

# Package: intccr (via r-universe)

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**Type** Package

**Title** Semiparametric Competing Risks Regression under Interval Censoring

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**Description** Semiparametric regression models on the cumulative incidence function for interval-censored competing risks data as described in Bakoyannis, Yu, & Yiannoutsos (2017) /doi{10.1002/sim.7350} and the models with missing event types as described in Park, Bakoyannis, Zhang, & Yiannoutsos (2021) \doi{10.1093/biostatistics/kxaa052}. The proportional subdistribution hazards model (Fine-Gray model), the proportional odds model, and other models that belong to the class of semiparametric generalized odds rate transformation models.

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bs.derivs	<i>Derivative of B-spline</i>
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**Description**

Generates the derivative of the B-splines basis matrix.

**Usage**

```
bs.derivs(  
  x,  
  derivs = 0,  
  df = NULL,  
  knots = NULL,  
  degree = 3,  
  intercept = FALSE,  
  Boundary.knots = range(x)  
)
```

**Arguments**

x	object of B-splines
derivs	a number of derivatives
df	degrees of freedom of B-splines
knots	a vector of internal knots
degree	degrees of B-splines
intercept	a logical vector
Boundary.knots	a vector of boundary knots

**Details**

The function `bs.derivs` performs derivatives of B-splines

**Value**

The function `bs.derivs` returns a component:

resmat	derivatives of B-spline
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bssmle

*B-spline Sieve Maximum Likelihood Estimation*


---

### Description

Routine that performs B-spline sieve maximum likelihood estimation with linear and nonlinear inequality/equality constraints

### Usage

```
bssmle(formula, data, alpha, k = 1)
```

### Arguments

formula	a formula object relating survival object <code>Surv2(v, u, event)</code> to a set of covariates
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the cause of failure 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the cause of failure 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$

### Details

The function `bssmle` performs B-spline sieve maximum likelihood estimation.

### Value

The function `bssmle` returns a list of components:

beta	a vector of the estimated coefficients for the B-splines
varnames	a vector containing variable names
alpha	a vector of the link function parameters
loglikelihood	a loglikelihood of the fitted model
convergence	an indicator of convergence
tms	a vector of the minimum and maximum observation times
Z	a set of covariates
Tv	a vector of v
Tu	a vector of u
Bv	a list containing the B-splines basis functions evaluated at v
Bu	a list containing the B-splines basis functions evaluated at u

dBv	a list containing the first derivative of the B-splines basis functions evaluated at v
dBu	a list containing the first derivative of the B-splines basis functions evaluated at u
dmat	a matrix of event indicator functions

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---

bssmle_aipw	<i>B-spline Sieve Maximum Likelihood Estimation for Interval-Censored Competing Risks Data and Missing Cause of Failure</i>
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**Description**

Routine that performs B-spline sieve maximum likelihood estimation with linear and nonlinear inequality and equality constraints

**Usage**

```
bssmle_aipw(formula, aux, data, alpha, k)
```

**Arguments**

formula	a formula object relating survival object <code>Surv2(v, u, event)</code> to a set of covariates
aux	auxiliary variables that may be associated with the missingness and the outcome of interest
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the event type 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the event type 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$

**Details**

The function `bssmle_aipw` performs B-spline sieve maximum likelihood estimation.

**Value**

The function `bssmle_aipw` returns a list of components:

<code>beta</code>	a vector of the estimated coefficients for the B-splines
<code>varnames</code>	a vector containing variable names
<code>varnames.aux</code>	a vector containing auxiliary variable names
<code>alpha</code>	a vector of the link function parameters
<code>loglikelihood</code>	a loglikelihood of the fitted model
<code>convergence</code>	an indicator of convergence
<code>tms</code>	a vector of the minimum and maximum observation times
<code>Bv</code>	a list containing the B-splines basis functions evaluated at <code>v</code>

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`bssmle_lse`

*Least-Squares Estimator of the Information Matrix*

---

**Description**

Performs the least-squares methods to estimate the information matrix for the estimated regression coefficients

**Usage**

```
bssmle_lse(obj)
```

**Arguments**

`obj` a list of objectives from `bssmle`

**Details**

The function `bssmle_lse` estimates the information matrix for the estimated regression coefficients from the function `bssmle` using the least-squares method.

