0.1 exp: Exponential Regression for Duration Dependent Variables

Use the exponential duration regression model if you have a dependent variable representing a duration (time until an event). The model assumes a constant hazard rate for all events. The dependent variable may be censored (for observations have not yet been completed when data were collected).

Syntax

```
> z.out <- zelig(Surv(Y, C) ~ X, model = "exp", data = mydata)
> x.out <- setx(z.out)
> s.out <- sim(z.out, x = x.out)</pre>
```

Exponential models require that the dependent variable be in the form Surv(Y, C), where Y and C are vectors of length n. For each observation i in $1, \ldots, n$, the value y_i is the duration (lifetime, for example), and the associated c_i is a binary variable such that $c_i = 1$ if the duration is not censored (e.g., the subject dies during the study) or $c_i = 0$ if the duration is censored (e.g., the subject is still alive at the end of the study and is know to live at least as long as y_i). If c_i is omitted, all Y are assumed to be completed; that is, time defaults to 1 for all observations.

Input Values

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

- robust: defaults to FALSE. If TRUE, zelig() computes robust standard errors based on sandwich estimators (see Huber (1981) and White (1980)) and the options selected in cluster.
- cluster: if robust = TRUE, you may select a variable to define groups of correlated observations. Let x3 be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then

means that the observations can be correlated within the strata defined by the variable x3, and that robust standard errors should be calculated according to those clusters. If robust = TRUE but cluster is not specified, zelig() assumes that each observation falls into its own cluster.

Example

Attach the sample data:

```
> data(coalition)
```

Estimate the model:

```
> z.out <- zelig(Surv(duration, ciep12) ~ fract + numst2, model = "exp",
+ data = coalition)</pre>
```

View the regression output:

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

Set the baseline values (with the ruling coalition in the minority) and the alternative values (with the ruling coalition in the majority) for X:

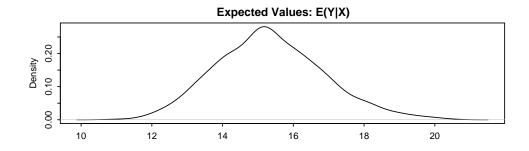
```
> x.low <- setx(z.out, numst2 = 0)
> x.high <- setx(z.out, numst2 = 1)</pre>
```

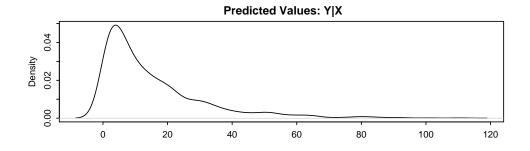
Simulate expected values (qi\$ev) and first differences (qi\$fd):

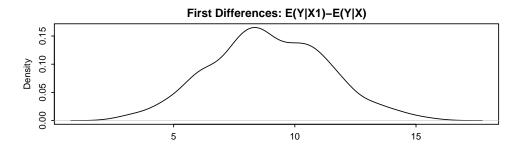
```
> s.out <- sim(z.out, x = x.low, x1 = x.high)
```

Summarize quantities of interest and produce some plots:

- > summary(s.out)
- > plot(s.out)







Model

Let Y_i^* be the survival time for observation i. This variable might be censored for some observations at a fixed time y_c such that the fully observed dependent variable, Y_i , is defined as

$$Y_i = \begin{cases} Y_i^* & \text{if } Y_i^* \le y_c \\ y_c & \text{if } Y_i^* > y_c \end{cases}$$

• The stochastic component is described by the distribution of the partially observed variable Y^* . We assume Y_i^* follows the exponential distribution whose density function is given by

$$f(y_i^* \mid \lambda_i) = \frac{1}{\lambda_i} \exp\left(-\frac{y_i^*}{\lambda_i}\right)$$

for $y_i^* \ge 0$ and $\lambda_i > 0$. The mean of this distribution is λ_i .

In addition, survival models like the exponential have three additional properties. The hazard function h(t) measures the probability of not surviving past time t given survival

up to t. In general, the hazard function is equal to f(t)/S(t) where the survival function $S(t) = 1 - \int_0^t f(s)ds$ represents the fraction still surviving at time t. The cumulative hazard function H(t) describes the probability of dying before time t. In general, $H(t) = \int_0^t h(s)ds = -\log S(t)$. In the case of the exponential model,

$$h(t) = \frac{1}{\lambda_i}$$

$$S(t) = \exp\left(-\frac{t}{\lambda_i}\right)$$

$$H(t) = \frac{t}{\lambda_i}$$

For the exponential model, the hazard function h(t) is constant over time. The Weibull model and lognormal models allow the hazard function to vary as a function of elapsed time (see Section ?? and Section ?? respectively).

• The systematic component λ_i is modeled as

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

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

Quantities of Interest

• The expected values (qi\$ev) for the exponential model are simulations of the expected duration given x_i and draws of β from its posterior,

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

- The predicted values (qi\$pr) are draws from the exponential distribution with rate equal to the expected value.
- The first difference (or difference in expected values, qi\$ev.diff), is

$$FD = E(Y \mid x_1) - E(Y \mid x), \tag{1}$$

where x and x_1 are different vectors of values for the explanatory variables.

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

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

where t_i is a binary explanatory variable defining the treatment $(t_i = 1)$ and control $(t_i = 0)$ groups. When $Y_i(t_i = 1)$ is censored rather than observed, we replace it with

a simulation from the model given available knowledge of the censoring process. Variation in the simulations is due to two factors: uncertainty in the imputation process for censored y_i^* and 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. When $Y_i(t_i = 1)$ is censored rather than observed, we replace it with a simulation from the model given available knowledge of the censoring process. Variation in the simulations is due to two factors: uncertainty in the imputation process for censored y_i^* and 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(Surv(Y, C) ~ X, model = "exp", 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.
 - icoef: parameter estimates for the intercept and scale parameter. While the scale parameter varies for the Weibull distribution, it is fixed to 1 for the exponential distribution (which is modeled as a special case of the Weibull).
 - var: the variance-covariance matrix for the estimates of β .
 - loglik: a vector containing the log-likelihood for the model and intercept only (respectively).
 - linear.predictors: the vector of $x_i\beta$.
 - 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.

- Most of this may be conveniently summarized using summary(z.out). From summary(z.out), you may additionally extract:
 - table: the parameter estimates with their associated standard errors, p-values, and t-statistics. For example, summary(z.out)\$table
- From the sim() output stored in s.out:
- 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 for the specified values of x.
 - qi\$pr: the simulated predicted values drawn from a distribution defined by the expected values.
 - qi\$fd: the simulated first differences between the simulated expected values for 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 *exp* Zelig model:

Kosuke Imai, Gary King, and Oliva Lau. 2007. "exp: Exponential Regression for Duration Dependent Variables" in Kosuke Imai, Gary King, and Olivia Lau, "Zelig: Everyone's Statistical Software," http://gking.harvard.edu/zelig

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See also

The exponential function is part of the survival library by Terry Therneau, ported to R by Thomas Lumley. Advanced users may wish to refer to help(survfit) in the survival library and Venables and Ripley (2002). Sample data are from King et al. (1990).

Bibliography

- Huber, P. J. (1981), Robust Statistics, Wiley.
- King, G., Alt, J., Burns, N., and Laver, M. (1990), "A Unified Model of Cabinet Dissolution in Parliamentary Democracies," *American Journal of Political Science*, 34, 846–871, http://gking.harvard.edu/files/abs/coal-abs.shtml.
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- White, H. (1980), "A Heteroskedasticity-Consistent Covariance Matrix Estimator and a Direct Test for Heteroskedasticity," *Econometrica*, 48, 817–838.