

Package ‘UdderQuarterInfectionData’

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Type Package

Title Udder Quarter Infection Data

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Description

The udder quarter infection data set contains infection times of individual cow udder quarters with *Corynebacterium bovis* (Laevens et al. 1997 <[DOI:10.3168/jds.S0022-0302\(97\)76295-7](https://doi.org/10.3168/jds.S0022-0302(97)76295-7)>). Obviously, the four udder quarters are clustered within a cow, and udder quarters are sampled only approximately monthly, generating interval-censored data. The data set contains both covariates that change within a cow (e.g., front and rear udder quarters) and covariates that change between cows (e.g., parity [the number of previous calvings]). The correlation between udder infection times within a cow also is of interest, because this is a measure of the infectivity of the agent causing the disease. Various models have been applied to address the problem of interdependence for right-censored event times. These models, as applied to this data set, can be found back in the publications found in the reference list.

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

Imports stats, utils

NeedsCompilation no

Repository CRAN

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Contents

Gamma_Frailty_Interval_Censoring	2
udderquarterinfection	3
UdderQuarterInfectionData	5

Index	7
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`Gamma_Frailty_Interval_Censoring`*Gamma Frailty Interval Censoring*

Description

Application of the Gamma Frailty Interval Censoring Model on the Udder Quarter Infection Data Set. For more information see Details.

Usage

```
Gamma_Frailty_Interval_Censoring(print.level = 2)
```

Arguments

<code>print.level</code>	Parameter of <code>nlm</code> (default=2): this argument determines the level of printing which is done during the minimization process. The default value of 0 means that no printing occurs, a value of 1 means that initial and final details are printed and a value of 2 means that full tracing information is printed.
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Details

This function fits a parametric Weibull baseline hazard frailty model with gamma distributed frailties for the udder quarter infection data taking into consideration the interval censored nature of the data. Further theoretical details can be found in the paper in the reference

Value

Returns a list with the NLM result in `nlm` and the covariance matrix in `covmat`.

R Code for Model

The source R code for this model can found:

- in the `doc/Models_R_Code.R` file in the package installation folder.
- by accessing the function by calling `Gamma_Frailty_Interval_Censoring` (without brackets) or `getAnywhere("Gamma_Frailty_Interval_Censoring")`.

Author(s)

Klara Goethals

Luc Duchateau

References

Goethals, K., Ampe, B., Berkvens, D., Laevens, H., Janssen, P. and Duchateau, L. (2009). Modeling interval-censored, clustered cow udder quarter infection times through the shared gamma frailty model. *Journal of Agricultural Biological and Environmental Statistics* 14, 1-14.

Examples

```
## Not run:
library(UdderQuarterInfectionData)
data("udderquarterinfection")

Gamma_Fraily_Interval_Censoring()
# $nlm
# $nlm$minimum
# [1] 5670.491
#
# $nlm$estimate
# [1] 3.7967246 0.1201593 1.9672298 0.8590531
#
# $nlm$gradient
# [1] 0.0002924871 0.0017653292 -0.0005460029 0.0003265086
#
# $nlm$hessian
# [,1] [,2] [,3] [,4]
# [1,] 23.22965 -117.7682 -39.93813 -10.10561
# [2,] -117.76825 15471.4753 567.24283 1228.87332
# [3,] -39.93813 567.2428 664.76359 24.63047
# [4,] -10.10561 1228.8733 24.63047 147.76479
#
# $nlm$code
# [1] 1
#
# $nlm$iterations
# [1] 22
#
#
# $covmat
# [,1] [,2] [,3] [,4]
# [1,] 0.049281911 0.0001242730 0.0027853686 0.001872592
# [2,] 0.000124273 0.0001982213 -0.0001015391 -0.001623066
# [3,] 0.002785369 -0.0001015391 0.0017306214 0.000746460
# [4,] 0.001872592 -0.0016230660 0.0007464600 0.020269244

## End(Not run)
```

udderquarterinfection *Udder Quarter Infection Data*

Description

The udder quarter infection data contains the times to infection of the individual udder quarters of 100 cows during one lactation period. The 'cowid' variable contains the cow number. Each cow has 4 different quarters, with the variable 'quarter' referring to the specific quarter, i.e., RF: right front, RR: right rear, LF: left front, LR: left rear. The lactation number, i.e., the number of calvings that the cow has experienced, is given by the variable 'lactation'. If at least one infection occurs in

a quarter during the lactation period, the 'status' variable is set to 1, and the interval [t1,t2] in which an infection takes place is given by the variables 'left' and 'right', where 'right' denotes the time with the first positive result, and 'left' the last time (with a negative result) before the first positive result. Only the first infection is contained in the data set. If no infection occurs, the 'status' variable is set to 0, and the last interval [t1,t2] in which the quarter was observed is put in the variables 'left' and 'right'.