**Value**

The function `bssmle_lse` returns a list of components:

<code>Sigma</code>	the estimated variance-covariance matrix for the estimated regression coefficients
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**Author(s)**

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**References**

Zhang, Y., Hua, L., and Huang, J. (2010), A spline-based semiparametric maximum likelihood estimation method for the Cox model with interval-censored data. *Scandinavian Journal of Statistics*, **37**:338-354.

---

`bssmle_lse_lt`*Least-Squares Estimator of the Information Matrix*

---

**Description**

Performs the least-squares methods to estimate the information matrix for the estimated regression coefficients

**Usage**

```
bssmle_lse_lt(obj)
```

**Arguments**

`obj` a list of objectives from `bssmle_lt`

**Details**

The function `bssmle_lse_lt` estimates the information matrix for the estimated regression coefficients from the function `bssmle_lt` using the least-squares method.

**Value**

The function `bssmle_lse_lt` returns a list of components:

`Sigma` the estimated information matrix for the estimated regression coefficients

**Author(s)**

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**References**

Zhang, Y., Hua, L., and Huang, J. (2010), A spline-based semiparametric maximum likelihood estimation method for the Cox model with interval-censored data. *Scandinavian Journal of Statistics*, **37**:338-354.

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bssmle_lt	<i>B-spline Sieve Maximum Likelihood Estimation for Left-Truncated and Interval-Censored Competing Risks Data</i>
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---

### Description

Routine that performs B-spline sieve maximum likelihood estimation with linear and nonlinear inequality/equality constraints

### Usage

```
bssmle_lt(formula, data, alpha, k = 1)
```

### Arguments

formula	a formula object relating survival object <code>Surv2(w, v, u, event)</code> to a set of covariates
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the event type 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the event type 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$

### Details

The function `bssmle_lt` performs B-spline sieve maximum likelihood estimation for left-truncated and interval-censored competing risks data.

### Value

The function `bssmle_lt` returns a list of components:

beta	a vector of the estimated coefficients
varnames	a vector containing variable names
alpha	a vector of the link function parameters
loglikelihood	a loglikelihood of the fitted model
convergence	an indicator of convergence
tms	a vector of the minimum and maximum observation times
Z	a design matrix
Tw	a vector of w
Tv	a vector of v
Tu	a vector of u



Bw	a list containing the B-splines basis functions evaluated at w
Bv	a list containing the B-splines basis functions evaluated at v
Bu	a list containing the B-splines basis functions evaluated at u
dBw	a list containing the first derivative of the B-splines basis functions evaluated at w
dBv	a list containing the first derivative of the B-splines basis functions evaluated at v
dBu	a list containing the first derivative of the B-splines basis functions evaluated at u
dmat	a matrix of event indicator functions

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---

bssmle\_se

*Bootstrap varince-covariance estimation*


---

**Description**

Bootstrap varince estimation for the estimated regression coefficients

**Usage**

```
bssmle_se(formula, data, alpha, k = 1, do.par, nboot, objfun)
```

**Arguments**

formula	a formula object relating survival object <code>Surv2(v, u, event)</code> to a set of covariates
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the cause of failure 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the cause of failure 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$
do.par	using parallel computing for bootstrap calculation. If <code>do.par = TRUE</code> , parallel computing will be used during the bootstrap estimation of the variance-covariance matrix for the regression parameter estimates.
nboot	a number of bootstrap samples for estimating variances and covariances of the estimated regression coefficients. If <code>nboot = 0</code> , the function <code>cieregic</code> does not perform bootstrap estimation of the variance matrix of the regression parameter estimates and returns NA in the place of the estimated variance matrix of the regression parameter estimates.
objfun	an option to select estimating function

**Details**

The function `bssmle_se` estimates bootstrap standard errors for the estimated regression coefficients from the function `bssmle`, `bssmle_lt`, or `bssmle_ltir`.

