Package ‘TwoStepCLogit’

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TwoStepCLogit-package  Conditional Logistic Regression: A Two-Step Estimation Method

Description

Conditional logistic regression with longitudinal follow up and individual-level random coefficients: A stable and efficient two-step estimation method (see Ts.estim).

Details

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Author(s)

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References


bison  Bison Dataset

Description

Bison data collected in Prince Albert National Park, Saskatchewan, Canada (Craiu et al. 2011).

Format

A data frame with 16818 observations on the following 10 variables.

Cluster  pair of animals (dyad) ID
Strata   stratum ID
Y        response variable: 1 for visited locations, 0 otherwise
**Details**

This data set was collected in order to study habitat selection by groups of free-ranging bison. For each observed group, two individuals (dyad) equipped with GPS radio-collars were followed simultaneously. A cluster is defined here as a pair of bison. This data set contains 20 clusters. The number of strata per cluster varies between 13 and 345 for a total of 1410 strata. A stratum is composed of two visited GPS locations (one for each individual) gathered at the same time, together with 10 random locations (five drawn within 700 m of each of the two focal bison). Therefore, there are 12 observations per stratum, with 2 cases (Y=1) and 10 controls (Y=0). However, due to problems in the data collection, 17 of the 1410 strata have only 6 observations (1 case and 5 controls).

**References**


**Examples**

```r
# Some descriptive statistics about the data set:
ddim(formula = y ~ strata(Strata) + cluster(Cluster), data = bison)

# Model 1: covariates meadow, biomass and biomass^2
# Random effects in front of biomass and biomass^2
# Main diagonal covariance structure for D
Fit1 <- Ts.estim(formula = Y ~ meadow + biomass + I(biomass^2) + strata(Strata) + cluster(Cluster), data = bison,
                   random = ~ biomass + I(biomass^2), all.m.1=FALSE, D="UN(1)"
)
Fit1

# Model 2: only covariates biomass and biomass^2
# Random effects in front of biomass and biomass^2
# Main diagonal covariance structure for D
Fit2 <- Ts.estim(formula = Y ~ biomass + I(biomass^2) + strata(Strata) +
                   cluster(Cluster), data = bison, all.m.1=FALSE, D="UN(1)"
)
Fit2
# Results reported in Table 2 of Craiu et al. (2011).
```

**Variables**

- **water** land cover indicator covariate: 1 for water, 0 otherwise
- **agric** land cover indicator covariate: 1 for agricultural locations, 0 otherwise
- **forest** land cover indicator covariate: 1 for forests, 0 otherwise
- **meadow** land cover indicator covariate: 1 for meadows, 0 otherwise
- **biomass** continuous covariate: above-ground vegetation biomass index measured (in kg/m^2) only at locations within meadows, 0 otherwise
- **pmeadow** continuous covariate: the proportion of meadow in a circular plot (700 m in radius) centered at the bison’s location
**ddim**

*Data Dimension Statistics*

**Description**

Function that computes dimension statistics for a data set with clusters and strata and its print method.

**Usage**

```r
ddim(formula, data)
```

```r
# S3 method for class 'ddim'
print(x, ...)
```

**Arguments**

- `formula` A formula object, with the response on the left of a `~` operator and, on the right hand side, a strata and a cluster term (ex. `formula = Y ~ strata(var_strata) + cluster(var_cluster)`). The strata and cluster functions (from the package `survival`) are used to identify the stratification and the cluster variables, respectively.
- `data` A data frame (or object coercible by `as.data.frame` to a data frame) containing the variables in the model.
- `x` An object, produced by the `ddim` function, to print.
- `...` Further arguments to be passed to `print.default`.

**Value**

- `sc` The number of strata in each cluster.
- `Ystat` A `data.frame` with `n`, the numbers of observations per stratum ($n_\xi$), and `m`, the sum of the responses per stratum ($m_\xi$).

**See Also**

- `Ts.estim`

**Examples**

```r
dimstat <- ddim(formula = Y ~ strata(Strata) + cluster(Cluster), data = bison)
dimstat
```
Description

Function that computes the two-step estimator proposed in Craiu et al. (2011) and its print method.

Usage

\begin{verbatim}
Ts.estim(formula, data, random, all.m.1 = FALSE, D = "UN(1)",
         itermax = 2000, tole = 1e-06)
\end{verbatim}

## S3 method for class 'Ts.estim'
print(x, ...)

Arguments

- **formula**: A formula object, with the response on the left of a \sim operator, and the covariates on the right. The right hand side of the model must also include two special terms: a strata and a cluster term (ex. \texttt{formula} = \texttt{Y \sim X1 + X2 + X3 + strata(var_strata) + cluster(var_cluster)}).

- **data**: A data frame (or object coercible by as.data.frame to a data frame) containing the variables in the model.

- **random**: A formula object, with a blank on the left of a \sim operator, and, on the right, the covariates with random coefficients among the covariate listed in the model formula (ex. \texttt{random} = \texttt{~ X1 + X3}). The default is to add random coefficients for every covariates listed in the model formula.

- **all.m.1**: TRUE if sum of Y’s in all strata is 1, FALSE otherwise (the default). When in doubt use FALSE (always works, but slower than necessary if all stratum sums are 1).

- **D**: The form of the between-cluster variance-covariance matrix of the regression coefficients (matrix \( D \)) : either "UN" for unstructured matrix \( D \) or "UN(1)" (the default) for diagonal matrix \( D \).

- **itermax**: maximal number of EM iterations (default = 2000)

- **toler**: maximal distance between successive EM iterations tolerated before declaring convergence (default = 0.000001)

- **x**: An object, produced by the \texttt{Ts.estim} function, to print.

- **...**: Further arguments to be passed to print.default.

Details

Calls \texttt{coxph} from the package \texttt{survival}. 

Value

- **beta**: A vector: the regression coefficients.
- **se**: A vector: the regression coefficients’ standard errors.
- **vcov**: A matrix: the variance-covariance matrix of the regression coefficients.
- **D**: A matrix: estimate of the between-cluster variance-covariance matrix of the regression coefficients (matrix D).
- **r.effect**: The random effect estimates.
- **coxph.warn**: A list of character string vectors. If the coxph function generates one or more warnings when fitting the Cox model to a cluster, a copy of these warnings are stored in coxph.warn$Cluster_name where Cluster_name is the identification value for the cluster in the data set. A NULL list element means that coxph did not produce any warnings for that cluster.

Call

The function call.

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References


See Also

- ddim

Examples

# Two ways for specifying the same model
# Data: bison
# Model: covariates forest, biomass and pmeadow
# Random effects in front of forest and biomass
# Main diagonal covariance structure for D (the default)
way1 <- Ts.estim(formula = Y ~ forest + biomass + pmeadow +
                   strata(Strata) + cluster(Cluster), data = bison,
                   random = ~ forest + biomass)

way1

way2 <- Ts.estim(formula = bison[,3] ~ as.matrix(bison[,c(6,8:9)]) +
                 strata(bison[,2]) + cluster(bison[,1]), data = bison,
                 random = ~ as.matrix(bison[,c(6,8)]))

way2

# Unstructured covariance for D
Fit <- Ts.estim(formula = Y ~ forest + biomass + pmeadow +
                strata(Strata) + cluster(Cluster), data = bison,
                random = ~ forest + biomass, D="UN")

Fit
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