_j = n)$
pr(a,b)	probability $Pr(a \leq y_j \leq b)$
хb	linear prediction
stdp	standard error of the linear prediction
score	first derivative of the log likelihood with respect to $\mathbf{x}_{j}\boldsymbol{\beta}$

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

- n, the default, calculates the predicted number of events, which is  $\exp(\mathbf{x}_j \boldsymbol{\beta})$  if neither offset() nor exposure() was specified when the model was fit;  $\exp(\mathbf{x}_j \boldsymbol{\beta} + \text{offset}_j)$  if offset() was specified; or  $\exp(\mathbf{x}_j \boldsymbol{\beta}) \times \exp(\mathbf{x}_j \boldsymbol{\beta})$  if exposure() was specified.
- ir calculates the incidence rate  $\exp(\mathbf{x}_j \boldsymbol{\beta})$ , which is the predicted number of events when exposure is 1. Specifying ir is equivalent to specifying n when neither offset() nor exposure() was specified when the model was fit.
- 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  $Pr(a \le y_j \le b)$ , where a and b are nonnegative integers that may be specified as numbers or variables;

```
b missing (b \ge .) means +\infty; pr(20,.) calculates \Pr(y_j \ge 20); pr(20,b) calculates \Pr(y_j \ge 20) in observations for which b \ge . and calculates \Pr(20 \le y_j \le 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, which is  $x_i\beta$  if neither offset() nor exposure() was specified;  $\mathbf{x}_i \boldsymbol{\beta} + \text{offset}_i$  if offset() was specified; or  $\mathbf{x}_i \boldsymbol{\beta} + \text{ln}(\text{exposure}_i)$  if exposure() was specified; see nooffset below.

stdp calculates the standard error of the linear prediction.

score calculates the equation-level score,  $\partial \ln L/\partial(\mathbf{x}_i\beta)$ .

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; the linear prediction is treated as  $\mathbf{x}_i \boldsymbol{\beta}$  rather than as  $\mathbf{x}_i \boldsymbol{\beta}$  + offset i or  $\mathbf{x}_i \boldsymbol{\beta}$  + ln(exposure i). Specifying predict ..., nooffset is equivalent to specifying predict ..., ir.

# margins

## Description for margins

margins estimates margins of response for numbers of events, incidence rates, probabilities, and linear predictions.

## Menu for margins

Statistics > Postestimation

## Syntax for margins

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

statistic	Description
n	number of events; the default
ir	incidence rate
pr(n)	probability $Pr(y_j = n)$
pr(a,b)	probability $Pr(a \le y_j \le b)$
хb	linear prediction
stdp	not allowed with margins
<u>sc</u> ore	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.

#### estat

## Description for estat

estat gof performs a goodness-of-fit test of the model. Both the deviance statistic and the Pearson statistic are reported. If the tests are significant, the Poisson regression model is inappropriate.

#### Menu for estat

Statistics > Postestimation

## Syntax for estat

estat gof

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

# Remarks and examples

stata.com

## Example 1

Continuing with example 2 of [R] **poisson**, we use estat gof to determine whether the model fits the data well.

- . use https://www.stata-press.com/data/r18/dollhill3 (Doll and Hill (1966))  $\,$
- . poisson deaths smokes i.agecat, exp(pyears) irr
  (output omitted)
- . estat gof