**Value**

The function `bssmle_se` returns a list of components:

<code>notconverged</code>	a list of number of bootstrap samples that did not converge
<code>numboot</code>	a number of bootstrap converged
<code>Sigma</code>	an estimated bootstrap variance-covariance matrix of the estimated regression coefficients

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---

<code>bssmle_se_aipw</code>	<i>Bootstrap variance-covariance estimation for interval-censored competing risks data and missing cause of failure</i>
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---

**Description**

Bootstrap variance estimation for the estimated regression coefficients

**Usage**

```
bssmle_se_aipw(formula, aux, data, alpha, k, do.par, nboot, w.cores = NULL)
```

**Arguments**

<code>formula</code>	a formula object relating survival object <code>mSurv(v, u, event)</code> to a set of covariates
<code>aux</code>	auxiliary variables that may be associated with the missingness and the outcome of interest
<code>data</code>	a data frame that includes the variables named in the <code>formula</code> argument
<code>alpha</code>	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the event type 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the event type 2.
<code>k</code>	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$

do.par	using parallel computing for bootstrap calculation. If do.par = TRUE, parallel computing will be used during the bootstrap estimation of the variance-covariance matrix for the regression parameter estimates.
nboot	a number of bootstrap samples for estimating variances and covariances of the estimated regression coefficients. If nboot = 0, the function ciregic does not perform bootstrap estimation of the variance matrix of the regression parameter estimates and returns NA in the place of the estimated variance matrix of the regression parameter estimates.
w.cores	a number of cores that are assigned (the default is NULL)

### Details

The function `bssmle_aipw_se` estimates bootstrap standard errors for the estimated regression coefficients from the function `bssmle`.

### Value

The function `bssmle_aipw_se` returns a list of components:

notconverged	a list of number of bootstrap samples that did not converge
numboot	a number of bootstrap converged
Sigma	an estimated bootstrap variance-covariance matrix of the estimated regression coefficients

### Author(s)

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---

ciregic

*Competing Risks Regression with Interval-Censored Data*

---

### Description

The function `ciregic` performs semiparametric regression on cumulative incidence function with interval-censored competing risks data. It fits the proportional subdistribution hazards model (Fine-Gray model), the proportional odds model, and other models that belong to the class of semiparametric generalized odds rate transformation models. The standard errors for the estimated regression coefficients are estimated by a choice of options: 1) the bootstrapping method or 2) the least-squares method.

### Usage

```
ciregic(formula, data, alpha, k = 1, do.par, nboot, ...)
```

**Arguments**

formula	a formula object relating the survival object <code>Surv2(v, u, event)</code> to a set of covariates
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the cause of failure 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the cause of failure 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$
do.par	an option to use parallel computing for bootstrap. If <code>do.par = TRUE</code> , parallel computing will be used during the bootstrap estimation of the variance-covariance matrix for the regression parameter estimates.
nboot	a number of bootstrap samples for estimating variances and covariances of the estimated regression coefficients. If <code>nboot = 0</code> , the function <code>ciregic</code> provides the variance estimator of the regression parameter estimates using the least-squares method and does not perform the bootstrap method.
...	further arguments

**Details**

The formula for the model has the form of `response ~ predictors`. The response in the formula is a `Surv2(v, u, event)` object where `v` is the last observation time prior to the failure, `u` is the first observation time after the failure, and `event` is the event or censoring indicator. `event` should include 0, 1 or 2, denoting right-censoring, failure from cause 1 and failure from cause 2, respectively. If `event=0` (i.e. right-censored observation) then `u` is not included in any calculation as it corresponds to  $\infty$ . The user can provide any value in `u` for the right-censored cases, even `NA`. The function `ciregic` fits models that belong to the class of generalized odds rate transformation models which includes the proportional subdistribution hazards or the Fine-Gray model and the proportional odds model. The parameter  $\alpha = (\alpha_1, \alpha_2)$  defines the link function/model to be fitted for cause of failure 1 and 2, respectively. A value of 0 corresponds to the Fine-Gray model and a value of 1 corresponds to the proportional odds model. For example, if  $\alpha = (0, 1)$  then the function `ciregic` fits the Fine-Gray model for cause 1 and the proportional odds model for cause 2.

