Title

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poisson postestimation - Postestimation tools for poisson

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Postestimation commands

The following postestimation commands are of special interest after poisson:

estat gofgoodness-of-fit testlassogofcalculate goodness-of-fit predictions	Command	Description
lassogof calculate goodness-of-fit predictions	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 information 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

*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(<i>n</i>)	probability $\Pr(y_i = n)$	
pr(<i>a</i> , <i>b</i>)	probability $\Pr(a \le y_j \le b)$	
xb	linear prediction	
stdp	standard error of the linear prediction	
score	first derivative of the log likelihood with respect to $\mathbf{x}_{i}\boldsymbol{\beta}$	

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 $\exp(\mathbf{x}_j\beta)$ if neither offset() nor exposure() was specified when the model was fit; $\exp(\mathbf{x}_j\beta + \text{offset}_j)$ if offset() was specified; or $\exp(\mathbf{x}_j\beta) \times \exp(\sup_i \beta)$ if exposure() was specified.
- ir calculates the incidence rate $\exp(\mathbf{x}_j\beta)$, 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 $\mathbf{x}_j\beta$ if neither offset() nor exposure() was specified; $\mathbf{x}_j\beta$ + offset_j if offset() was specified; or $\mathbf{x}_j\beta$ + ln(exposure_j) if exposure() was specified; see nooffset below.

stdp calculates the standard error of the linear prediction.

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

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 $\mathbf{x}_j\beta$ rather than as $\mathbf{x}_j\beta$ +offset_j or $\mathbf{x}_j\beta$ +ln(exposure_j). 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(<i>n</i>)	probability $Pr(y_j = n)$
pr(<i>a</i> , <i>b</i>)	probability $Pr(a \le y_j \le b)$
xb	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

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Example 1

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

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
```

.t	M - H weight	interval]	[95% conf.	IRR	Age category
9 (exact) 7 (exact) 6 (exact) 5 (exact) 5 (exact)	9.624747 23.34176 23.25315	49.40468 4.272545 2.264107 2.096412 1.399687	1.463557 1.173714 .9863624 .9081925 .6000757	5.736638 2.138812 1.46824 1.35606 .9047304	35-44 45-54 55-64 65-74 75-84
- (exact)		2.14353 1.757784	1.391992 1.154703	1.719823 1.424682	Crude M-H combined

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 = -27.572645
Iteration 3: Log likelihood = -27.572645
Iteration 4: Log likelihood = -27.572645
Poisson regression Number of obs = 10
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						
45-54	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
65-74	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 ln(pyears)	.0001064 1	.0000753 (exposure)	-12.94	0.000	.0000266	.0004256

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?

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
 (1)
       [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 = -31.550722
Iteration 1: Log likelihood = -28.525057
Iteration 2: Log likelihood = -28.514535
Iteration 3: Log likelihood = -28.514535
Poisson regression
                                                        Number of obs =
                                                                            10
                                                        Wald chi2(7) = 642.25
```

Log likelihood = -28.514535

```
= 0.0000
Prob > chi2
```

(1) [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						
35-44	2.636259	.7408403	3.45	0.001	1.519791	4.572907
45-54	2.636259	.7408403	3.45	0.001	1.519791	4.572907
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						
45-54	4.294559	.8385329	7.46	0.000	2.928987	6.296797
55-64	23.42263	7.787716	9.49	0.000	12.20738	44.94164
65-74	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
ln(pyears)	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.	32 104 206	.6891766 .4456625 .5455328
4. 5.	186 102	.4910622
6. 7. 8. 9. 10.	2 12 28 28 31	.227953 .7981917 .4772961 .6227565 .5475718
	-	

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

Stored results

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

Scalars

r(df)	degrees of freedom (Pearson and deviance)
r(chi2_p)	χ^2 (Pearson)
r(chi2_d)	χ^2 (deviance)
r(p_p)	<i>p</i> -value for χ^2 test (Pearson)
r(p_d)	<i>p</i> -value for χ^2 test (deviance)

Methods and formulas

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

The equation-level score is given by

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

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

$$\begin{aligned} \ln L_{\max} &= \sum_{j=1}^{n} w_j \left[y_j \{ \ln(y_j) - 1 \} - \ln(y_j!) \right] \\ \chi_D^2 &= -2 \{ \ln L - \ln L_{\max} \} \\ \chi_P^2 &= \sum_{j=1}^{n} \frac{w_j (y_j - e^{\xi_j})^2}{e^{\xi_j}} \end{aligned}$$

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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