```
Deviance goodness-of-fit = 12.13237

Prob > chi2(4) = 0.0164

Pearson goodness-of-fit = 11.15533

Prob > chi2(4) = 0.0249
```

The deviance goodness-of-fit test tells us that, given the model, we can reject the hypothesis that these data are Poisson distributed at the 1.64% significance level. The Pearson goodness-of-fit test tells us that we can reject the hypothesis at the 2.49% significance level.

So let us now back up and be more careful. We can most easily obtain the incidence-rate ratios within age categories by using ir; see [R] **Epitab**:

. ir deaths smokes pyears, by(agecat) nohet Stratified incidence-rate analysis

Age category	IRR	[95% conf.	interval]	M-H weight	
35-44 45-54 55-64 65-74 75-84	5.736638 2.138812 1.46824 1.35606 .9047304	1.463557 1.173714 .9863624 .9081925 .6000757	49.40468 4.272545 2.264107 2.096412 1.399687	1.472169 9.624747 23.34176 23.25315 24.31435	(exact) (exact) (exact)
Crude M-H combined	1.719823 1.424682	1.391992 1.154703	2.14353 1.757784		(exact)

We find that the mortality incidence ratios are greatly different within age category, being highest for the youngest categories and actually dropping below 1 for the oldest. (In the last case, we might argue that those who smoke and who have not died by age 75 are self-selected to be particularly robust.)

Seeing this, we will now parameterize the smoking effects separately for each category, although we will begin by constraining the smoking effects on third and fourth age categories to be equivalent:

```
. constraint 1 smokes#3.agecat = smokes#4.agecat
. poisson deaths c.smokes#agecat i.agecat, exposure(pyears) irr constraints(1)
Iteration 0: Log likelihood = -31.95424
Iteration 1: Log likelihood = -27.796801
Iteration 2: Log likelihood = -27.574177
Iteration 3: Log likelihood = -27.572645
Iteration 4: Log likelihood = -27.572645
Poisson regression
                                                       Number of obs =
                                                       Wald chi2(8) = 632.14
Log likelihood = -27.572645
                                                       Prob > chi2
                                                                     = 0.0000
 (1) [deaths]3.agecat#c.smokes - [deaths]4.agecat#c.smokes = 0
```

deaths	IRR	Std. err.	z	P> z	[95% conf.	interval]
agecat#						
c.smokes						
35-44	5.736637	4.181256	2.40	0.017	1.374811	23.93711
45-54	2.138812	.6520701	2.49	0.013	1.176691	3.887609
55-64	1.412229	.2017485	2.42	0.016	1.067343	1.868557
65-74	1.412229	.2017485	2.42	0.016	1.067343	1.868557
75-84	.9047304	.1855513	-0.49	0.625	.6052658	1.35236
agecat						
45 <del>-</del> 54	10.5631	8.067701	3.09	0.002	2.364153	47.19623
55-64	47.671	34.37409	5.36	0.000	11.60056	195.8978
65 <del>-</del> 74	98.22765	70.85012	6.36	0.000	23.89324	403.8244
75-84	199.2099	145.3356	7.26	0.000	47.67693	832.3648
_cons	.0001064	.0000753	-12.94	0.000	.0000266	.0004256
ln(pyears)	1	(exposure)				

Note: \_cons estimates baseline incidence rate.

. estat gof

```
Deviance goodness-of-fit = .0773491
Prob > chi2(1) =
                          0.7809
Pearson goodness-of-fit = .0773885
Prob > chi2(1)
                           0.7809
```

The goodness-of-fit is now small; we are no longer running roughshod over the data. Let us now consider simplifying the model. The point estimate of the incidence-rate ratio for smoking in age category 1 is much larger than that for smoking in age category 2, but the confidence interval for smokes#1.agecat is similarly wide. Is the difference real?

```
. test smokes#1.agecat = smokes#2.agecat
      [deaths]1b.agecat#c.smokes - [deaths]2.agecat#c.smokes = 0
          chi2(1) =
                         1.56
        Prob > chi2 =
                         0.2117
```

The point estimates of the incidence-rate ratio for smoking in the 35-44 age category is much larger than that for smoking in the 45-54 age category, but there is insufficient data, and we may be observing random differences. With that success, might we also combine the smokers in the third and fourth categories with those in the first and second categories?

- . test smokes#2.agecat = smokes#3.agecat, accum
  - [deaths]1b.agecat#c.smokes [deaths]2.agecat#c.smokes = 0
  - [deaths]2.agecat#c.smokes [deaths]3.agecat#c.smokes = 0 (2)

