Package ‘aftgee’

July 24, 2018

Title Accelerated Failure Time Model with Generalized Estimating
Equations

Version 1.1.3

Description A collection of methods for both the rank-based estimates and least-square estimates
to the Accelerated Failure Time (AFT) model.
For rank-based estimation, it provides approaches that include the computationally
efficient Gehan's weight and the general's weight such as the logrank weight.
Details of the rank-based estimation can be found in
Chiou et al. (2014) <doi:10.1007/s11222-013-9388-2> and
Chiou et al. (2015) <doi:10.1002/sim.6415>.
For the least-square estimation, the estimating equation is solved with
generalized estimating equations (GEE).
Moreover, in multivariate cases, the dependence working correlation structure
can be specified in GEE's setting.
Details on the least-squares estimation can be found in
Chiou et al. (2014) <doi:10.1007/s10985-014-9292-x>.

Depends R (>= 3.4.0)

License GPL (>= 3)

URL http://github.com/stc04003/aftgee

BugReports http://github.com/stc04003/aftgee/issues

Encoding UTF-8

LazyData true

LazyLoad yes

Suggests copula

Imports methods, parallel, geepack, survival, BB, MASS

RoxygenNote 6.0.1

NeedsCompilation yes

Author Sy Han Chiou [aut, cre],
Sangwook Kang [aut],
Jun Yan [aut]

Maintainer Sy Han Chiou <schiou@utdallas.edu>

Repository CRAN

Date/Publication 2018-07-24 04:50:03 UTC

R topics documented:

`aftgee-package` *aftgee: Accelerated Failure Time with Generalized Estimating Equation*

Description

A package that uses Generalized Estimating Equations (GEE) to estimate Multivariate Accelerated Failure Time Model (AFT). This package implements recently developed inference procedures for AFT models with both the rank-based approach and the least squares approach. For the rank-based approach, the package allows various weight choices and uses an induced smoothing procedure that leads to much more efficient computation than the linear programming method. With the rank-based estimator as an initial value, the generalized estimating equation approach is used as an extension of the least squares approach to the multivariate case. Additional sampling weights are incorporated to handle missing data needed as in case-cohort studies or general sampling schemes.

Author(s)

Maintainer: Sy Han Chiou <`schiou@utdallas.edu`>

Authors:

- Sangwook Kang
- Jun Yan

References

Chiou, S., Kim, J. and Yan, J. (2014) Marginal Semiparametric Multivariate Accelerated Failure Time Model with Generalized Estimating Equation. *Life Time Data*, **20**(4): 599–618.

Chiou, S., Kang, S. and Yan, J. (2014) Fast Accelerated Failure Time Modeling for Case-Cohort Data. *Statistics and Computing*, **24**(4): 559–568.

Chiou, S., Kang, S. and Yan, J. (2014) Fitting Accelerated Failure Time Model in Routine Survival Analysis with R Package **aftgee**. *Journal of Statistical Software*, **61**(11): 1–23.

Huang, Y. (2002) Calibration Regression of Censored Lifetime Medical Cost. *Journal of American Statistical Association*, **97**, 318–327.

Jin, Z. and Lin, D. Y. and Ying, Z. (2006) On Least-squares Regression with Censored Data. *Biometrika*, **90**, 341–353.

Johnson, L. M. and Strawderman, R. L. (2009) Induced Smoothing for the Semiparametric Accelerated Failure Time Model: Asymptotic and Extensions to Clustered Data. *Biometrika*, **96**, 577 – 590.

Zeng, D. and Lin, D. Y. (2008) Efficient Resampling Methods for Nonsmooth Estimating Functions. *Biostatistics*, **9**, 355–363

See Also

Useful links:

- `http://github.com/stc04003/aftgee`
- Report bugs at `http://github.com/stc04003/aftgee/issues`

`aftgee` *Least-Squares Approach for Accelerated Failure Time with Generalized Estimating Equation*

Description

Fits a semiparametric accelerated failure time (AFT) model with least-squares approach. Generalized estimating equation is generalized to multivariate AFT modeling to account for multivariate dependence through working correlation structures to improve efficiency.

Usage

```
aftgee(formula, data, subset, id = NULL, contrasts = NULL, weights = NULL,
  margin = NULL, corstr = "independence", binit = "srrgehan", B = 100,
  control = aftgee.control())
```

Arguments

`formula` a formula expression, of the form `response ~ predictors`. The `response` is a `Surv` object object with right censoring. In the case of no censoring, `aftgee` will return an ordinary least estimate when `corstr = "independence"`. See the documentation of `lm`, `coxph` and `formula` for details.

`data` an optional data.frame in which to interpret the variables occurring in the `formula`.

`subset` an optional vector specifying a subset of observations to be used in the fitting process.

`id` an optional vector used to identify the clusters. If missing, then each individual row of `data` is presumed to represent a distinct subject. The length of `id` should be the same as the number of observations.

`contrasts` an optional list.