**Value**

The function `ciregic` provides an object of class `ciregic` with components:

varnames	a vector containing variable names
coefficients	a vector of the regression coefficient estimates
gamma	a vector of the estimated coefficients for the B-splines
vcov	a variance-covariance matrix of the estimated regression coefficients
alpha	a vector of the link function parameters
loglikelihood	a loglikelihood of the fitted model
convergence	an indicator of convergence

tms	a vector of the minimum and maximum observation times
Bv	a list containing the B-splines basis functions evaluated at v
numboot	a number of converged bootstrap
notconverged	a list of number of bootstrap samples that did not converge
call	a matched call

**Author(s)**

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**References**

Bakoyannis, G., Yu, M., and Yiannoutsos C. T. (2017). Semiparametric regression on cumulative incidence function with interval-censored competing risks data. *Statistics in Medicine*, **36**:3683-3707.

Fine, J. P. and Gray, R. J. (1999). A proportional hazards model for the subdistribution of a competing risk. *Journal of the American Statistical Association*, **94**:496-509.

**See Also**

[summary.ciregic](#) for the summarized results and [predict.ciregic](#) for value of the predicted cumulative incidence functions. [coef](#) and [vcov](#) are the generic functions. [dataprep](#) for reshaping data from a long format to a suitable format to be used in the function `ciregic`.

**Examples**

```
## Not run:
## Set seed in order to have reproducibility of the bootstrap standard error estimate
set.seed(1234)

## Reshaping data from a long format to a suitable format
newdata <- dataprep(data = longdata, ID = id, time = t,
                    event = c, Z = c(z1, z2))
## Estimation of regression parameters only. No bootstrap variance estimation.
## with 'newdata'
fit <- ciregic(formula = Surv2(v = v, u = u, event = c) ~ z1 + z2, data = newdata,
              alpha = c(1, 1), nboot = 0, do.par = FALSE)
fit

## Bootstrap variance estimation based on 50 replications
fit <- ciregic(formula = Surv2(v = v, u = u, event = c) ~ z1 + z2, data = newdata,
              alpha = c(1, 1), nboot = 50, do.par = FALSE)

## End(Not run)
## Note that the user can use parallel computing to decrease
## the computation time of the bootstrap variance-covariance
## estimation (e.g. nboot = 50)
```

```
## Summarize semiparametric regression model
summary(fit)

## Predict and draw plot the cumulative incidence function evaluated at z1 = 1 and z2 = 0.5
t <- seq(from = 0, to = 2.8, by = 2.8 / 99)
pred <- predict(object = fit, covp = c(1, 0.5), times = t)
pred
plot(pred$t, pred$CIF1, type = "l", ylim = c(0, 1))
points(pred$t, pred$CIF2, type = "l", col = 2)
```

---

ciregic\_aipw

*Competing Risks Regression with Interval-Censored Data and Missing Cause of Failure*

---

### Description

The function `ciregic_aipw` performs semiparametric regression on cumulative incidence function with interval-censored competing risks data in the presence of missing cause of failure. It fits the proportional subdistribution hazards model (Fine-Gray model), the proportional odds model, and other models that belong to the class of semiparametric generalized odds rate transformation models. The estimates have double robustness property, which means that the estimators are consistent even if either the model for the probability of missingness or the model for the probability of the cause of failure is misspecified under the missing at random assumption.

