

## 0.1 negbin: Negative Binomial Regression for Event Count Dependent Variables

Use the negative binomial regression if you have a count of events for each observation of your dependent variable. The negative binomial model is frequently used to estimate over-dispersed event count models.

### Syntax

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

### Additional Inputs

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

- **robust**: defaults to **FALSE**. If **TRUE** is selected, `zelig()` computes robust standard errors via the `sandwich` package (see Zeileis (2004)). 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 Andrews (1991).
  - \* **"weave"**: HAC standard errors using the weights given in Lumley and Heagerty (1999).
- **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 Zeileis (2004) for more options.

### Example

Load sample data:

```
> data(sanction)
```

Estimate the model:

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

```
> summary(z.out)
```

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

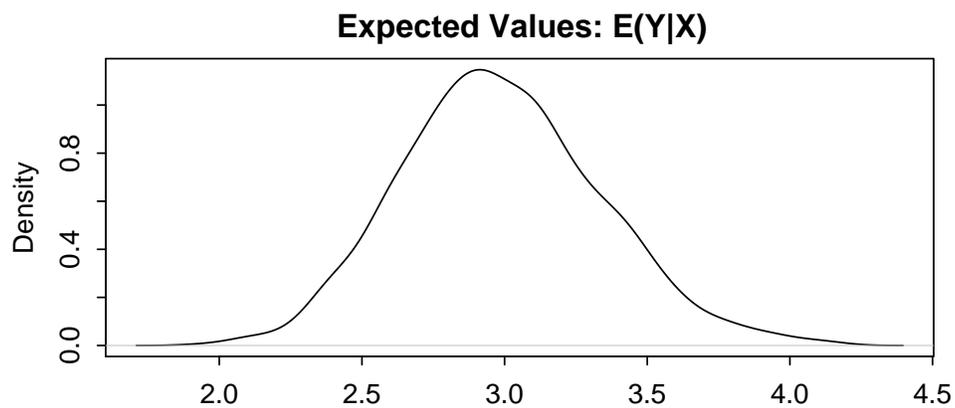
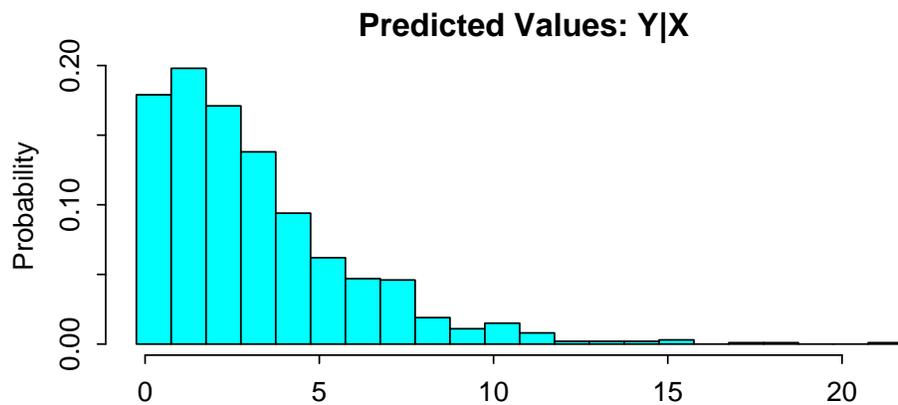
```
> x.out <- setx(z.out)
```

Simulate fitted values:

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

```
> summary(s.out)
```

```
> plot(s.out)
```



## 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 value.

- The negative binomial distribution is derived by letting the mean of the Poisson distribution vary according to a fixed parameter  $\zeta$  given by the Gamma distribution. The *stochastic component* is given by

$$\begin{aligned} Y_i | \zeta_i &\sim \text{Poisson}(\zeta_i \mu_i), \\ \zeta_i &\sim \frac{1}{\theta} \text{Gamma}(\theta). \end{aligned}$$

The marginal distribution of  $Y_i$  is then the negative binomial with mean  $\mu_i$  and variance  $\mu_i + \mu_i^2/\theta$ :

$$\begin{aligned} Y_i &\sim \text{NegBin}(\mu_i, \theta), \\ &= \frac{\Gamma(\theta + y_i)}{y! \Gamma(\theta)} \frac{\mu_i^{y_i} \theta^\theta}{(\mu_i + \theta)^{\theta + y_i}}, \end{aligned}$$

where  $\theta$  is the systematic parameter of the Gamma distribution modeling  $\zeta_i$ .