`weights` an optional vector of observation weights.

`margin` a `sformula` vector; default at 1.

`corstr` a character string specifying the correlation structure. The following are permitted:

- `independence`
- `exchangeable`
- `ar1`
- `unstructured`
- `userdefined`
- `fixed`

`binit` an optional vector can be either a numeric vector or a character string specifying the initial slope estimator.

- When `binit` is a vector, its length should be the same as the dimension of covariates.
- When `binit` is a character string, it should be either `lm` for simple linear regression, or `srrgehan` for smoothed Gehan weight estimator.

The default value is "srrgehan".

`B` a numeric value specifies the resampling number. When B = 0, only the beta estimate will be displayed.

`control` controls maxiter and tolerance.

Value

An object of class "`aftgee`" representing the fit. The `aftgee` object is a list containing at least the following components:

coefficients a vector of initial value and a vector of point estimates

coef.res a vector of point estimates

var.res estimated covariance matrix

coef.init a vector of initial value

var.init.mat estimated initial covariance matrix

binit a character string specifying the initial estimator.

conv An integer code indicating type of convergence after GEE iteration. 0 indicates successful convergence; 1 indicates that the iteration limit `maxit` has been reached

ini.conv An integer code indicating type of convergence for initial value. 0 indicates successful convergence; 1 indicates that the iteration limit `maxit` has been reached

conv.step An integer code indicating the step until convergence

References

Chiou, S., Kim, J. and Yan, J. (2014) Marginal Semiparametric Multivariate Accelerated Failure Time Model with Generalized Estimating Equation. *Life Time Data*, **20**(4): 599–618.

Jin, Z. and Lin, D. Y. and Ying, Z. (2006) On Least-squares Regression with Censored Data. *Biometrika*, **90**, 341–353.

Examples

```
library(survival)
library(copula)
datgen <- function(n = 100, tau = 0.3, cen = 75.4, dim = 2) {
    kt <- iTau(claytonCopula(1), tau)
    copula <- claytonCopula(kt, dim = dim)
    id <- rep(1:n, rep(dim, n))
    x1 <- rbinom(dim * n, 1, 0.5)
    x2 <- rnorm(dim * n)
    ed <- mvdc(copula, rep("weibull", dim), rep(list(list(shape = 1)), dim))
    e <- c(t(rMvdc(n, ed)))
    T <- exp(2 + x1 + x2 + e)
    cstime <- runif(n, 0, cen)
    delta <- (T < cstime) * 1
    Y <- pmin(T, cstime)
   out <- data.frame(T = T, Y = Y, delta = delta, x1 = x1, x2 = x2, id = rep(1:n, each = dim))
    out
}
set.seed(1)
mydata <- datgen(n = 50, dim = 2)
summary(aftgee(Surv(Y, delta) ~ x1 + x2, data = mydata,
               id = id, corstr = "ind", B = 8))
summary(aftgee(Surv(Y, delta) ~ x1 + x2, data = mydata,
               id = id, corstr = "ex", B = 8))
```

`aftgee.control` *Auxiliary for Controlling AFTGEE Fitting*

Description

Auxiliary function as user interface for `aftgee` and `aftsrr` fitting.

Usage

```
aftgee.control(maxiter = 50, reltol = 0.001, trace = FALSE,
  seIni = FALSE, parallel = FALSE, parCl = parallel::detectCores()/2)
```

Arguments

`maxiter` max number of iteration.

`reltol` relative error tolerance.

`trace` a binary variable, determine whether to display output for each iteration.

`seIni` a logical value indicating whether a new rank-based initial value is computed for each resampling sample in variance estimation.

`parallel` an logical value indicating whether parallel computing is used for resampling and bootstrap.

`parCl` an integer value indicating the number of CPU cores used when `parallel = TRUE`. The default value is half the CPU cores on the current host.

Details

When `trace` is TRUE, output for each iteration is printed to the screen.

Value

A list with the arguments as components.

See Also

`aftgee`

`aftsrr` *Accelerated Failure Time with Smooth Rank Regression*

Description

Fits a semiparametric accelerated failure time (AFT) model with rank-based approach. General weights, additional sampling weights and fast sandwich variance estimations are also incorporated. Estimating equations are solved with Barzilar-Borwein spectral method implemented as `BBsolve` in package **BB**.

Usage

```
aftsrr(formula, data, subset, id = NULL, contrasts = NULL, weights = NULL,
  B = 100, rankWeights = c("gehan", "logrank", "pw", "gp", "userdefined"),
  eqType = c("is", "ns", "mns", "mis"), se = c("NULL", "bootstrap", "MB",
  "ZLCF", "ZLMB", "sHCF", "sHMB", "ISCF", "ISMB"), control = list())
```

Arguments

`formula` a formula expression, of the form `response ~ predictors`. The `response` is a `Surv` object object with right censoring. See the documentation of `lm`, `coxph` and `formula` for details.