```
chi2(2) =
                4.73
Prob > chi2 =
                0.0938
```

Combining the first four categories may be overdoing it—the 9.38% significance level is enough to stop us, although others may disagree.

Thus, we now fit our final model:

- . constraint 2 smokes#1.agecat = smokes#2.agecat
- . poisson deaths c.smokes#agecat i.agecat, exposure(pyears) irr constraints(1/2)

Iteration 0: Log likelihood = -31.550722Iteration 1: Log likelihood = -28.525057 Iteration 2: Log likelihood = -28.514535 Iteration 3: Log likelihood = -28.514535

Poisson regression

Number of obs = Wald chi2(7) = 642.25

Log likelihood = -28.514535

- Prob > chi2 = 0.0000[deaths]3.agecat#c.smokes - [deaths]4.agecat#c.smokes = 0
- [deaths]1b.agecat#c.smokes [deaths]2.agecat#c.smokes = 0 (2)

deaths	IRR	Std. err.	z	P> z	[95% conf.	interval]
agecat#						
c.smokes						
35 <del>-</del> 44	2.636259	.7408403	3.45	0.001	1.519791	4.572907
45 <del>-</del> 54	2.636259	.7408403	3.45	0.001	1.519791	4.572907
55 <del>-</del> 64	1.412229	.2017485	2.42	0.016	1.067343	1.868557
65-74	1.412229	.2017485	2.42	0.016	1.067343	1.868557
75 <del>-</del> 84	.9047304	.1855513	-0.49	0.625	.6052658	1.35236
agecat						
45 <del>-</del> 54	4.294559	.8385329	7.46	0.000	2.928987	6.296797
55 <del>-</del> 64	23.42263	7.787716	9.49	0.000	12.20738	44.94164
65 <del>-</del> 74	48.26309	16.06939	11.64	0.000	25.13068	92.68856
75-84	97.87965	34.30881	13.08	0.000	49.24123	194.561
_cons	.0002166	.0000652	-28.03	0.000	.0001201	.0003908
<pre>ln(pyears)</pre>	1	(exposure)				

Note: \_cons estimates baseline incidence rate.

The above strikes us as a fair representation of the data. The probabilities of observing the deaths seen in these data are estimated using the following predict command:

- . predict p, pr(0, deaths)
- . list deaths p

	deaths	р
1. 2. 3. 4. 5.	32 104 206 186 102	.6891766 .4456625 .5455328 .4910622 .5263011
6. 7. 8. 9.	2 12 28 28 31	.227953 .7981917 .4772961 .6227565 .5475718

The probability  $Pr(y \le \text{deaths})$  ranges from 0.23 to 0.80.

Stored results

estat gof after poisson stores the following in r():

Scalars degrees of freedom (Pearson and deviance) r(df)  $\chi^2$  (Pearson)  $\chi^2$  (deviance) p-value for  $\chi^2$  test (Pearson) p-value for  $\chi^2$  test (deviance) r(chi2\_p) r(chi2\_d)  $r(p_p)$  $r(p_d)$ 

## Methods and formulas

In the following, we use the same notation as in [R] **poisson**.

The equation-level score is given by

$$\operatorname{score}(\mathbf{x}\boldsymbol{\beta})_j = y_j - e^{\xi_j}$$

The deviance (D) and Pearson (P) goodness-of-fit statistics are given by

$$\ln L_{\text{max}} = \sum_{j=1}^{n} w_j \left[ y_j \{ \ln(y_j) - 1 \} - \ln(y_j!) \right]$$

$$\chi_D^2 = -2 \{ \ln L - \ln L_{\text{max}} \}$$

$$\chi_P^2 = \sum_{j=1}^{n} \frac{w_j (y_j - e^{\xi_j})^2}{e^{\xi_j}}$$

## Reference

Manjón, M., and O. Martínez. 2014. The chi-squared goodness-of-fit test for count-data models. *Stata Journal* 14: 798-816.

## Also see

- [R] **poisson** Poisson regression
- [LASSO] lassogof Goodness of fit after lasso for prediction
- [U] 20 Estimation and postestimation commands

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