- The *systematic component* is given by

$$\mu_i = \exp(x_i \beta)$$

where  $x_i$  is the vector of  $k$  explanatory variables and  $\beta$  is the vector of coefficients.

## Quantities of Interest

- The expected values (**qi\$ev**) are simulations of the mean of the stochastic component. Thus,

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

given simulations of  $\beta$ .

- The predicted value (**qi\$pr**) drawn from the distribution defined by the set of parameters  $(\mu_i, \theta)$ .
- The first difference (**qi\$fd**) is

$$\text{FD} = E(Y|x_1) - E(Y|x)$$

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

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

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  $\widehat{Y_i(t_i = 0)}$ , 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  $t_i = 0$ .

## 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 = "negbin", 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.
  - `theta`: the maximum likelihood estimate for the stochastic parameter  $\theta$ .
  - `SE.theta`: the standard error for `theta`.
  - `residuals`: the working residuals in the final iteration of the IWLS fit.
  - `fitted.values`: a vector of the fitted values for the systemic component  $\lambda$ .
  - `linear.predictors`: a vector of  $x_i\beta$ .
  - `aic`: Akaike's Information Criterion (minus twice the maximized log-likelihood 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  $\times$   $\mathbf{x}$ -observation (for more than one  $\mathbf{x}$ -observation). Available quantities are:
    - `qi$ev`: the simulated expected values given the specified values of  $\mathbf{x}$ .
    - `qi$pr`: the simulated predicted values drawn from the distribution defined by  $(\mu_i, \theta)$ .
    - `qi$fd`: the simulated first differences in the simulated expected values given the specified values of  $\mathbf{x}$  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

To cite the *negbin* Zelig model:

Kosuke Imai, Gary King, and Olivia Lau. 2007. “negbin: Negative Binomial Regression for Event Count Dependent Variables,” in Kosuke Imai, Gary King, and Olivia Lau, “Zelig: Everyone’s Statistical Software,” <http://gking.harvard.edu/zelig>.

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>.

Kosuke Imai, Gary King, and Olivia Lau. 2008. “Toward A Common Framework for Statistical Analysis and Development,” *Journal of Computational and Graphical Statistics*, forthcoming, <http://gking.harvard.edu/files/abs/z-abs.shtml>.

## See also

The negative binomial model is part of the MASS package by William N. Venables and Brian D. Ripley (Venables and Ripley 2002). Advanced users may wish to refer to `help(glm.nb)` as well as McCullagh and Nelder (1989). Robust standard errors are implemented via sandwich package by Achim Zeileis (Zeileis 2004). Sample data are from Martin (1992).

# Bibliography

- Andrews, D. W. (1991), “Heteroskedasticity and Autocorrelation Consistent Covariance Matrix Estimation,” *Econometrica*, 59, 817–858.
- Lumley, T. and Heagerty, P. (1999), “Weighted Empirical Adaptive Variance Estimators for Correlated Data Regression,” *jrssb*, 61, 459–477.
- Martin, L. (1992), *Coercive Cooperation: Explaining Multilateral Economic Sanctions*, Princeton University Press, please inquire with Lisa Martin before publishing results from these data, as this dataset includes errors that have since been corrected.
- McCullagh, P. and Nelder, J. A. (1989), *Generalized Linear Models*, no. 37 in Monograph on Statistics and Applied Probability, Chapman & Hall, 2nd ed.
- Venables, W. N. and Ripley, B. D. (2002), *Modern Applied Statistics with S*, Springer-Verlag, 4th ed.
- Zeileis, A. (2004), “Econometric Computing with HC and HAC Covariance Matrix Estimators,” *Journal of Statistical Software*, 11, 1–17.