1 poisson: Poisson Regression for Event Count Dependent Variables

Use the Poisson regression model if the observations of your dependent variable represents the number of independent events that occur during a fixed period of time (see the negative binomial model, Section ??, for over-dispersed event counts.).

1.0.1 Syntax

```
> z.out <- zelig(Y ~ X1 + X2, model = "poisson", data = mydata)
> x.out <- setx(z.out)
> s.out <- sim(z.out, x = x.out)</pre>
```

1.0.2 Additional Inputs

In addition to the standard inputs, zelig() takes the following additional options for poisson regression:

• robust: defaults to FALSE. If TRUE is selected, zelig() computes robust standard errors via the sandwich package (see [7]). The default type of robust standard error is heteroskedastic and autocorrelation consistent (HAC), and assumes that observations are ordered by time index.

In addition, robust may be a list with the following options:

- method: Choose from
 - * "vcovHAC": (default if robust = TRUE) HAC standard errors.
 - * "kernHAC": HAC standard errors using the weights given in [1].
 - * "weave": HAC standard errors using the weights given in [3].
- order.by: defaults to NULL (the observations are chronologically ordered as in the original data). Optionally, you may specify a vector of weights (either as order.by = z, where z exists outside the data frame; or as order.by = z, where z is a variable in the data frame). The observations are chronologically ordered by the size of z.
- ...: additional options passed to the functions specified in method.
 See the sandwich library and [7] for more options.

1.0.3 Example

Load sample data:

> data(sanction)

Estimate Poisson model:

```
> z.out <- zelig(num ~ target + coop, model = "poisson", data = sanction)
```

> summary(z.out)

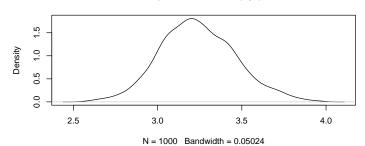
Set values for the explanatory variables to their default mean values:

> x.out <- setx(z.out)

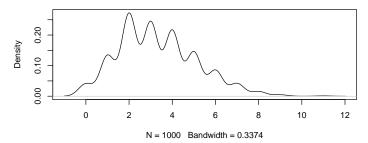
Simulate fitted values:

- > s.out <- sim(z.out, x = x.out)
- > summary(s.out)
- > plot(s.out)

Expected Values: E(Y|X)



Predicted Values: Y|X



1.0.4 Model

Let Y_i be the number of independent events that occur during a fixed time period. This variable can take any non-negative integer.

• The Poisson distribution has stochastic component

$$Y_i \sim \text{Poisson}(\lambda_i),$$

where λ_i is the mean and variance parameter.

• The systematic component is

$$\lambda_i = \exp(x_i \beta),$$

where x_i is the vector of explanatory variables, and β is the vector of coefficients.

1.0.5 Quantities of Interest

• The expected value (qi\$ev) is the mean of simulations from the stochastic component,

$$E(Y) = \lambda_i = \exp(x_i \beta),$$

given draws of β from its sampling distribution.

- The predicted value (qi\$pr) is a random draw from the poisson distribution defined by mean λ_i .
- The first difference in the expected values (qi\$fd) is given by:

$$FD = E(Y|x_1) - E(Y \mid x)$$

• In conditional prediction models, the average expected treatment effect (att.ev) for the treatment group is

$$\frac{1}{\sum_{i=1}^{n} t_i} \sum_{i:t=1}^{n} \left\{ Y_i(t_i = 1) - E[Y_i(t_i = 0)] \right\},\,$$

where t_i is a binary explanatory variable defining the treatment $(t_i = 1)$ and control $(t_i = 0)$ groups. Variation in the simulations are due to uncertainty in simulating $E[Y_i(t_i = 0)]$, the counterfactual expected value of Y_i for observations in the treatment group, under the assumption that everything stays the same except that the treatment indicator is switched to $t_i = 0$.