### Usage

```
ciregic_aipw(
  formula,
  aux = NULL,
  data,
  alpha,
  k = 1,
  do.par,
  nboot,
  w.cores = NULL,
  ...
)
```

### Arguments

<code>formula</code>	a formula object relating the survival object <code>Surv2(v, u, event)</code> to a set of covariates
<code>aux</code>	auxiliary variable(s) that may be associated with the missingness and the outcome of interest
<code>data</code>	a data frame that includes the variables named in the formula argument

alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the event type 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the event type 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$
do.par	an option to use parallel computing for bootstrap. If <code>do.par = TRUE</code> , parallel computing will be used during the bootstrap estimation of the variance-covariance matrix for the regression parameter estimates.
nboot	a number of bootstrap samples for estimating variances and covariances of the estimated regression coefficients. If <code>nboot = 0</code> , the function <code>ciregic_aipw</code> does not perform bootstrap estimation of the variance-covariance matrix of the regression parameter estimates and returns NA in the place of the estimated variance-covariance matrix of the regression parameter estimates.
w.cores	a number of cores that are assigned (the default is NULL)
...	further arguments

### Details

The formula for the model has the form of `response ~ predictors`. The response in the formula is a `Surv2(v, u, event)` object where `v` is the last observation time prior to the event, `u` is the first observation time after the event, and `event` is the event or censoring indicator. `event` should include 0, 1 or 2, denoting right-censoring, event type 1 and 2, respectively. If `event=0` (i.e. right-censored observation) then `u` is not included in any calculation as it corresponds to  $\infty$ . The user can provide any value in `u` for the right-censored cases, even NA. The function `ciregic_aipw` fits models that belong to the class of generalized odds rate transformation models which includes the proportional subdistribution hazards or the Fine-Gray model and the proportional odds model. The parameter  $\alpha = (\alpha_1, \alpha_2)$  defines the link function/model to be fitted for event 1 and 2, respectively. A value of 0 corresponds to the Fine-Gray model and a value of 1 corresponds to the proportional odds model. For example, if  $\alpha = (0, 1)$  then the function `ciregic_aipw` fits the Fine-Gray model for the event type 1 and the proportional odds model for the event type 2.

### Value

The function `ciregic_aipw` provides an object of class `ciregic_aipw` with components:

<code>varnames</code>	a vector containing variable names
<code>varnames.aux</code>	a vector containing auxiliary variable names
<code>coefficients</code>	a vector of the regression coefficient estimates
<code>gamma</code>	a vector of the estimated coefficients for the B-splines
<code>vcov</code>	a variance-covariance matrix of the estimated regression coefficients
<code>alpha</code>	a vector of the link function parameters
<code>loglikelihood</code>	a loglikelihood of the fitted model
<code>convergence</code>	an indicator of convergence
<code>tms</code>	a vector of the minimum and maximum observation times

Bv                    a list containing the B-splines basis functions evaluated at v  
 numboot            a number of converged bootstrap  
 notconverged      a list of number of bootstrap samples that did not converge  
 call                a matched call

### Author(s)

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### References

Bakoyannis, G., Yu, M., and Yiannoutsos C. T. (2017). Semiparametric regression on cumulative incidence function with interval-censored competing risks data. *Statistics in Medicine*, **36**:3683-3707.

Fine, J. P. and Gray, R. J. (1999). A proportional hazards model for the subdistribution of a competing risk. *Journal of the American Statistical Association*, **94**:496-509.

### See Also

[summary.ciregic\\_aipw](#) for the summarized results and [predict.ciregic\\_aipw](#) for value of the predicted cumulative incidence functions. `coef` and `vcov` are the generic functions. `dataprep` function for reshaping data from a long format to a suitable format to be used in the function `ciregic_aipw`.