`data` an optional data frame in which to interpret the variables occurring in the `formula`.

`subset` an optional vector specifying a subset of observations to be used in the fitting process.

`id` an optional vector used to identify the clusters. If missing, then each individual row of `data` is presumed to represent a distinct subject. The length of `id` should be the same as the number of observation.

`contrasts` an optional list.

`weights` an optional vector of observation weights.

`B` a numeric value specifies the resampling number. When B = 0, only the beta estimate will be displayed.

`rankWeights` a character string specifying the type of general weights. The following are permitted:

- `logrank` logrank weight
- `gehan` Gehan's weight
- `PW` Prentice-Wilcoxon weight
- `GP` GP class weight

`eqType` a character string specifying the type of the estimating equation used to obtain the regression parameters. The following are permitted:

- `nonsm` Regression parameters are estimated by directly solving the nonsmooth estimating equations.
- `sm` Regression parameters are estimated by directly solving the induced-smoothing estimating equations.
- `monosm` Regression parameters are estimated by iterating the monotonic smoothed estimating equations. This is typical when `rankWeights = "PW"` and `rankWeights = "GP"`.

`se` a character string specifying the estimating method for the variance-covariance matrix. The following are permitted:

- `bootstrap` nonparametric bootstrap,
- `MB` multiplier resampling.
- `ZLCF` Zeng and Lin's approach with closed form Si.
- `ZLMB` Zeng and Lin's approach with empirical Si.
- `sHCF` Huang's approach with closed form Si.
- `sHMB` Huang's approach with empirical Si.
- `ISCF` Johnson and Strawderman's sandwich variance estimates with closed form Si.
- `ISMB` Johnson and Strawderman's sandwich variance estimates with empirical Si.
- `js` Johnson and Strawderman's iterating approach.

`control` controls equation solver, maxiter, tolerance, and resampling variance estimation. The available equation solvers are `BBsolve` and `dfsane` of the **BB** package. Instead of searching for the zero crossing, options including `BBoptim` and `optim` will return solution from maximizing the corresponding objective function.

Value

`aftsrr` returns an object of class "`aftsrr`" representing the fit. An object of class "`aftsrr`" is a list containing at least the following components:

beta A vector of beta estimates

covmat A list of covariance estimates

convergence An integer code indicating type of convergence.

- **0** indicates successful convergence.
- **1** indicates that the iteration limit `maxit` has been reached.
- **2** indicates failure due to stagnation.
- **3** indicates error in function evaluation.
- **4** is failure due to exceeding 100 step length reductions in line-search.
- **5** indicates lack of improvement in objective function.

bhist When `variance = "MB"`, `bhist` gives the bootstrap samples.

References

Chiou, S., Kang, S. and Yan, J. (2014) Fast Accelerated Failure Time Modeling for Case-Cohort Data. *Statistics and Computing*, **24**(4): 559–568.

Chiou, S., Kang, S. and Yan, J. (2014) Fitting Accelerated Failure Time Model in Routine Survival Analysis with R Package **Aftgee**. *Journal of Statistical Software*, **61**(11): 1–23.

Huang, Y. (2002) Calibration Regression of Censored Lifetime Medical Cost. *Journal of American Statistical Association*, **97**, 318–327.

Johnson, L. M. and Strawderman, R. L. (2009) Induced Smoothing for the Semiparametric Accelerated Failure Time Model: Asymptotic and Extensions to Clustered Data. *Biometrika*, **96**, 577 – 590.

Varadhan, R. and Gilbert, P. (2009) BB: An R Package for Solving a Large System of Nonlinear Equations and for Optimizing a High-Dimensional Nonlinear Objective Function. *Journal of Statistical Software*, **32**(4): 1–26

Zeng, D. and Lin, D. Y. (2008) Efficient Resampling Methods for Nonsmooth Estimating Functions. *Biostatistics*, **9**, 355–363

Examples

```
## kidney data
library(survival)
data(kidney)
foo <- aftsrr(Surv(time, status) ~ age + sex, id = id,
                data = kidney, se = c("ISMB", "ZLMB"), B = 10)
foo

## nwtco data
library(survival)
data(nwtco)
subinx <- sample(1:nrow(nwtco), 668, replace = FALSE)
nwtco$subcohort <- 0
nwtco$subcohort[subinx] <- 1
pn <- table(nwtco$subcohort)[[2]] / sum(table(nwtco$subcohort))
nwtco$hi <- nwtco$rel + ( 1 - nwtco$rel) * nwtco$subcohort / pn
nwtco$age12 <- nwtco$age / 12
nwtco$study <- nwtco$study - 3
nwtco$histol = nwtco$histol - 1
sub <- nwtco[subinx,]
fit <- aftsrr(Surv(edrel, rel) ~ histol + age12 + study, id = seqno,
       weights = hi, data = sub, B = 10, se = c("ISMB", "ZLMB"),
       subset = stage == 4)
summary(fit)
```

Index