0.1 gamma.mixed: Mixed effects gamma regression

Use generalized multi-level linear regression if you have covariates that are grouped according to one or more classification factors. Gamma regression models a continuous, positive dependent variable.

While generally called multi-level models in the social sciences, this class of models is often referred to as mixed-effects models in the statistics literature and as hierarchical models in a Bayesian setting. This general class of models consists of linear models that are expressed as a function of both *fixed effects*, parameters corresponding to an entire population or certain repeatable levels of experimental factors, and *random effects*, parameters corresponding to individual experimental units drawn at random from a population.

Syntax

```
z.out <- zelig(formula= y ~ x1 + x2 + tag(z1 + z2 | g),
               data=mydata, model="gamma.mixed")

z.out <- zelig(formula= list(mu=y ~ xl + x2 + tag(z1, delta | g),
               delta= ~ tag(w1 + w2 | g)), data=mydata, model="gamma.mixed")
```

Inputs

`zelig()` takes the following arguments for `mixed`:

- `formula`: a two-sided linear formula object describing the systematic component of the model, with the response on the left of a ~ operator and the fixed effects terms, separated by `+` operators, on the right. Any random effects terms are included with the notation `tag(z1 + ... + zn | g)` with `z1 + ... + zn` specifying the model for the random effects and `g` the grouping structure. Random intercept terms are included with the notation `tag(1 | g)`.
  Alternatively, `formula` may be a list where the first entry, `mu`, is a two-sided linear formula object describing the systematic component of the model, with the repsonse on the left of a ~ operator and the fixed effects terms, separated by `+` operators, on the right. Any random effects terms are included with the notation `tag(z1, delta | g)` with `z1` specifying the individual level model for the random effects, `g` the grouping structure and `delta` references the second equation in the list. The `delta` equation is one-sided linear formula object with the group level model for the random effects on the right side of a ~ operator. The model is specified with the notation `tag(w1 + ... + wn | g)` with `w1 + ... + wn` specifying the group level model and `g` the grouping structure.

Additional Inputs

In addition, `zelig()` accepts the following additional arguments for model specification:

- `data`: An optional data frame containing the variables named in `formula`. By default, the variables are taken from the environment from which `zelig()` is called.
- `method`: a character string. The criterion is always the log-likelihood but this criterion does not have a closed form expression and must be approximated. The default approximation is `"PQL"` or penalized quasi-likelihood. Alternatives are `"Laplace"` or `"AGQ"` indicating the Laplacian and adaptive Gaussian quadrature approximations respectively.
- `na.action`: A function that indicates what should happen when the data contain `NA`s. The default action (`na.fail`) causes `zelig()` to print an error message and terminate if there are any incomplete observations.

Additionally, users may with to refer to `lmer` in the package `lme4` for more information, including control parameters for the estimation algorithm and their defaults.

Examples

1. Basic Example with First Differences

   Attach sample data:

   ```
   > data(coalition2)
   ```

   Estimate model using optional arguments to specify approximation method for the log-likelihood, and the log link function for the Gamma family:

   ```
   > z.out1 <- zelig(duration ~ invest + fract + polar + numst2 +
   +      crisis + tag(1 | country), data = coalition2, model = "gamma.mixed",
   +      method = "PQL", family = Gamma(link = log))
   ```

   Summarize regression coefficients and estimated variance of random effects:

   ```
   > summary(z.out1)
   ```

   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.high <- setx(z.out1, numst2 = 1)
   > x.low <- setx(z.out1, numst2 = 0)
   ```

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