### Examples

```
## Not run:
## Set seed in order to have reproducibility of the bootstrap standard error estimate
set.seed(1234)

## Estimation of regression parameters only. No bootstrap variance estimation.
## with 'simdata_aipw'
data(simdata_aipw)
fit_aipw <- ciregic_aipw(formula = Surv2(v = v, u = u, event = c) ~ z1 + z2, aux = a,
                        data = simdata_aipw, alpha = c(1, 1), nboot = 0,
                        do.par = FALSE)

fit_aipw
## Bootstrap variance estimation based on 50 replications
fit_aipw <- ciregic_aipw(formula = Surv2(v = v, u = u, event = c) ~ z1 + z2, aux = a,
                        data = simdata_aipw, alpha = c(1, 1), k = 1, nboot = 50,
                        do.par = FALSE)

## End(Not run)
## Note that the user can use parallel computing to decrease
## the computation time of the bootstrap variance-covariance
## estimation (e.g. nboot = 50)

## Summarize semiparametric regression model
summary(fit_aipw)
```



```
## Predict and draw plot the cumulative incidence function evaluated at z1 = 1 and z2 = 0.5
t <- seq(from = 0, to = 2.8, by = 2.8 / 99)
pred <- predict(object = fit_aipw, covp = c(1, 0.5), times = t)
pred
plot(pred$t, pred$CIF1, type = "l", ylim = c(0, 1))
points(pred$t, pred$CIF2, type = "l", col = 2)
```

---

ciregic_lt	<i>Competing Risks Regression with Left-truncated and Interval-Censored Data</i>
------------	--

---

## Description

The function `ciregic_lt` performs semiparametric regression on cumulative incidence function with left-truncated and interval-censored competing risks data. It fits the proportional subdistribution hazards model (Fine-Gray model), the proportional odds model, and other models that belong to the class of semiparametric generalized odds rate transformation models. The least-square method is implemented to estimate the standard error of the regression coefficients.

## Usage

```
ciregic_lt(formula, data, alpha, k = 1, do.par, nboot, ...)
```

## Arguments

formula	a formula object relating the survival object <code>Surv2(v, u, w, event)</code> to a set of covariates
data	a data frame that includes the variables named in the formula argument
alpha	$\alpha = (\alpha_1, \alpha_2)$ contains parameters that define the link functions from class of generalized odds-rate transformation models. The components $\alpha_1$ and $\alpha_2$ should both be $\geq 0$ . If $\alpha_1 = 0$ , the user assumes the proportional subdistribution hazards model or the Fine-Gray model for the cause of failure 1. If $\alpha_2 = 1$ , the user assumes the proportional odds model for the cause of failure 2.
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$
do.par	an option to use parallel computing for bootstrap. If <code>do.par = TRUE</code> , parallel computing will be used during the bootstrap estimation of the variance-covariance matrix for the regression parameter estimates.
nboot	a number of bootstrap samples for estimating variances and covariances of the estimated regression coefficients. If <code>nboot = 0</code> , the function <code>ciregic_lt</code> returns a closed-form variance estimator using the least-squares method and does not perform bootstrap estimation of the variance-covariance matrix of the regression parameter estimates. For <code>nboot <math>\geq 1</math></code> , the function <code>ciregic_lt</code> returns the bootstrap variance estimator of the regression parameter estimates.
...	further arguments

## Details

The function `ciregic_lt` is capable of analyzing left-truncated and interval-censored competing risks data. A triplet of time points ( $w, v, u$ ) is required if an observation is left-truncated and interval-censored. A part of left-truncation is also allowed by defining  $w = 0$  for interval-censored only observation. The formula for the model has the form of `response ~ predictors`. The response in the formula is a `Surv2(v, u, w, event)` object where  $w$  is a left-truncation time,  $v$  is the last observation time prior to the failure,  $u$  is the first observation time after the failure, and `event` is the event or censoring indicator. `event` should include 0, 1 or 2, denoting right-censoring, failure from cause 1 and failure from cause 2, respectively. If `event=0` (i.e. right-censored observation) then  $u$  is not included in any calculation as it corresponds to  $\infty$ . The user can provide any value in  $u$  for the right-censored cases, even NA. The function `ciregic_lt` fits models that belong to the class of generalized odds rate transformation models which includes the proportional subdistribution hazards or the Fine-Gray model and the proportional odds model. The parameter  $\alpha = (\alpha_1, \alpha_2)$  defines the link function/model to be fitted for cause of failure 1 and 2, respectively. A value of 0 corresponds to the Fine-Gray model and a value of 1 corresponds to the proportional odds model. For example, if  $\alpha = (0, 1)$  then the function `ciregic_lt` fits the Fine-Gray model for cause 1 and the proportional odds model for cause 2.