• In conditional prediction models, the average predicted treatment effect (att.pr) for the treatment group is

$$\frac{1}{\sum_{i=1}^{n} t_i} \sum_{i:t_i=1}^{n} \left\{ Y_i(t_i=1) - \widehat{Y_i(t_i=0)} \right\},\,$$

where t_i is a binary explanatory variable defining the treatment $(t_i = 1)$ and control $(t_i = 0)$ groups. Variation in the simulations are due to uncertainty in simulating $Y_i(\widehat{t_i} = 0)$, the counterfactual predicted value of Y_i for observations in the treatment group, under the assumption that everything stays the same except that the treatment indicator is switched to $t_i = 0$.

1.0.6 Output Values

The output of each Zelig command contains useful information which you may view. For example, if you run z.out <- zelig(y ~ x, model = "poisson", data), then you may examine the available information in z.out by using names(z.out), see the coefficients by using z.out\$coefficients, and a default summary of information through summary(z.out). Other elements available through the \$ operator are listed below.

- From the zelig() output object z.out, you may extract:
 - coefficients: parameter estimates for the explanatory variables.
 - residuals: the working residuals in the final iteration of the IWLS fit.
 - fitted.values: a vector of the fitted values for the systemic component λ .
 - linear.predictors: a vector of $x_i\beta$.
 - aic: Akaike's Information Criterion (minus twice the maximized loglikelihood plus twice the number of coefficients).
 - df.residual: the residual degrees of freedom.
 - df.null: the residual degrees of freedom for the null model.
 - zelig.data: the input data frame if save.data = TRUE.
- From summary(z.out), you may extract:
 - coefficients: the parameter estimates with their associated standard errors, p-values, and t-statistics.
 - cov.scaled: a $k \times k$ matrix of scaled covariances.
 - cov.unscaled: a $k \times k$ matrix of unscaled covariances.
- From the sim() output object s.out, you may extract quantities of interest arranged as matrices indexed by simulation × x-observation (for more than one x-observation). Available quantities are:
 - qi\$ev: the simulated expected values given the specified values of x.
 - qi\$pr: the simulated predicted values drawn from the distributions defined by λ_i .
 - qifd: the simulated first differences in the expected values given the specified values of x and x1.
 - qi\$att.ev: the simulated average expected treatment effect for the treated from conditional prediction models.
 - qi\$att.pr: the simulated average predicted treatment effect for the treated from conditional prediction models.

How to Cite the Poisson Regression Model

Kosuke Imai, Olivia Lau, and Gary King. poisson: Poisson Regression for Event Count Dependent Variables, 2011

How to Cite the Zelig Software Package

To cite Zelig as a whole, please reference these two sources:

Kosuke Imai, Gary King, and Olivia Lau. 2007. "Zelig: Everyone's Statistical Software," http://GKing.harvard.edu/zelig.

Imai, Kosuke, Gary King, and Olivia Lau. (2008). "Toward A Common Framework for Statistical Analysis and Development." Journal of Computational and Graphical Statistics, Vol. 17, No. 4 (December), pp. 892-913.

See also

The poisson model is part of the stats package by (author?) [6]. Advanced users may wish to refer to help(glm) and help(family), as well as [5]. Robust standard errors are implemented via the sandwich package by (author?) [7]. Sample data are from [4].

References

- [1] Donald W.K. Andrews. Heteroskedasticity and autocorrelation consistent covariance matrix estimation. *Econometrica*, 59(3):817–858, May 1991.
- [2] Kosuke Imai, Olivia Lau, and Gary King. poisson: Poisson Regression for Event Count Dependent Variables, 2011.
- [3] Thomas Lumley and Patrick Heagerty. Weighted empirical adaptive variance estimators for correlated data regression. *jrssb*, 61(2):459–477, 1999.
- [4] Lisa Martin. Coercive Cooperation: Explaining Multilateral Economic Sanctions. Princeton University Press, 1992. Please inquire with Lisa Martin before publishing results from these data, as this dataset includes errors that have since been corrected.
- [5] Peter McCullagh and James A. Nelder. *Generalized Linear Models*. Number 37 in Monograph on Statistics and Applied Probability. Chapman & Hall, 2nd edition, 1989.
- [6] William N. Venables and Brian D. Ripley. Modern Applied Statistics with S. Springer-Verlag, 4th edition, 2002.
- [7] Achim Zeileis. Econometric computing with hc and hac covariance matrix estimators. *Journal of Statistical Software*, 11(10):1–17, 2004.