```
> s.out1 <- sim(z.out1, x = x.high, x1 = x.low)
> summary(s.out1)
```

Mixed effects gamma regression Model

Let Y_{ij} be the continuous, positive dependent variable, realized for observation j in group i as y_{ij}, for $i = 1, \ldots, M$, $j = 1, \ldots, n_i$.

- The *stochastic component* is described by a Gamma model with scale parameter α.

$$Y_{ij} \sim \text{Gamma}(y_{ij}|\lambda_{ij}, \alpha)$$

  where

$$Gamma(y_{ij}|\lambda_{ij}, \alpha) = \frac{1}{\alpha^{\lambda_{ij}}\Gamma\lambda_{ij}} y_{ij}^{\lambda_{ij}-1} \exp(-\{\frac{y_{ij}}{\alpha}\})$$

  for $\alpha,\ \lambda_{ij},\ y_{ij}\ > 0$.

- The q-dimensional vector of *random effects*, b_i, is restricted to be mean zero, and therefore is completely characterized by the variance covarance matrix Ψ, a $(q \times q)$ symmetric positive semi-definite matrix.

$$b_i \sim Normal(0, \Psi)$$

- The *systematic component* is

$$\lambda_{ij} \equiv \frac{1}{X_{ij}\beta + Z_{ij}b_i}$$

  where X_{ij} is the $(n_i \times p \times M)$ array of known fixed effects explanatory variables, β is the p-dimensional vector of fixed effects coefficients, Z_{ij} is the $(n_i \times q \times M)$ array of known random effects explanatory variables and b_i is the q-dimensional vector of random effects.

Quantities of Interest

- The predicted values (`qi$pr`) are draws from the gamma distribution for each given set of parameters (α, λ_{ij}), for

$$\lambda_{ij} = \frac{1}{X_{ij}\beta + Z_{ij}b_i}$$

  given X_{ij} and Z_{ij} and simulations of of β and b_i from their posterior distributions. The estimated variance covariance matrices are taken as correct and are themselves not simulated.

- The expected values (qi$ev) are simulations of the mean of the stochastic component given draws of α, β from their posteriors:

$$E(Y_{ij}|X_{ij}) = \alpha\lambda_{ij} = \frac{\alpha}{X_{ij}\beta}.$$

- The first difference (qi$fd) is given by the difference in expected values, conditional on X_{ij} and X'_{ij}, representing different values of the explanatory variables.

$$FD(Y_{ij}|X_{ij}, X'_{ij}) = E(Y_{ij}|X_{ij}) - E(Y_{ij}|X'_{ij})$$

- In conditional prediction models, the average predicted treatment effect (qi$att.pr) for the treatment group is given by

$$\frac{1}{\sum_{i=1}^{M}\sum_{j=1}^{n_i} t_{ij}} \sum_{i=1}^{M} \sum_{j:t_{ij}=1}^{n_i} \{Y_{ij}(t_{ij}=1) - Y_{ij}\widehat{(t_{ij}=0)}\},$$

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

- In conditional prediction models, the average expected treatment effect (qi$att.ev) for the treatment group is given by

$$\frac{1}{\sum_{i=1}^{M}\sum_{j=1}^{n_i} t_{ij}} \sum_{i=1}^{M} \sum_{j:t_{ij}=1}^{n_i} \{Y_{ij}(t_{ij}=1) - E[Y_{ij}(t_{ij}=0)]\},$$

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

Output Values

The output of each Zelig command contains useful information which you may view. You may examine the available information in `z.out` by using `slotNames(z.out)`, see the fixed effect coefficients by using `summary(z.out)@coefs`, and a default summary of information through `summary(z.out)`. Other elements available through the operator are listed below.

- From the `zelig()` output stored in `summary(z.out)`, you may extract:
  - `fixef`: numeric vector containing the conditional estimates of the fixed effects.

- `ranef`: numeric vector containing the conditional modes of the random effects.
- `frame`: the model frame for the model.

- From the `sim()` output stored in `s.out`, you may extract quantities of interest stored in a data frame:
  - `qi$pr`: the simulated predicted values drawn from the distributions defined by the expected values.
  - `qi$ev`: the simulated expected values for the specified values of x.
  - `qi$fd`: the simulated first differences in the expected values for the values specified in x and x1.
  - `qi$ate.pr`: the simulated average predicted treatment effect for the treated from conditional prediction models.
  - `qi$ate.ev`: the simulated average expected treatment effect for the treated from conditional prediction models.

How to Cite

To cite the *gamma.mixed* Zelig model:

> Delia Bailey and Ferdinand Alimadhi. 2007. "gamma.mixed: Mixed effects gamma model" 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`.

> 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

Mixed effects gamma regression is part of `lme4` package by Douglas M. Bates (Bates 2007). For a detailed discussion of mixed-effects models, please see **?**

Bibliography

Bates, D. (2007), *lme4: Fit linear and generalized linear mixed-effects models.*