## Value

The function `ciregic_lt` provides an object of class `ciregic_lt` with components:

<code>varnames</code>	a vector containing variable names
<code>coefficients</code>	a vector of the regression coefficient estimates
<code>gamma</code>	a vector of the estimated coefficients for the B-splines
<code>vcov</code>	a variance-covariance matrix of the estimated regression coefficients
<code>alpha</code>	a vector of the link function parameters
<code>loglikelihood</code>	a loglikelihood of the fitted model
<code>convergence</code>	an indicator of convergence
<code>tms</code>	a vector of the minimum and maximum observation times
<code>Bv</code>	a list containing the B-splines basis functions evaluated at $v$
<code>numboot</code>	a number of converged bootstrap
<code>notconverged</code>	a list of number of bootstrap samples that did not converge
<code>call</code>	a matched call

## Author(s)

Jun Park, <jun.park@alumni.iu.edu>  
Giorgos Bakoyannis, <gbakogia@iu.edu>

## References

- Bakoyannis, G., Yu, M., and Yiannoutsos C. T. (2017). Semiparametric regression on cumulative incidence function with interval-censored competing risks data. *Statistics in Medicine*, **36**:3683-3707.
- Fine, J. P. and Gray, R. J. (1999). A proportional hazards model for the subdistribution of a competing risk. *Journal of the American Statistical Association*, **94**:496-509.

**See Also**

[summary.ciregic\\_lt](#) for the summarized results and [predict.ciregic\\_lt](#) for value of the predicted cumulative incidence functions. `coef` and `vcov` are the generic functions. [dataprep](#) for reshaping data from a long format to a suitable format to be used in the function `ciregic_lt`.

**Examples**

```
## Not run:
## Set seed in order to have reproducibility of the bootstrap standard error estimate
set.seed(1234)

## Reshaping data from a long format to a suitable format
newdata <- dataprep_lt(data = longdata_lt, ID = id, time = t, W = w,
                      event = c, Z = c(z1, z2))
## Estimation of regression parameters only. No bootstrap variance estimation.
## with 'newdata'
fit_lt <- ciregic_lt(formula = Surv2(v = v, u = u, w = w, event = c) ~ z1 + z2, data = newdata,
                    alpha = c(1, 1), nboot = 0, do.par = FALSE)
fit_lt

## Bootstrap variance estimation based on 50 replications
fit_lt <- ciregic_lt(formula = Surv2(v = v, u = u, w = w, event = c) ~ z1 + z2, data = newdata,
                    alpha = c(1, 1), nboot = 50, do.par = FALSE)

## End(Not run)
## Note that the user can use parallel computing to decrease
## the computation time of the bootstrap variance-covariance
## estimation (e.g. nboot = 50)

## Summarize semiparametric regression model
summary(fit_lt)

## Predict and draw plot the cumulative incidence function evaluated at z1 = 1 and z2 = 0.5
mint <- fit_lt$tms[1]
maxt <- fit_lt$tms[2]
pred <- predict(object = fit_lt, covp = c(1, 0.5),
               times = seq(mint, maxt, by = (maxt - mint) / 99))
pred
plot(pred$t, pred$cif1, type = "l", ylim = c(0, 1))
points(pred$t, pred$cif2, type = "l", col = 2)
```

**Description**

The function `dataprep` reshapes data from a long format to a ready-to-use format to be used directly in the function `ciregic`.

**Usage**

```
dataprep(data, ID, time, event, Z)
```

**Arguments**

data	a data frame that includes the variables named in the ID, time, event, and z arguments
ID	a variable indicating individuals' ID
time	a variable indicating observed time points
event	a vector of event indicator. If an observation is right-censored, event = 0; otherwise, event = 1 or event = 2, where 1 represents the first cause of failure, and 2 represents the second cause of failure. The current version of package only allows two causes of failure.
Z	a vector of variables indicating name of covariates

**Details**

The function `dataprep` provides a ready-to-use data format that can be directly used in the function `ciregic`. The returned data frame consists of `id`, `v`, `u`, `c`, and `covariates` as columns. The `v` and `u` indicate time window with the last observation time before the event and the first observation after the event. The `c` represents a type of event, for example, `c = 1` for the first cause of failure, `c = 2` for the second cause of failure, and `c = 0` for the right-censored. For individuals having one time record with the event, the lower bound `v` will be replaced by zero, for example  $(0, v]$ . For individuals having one time record without the event, the upper bound `u` will be replaced by `Inf`, for example  $(v, \text{Inf}]$ .