The following example analyses of this data set are currently available in the package:

- [Gamma_Fraily_Interval_Censoring](#)

Format

A dataframe with 4784 rows and 6 variables.

Examples

```
## Not run:
## SHORT DATASET ANALYSIS ##

library(UdderQuarterInfectionData)
data("udderquarterinfection")

Gamma_Fraily_Interval_Censoring()
# $nlm
# $nlm$minimum
# [1] 5670.491
#
# $nlm$estimate
# [1] 3.7967246 0.1201593 1.9672298 0.8590531
#
# $nlm$gradient
# [1] 0.0002924871 0.0017653292 -0.0005460029 0.0003265086
#
# $nlm$hessian
# [,1] [,2] [,3] [,4]
# [1,] 23.22965 -117.7682 -39.93813 -10.10561
# [2,] -117.76825 15471.4753 567.24283 1228.87332
# [3,] -39.93813 567.2428 664.76359 24.63047
# [4,] -10.10561 1228.8733 24.63047 147.76479
#
# $nlm$code
# [1] 1
#
# $nlm$iterations
# [1] 22
#
# $covmat
# [,1] [,2] [,3] [,4]
# [1,] 0.049281911 0.0001242730 0.0027853686 0.001872592
# [2,] 0.000124273 0.0001982213 -0.0001015391 -0.001623066
# [3,] 0.002785369 -0.0001015391 0.0017306214 0.000746460
```

```
# [4,] 0.001872592 -0.0016230660 0.0007464600 0.020269244
## End(Not run)
```

UdderQuarterInfectionData

Udder Quarter Infection Data

Description

This package is built up around a data set consisting of clustered infection times in the four udder quarters of cows. Due to the special structure of these data, i.e., quadruples of four clustered event times, they have been used to develop new techniques in the area of multivariate survival data (see references). This package avails the data on which the analyses presented in the papers were based, and also, where possible, provides the R-code to fit the proposed models.

Author(s)

Hans Laevens
 Luc Duchateau
 Klara Goethals
 Ewoud de Troyer
 Paul Janssen

References

- Laevens, H., Deluyker, H., Schukken, Y.H., De Meulemeester, L., Vandermeersch, R., De Muele-naere, E. and De Kruif, A. (1997). Influence of parity and stage of lactation on the somatic cell count in bacteriologically negative dairy cows. *Journal of Dairy Science* 80, 3219-3226.
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- Goethals, K., Ampe, B., Berkvens, D., Laevens, H., Janssen, P. and Duchateau, L. (2009). Modeling interval-censored, clustered cow udder quarter infection times through the shared gamma frailty model. *Journal of Agricultural Biological and Environmental Statistics* 14, 1-14.
- Janssen, P. and Duchateau, L. (2011). Comments on: Inference in multivariate Archimedean copula models. *Test* 20, 271-275.
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- Geerdens, C., Claeskens, G., Janssen, P. (2016). Copula based flexible modeling of associations between clustered event times. *Lifetime data analysis* 22, 363-381.

Verbelen, R., Antonio, K. and Claeskens, G. (2016). Multivariate mixtures of Erlangs for density estimation under censoring. *Lifetime data analysis* 22, 429-455.

Prenen, L., Duchateau, L. and Braekers, R. (2018). Investigating the correlation structure of quadri-variate udder infection times through hierarchical Archimedean copulas. Accepted for publication in *Lifetime Data Analysis*.

Index

Gamma_Frailty_Interval_Censoring, [2](#), [4](#)

nlm, [2](#)

udderquarterinfection, [3](#)

UdderQuarterInfectionData, [5](#)

UdderQuarterInfectionData-package
 (UdderQuarterInfectionData), [5](#)