**Value**

a data frame

**Author(s)**

Jun Park, <jun.park@alumni.iu.edu>

Giorgos Bakoyannis, <gbakogia@iu.edu>

**Examples**

```
library(intccr)
dataprep(data = longdata, ID = id, time = t, event = c, Z = c(z1, z2))
```

---

dataprep_lt	<i>Data preparation</i>
-------------	-------------------------

---

### Description

The function `dataprep_lt` reshapes data from a long format to a ready-to-use format to be used directly in the function `ciregic_lt`.

### Usage

```
dataprep_lt(data, ID, W, time, event, Z)
```

### Arguments

<code>data</code>	a data frame that includes the variables named in the <code>ID</code> , <code>time</code> , <code>event</code> , and <code>Z</code> arguments
<code>ID</code>	a variable indicating individuals' ID
<code>W</code>	a vector of left-truncated time points
<code>time</code>	a variable indicating observed time points
<code>event</code>	a vector of event indicator. If an observation is right-censored, <code>event = 0</code> ; otherwise, <code>event = 1</code> or <code>event = 2</code> , where 1 represents the first cause of failure, and 2 represents the second cause of failure. The current version of package only allows two causes of failure.
<code>Z</code>	a vector of variables indicating name of covariates

### Details

The function `dataprep_lt` provides a ready-to-use data format that can be directly used in the function `ciregic_lt`. The returned data frame consists of `id`, `v`, `u`, `c`, and `covariates` as columns. The `v` and `u` indicate time window with the last observation time before the event and the first observation after the event. The `c` represents a type of event, for example, `c = 1` for the first cause of failure, `c = 2` for the second cause of failure, and `c = 0` for the right-censored. For individuals having one time record with the event, the lower bound `v` will be replaced by zero, for example  $(0, v]$ . For individuals having one time record without the event, the upper bound `u` will be replaced by `Inf`, for example  $(v, \text{Inf}]$ .

### Value

a data frame

### Author(s)

Jun Park, <jun.park@alumni.iu.edu>

Giorgos Bakoyannis, <gbakogia@iu.edu>

---

dbs *Derivative of B-spline*

---

**Description**

Generates the derivative of the B-splines basis matrix.

**Usage**

```
dbs(  
  x,  
  derivs = 1L,  
  df = NULL,  
  knots = NULL,  
  degree = 3L,  
  intercept = FALSE,  
  Boundary.knots = range(x, na.rm = TRUE)  
)
```

**Arguments**

x	object of B-splines
derivs	a number of derivatives
df	degrees of freedom of B-splines
knots	a vector of internal knots
degree	degrees of B-splines
intercept	a logical vector
Boundary.knots	a vector of boundary knots

**Details**

The function `dbs` performs derivatives of B-splines

**Value**

The function `dbs` returns a component:

dMat	B-spline matrix
------	-----------------

**Author(s)**

Jun Park, <jun.park@alumni.iu.edu>

Giorgos Bakoyannis, <gbakogia@iu.edu>

---

fit	<i>Output of ciregic</i>
-----	--------------------------

---

**Description**

Object contains the output of the function `ciregic`. Standard errors were estimated by the least-squares method.

**Usage**

```
fit
```

**Format**

A list of components.

**Examples**

```
fit
```

---

fit_aipw	<i>Output of ciregic_aipw</i>
----------	-------------------------------

---

**Description**

A list of outputs containing the last time prior to the event, the first time after the event, cause of failure with 50% of missingness, and covariates.

**Usage**

```
fit_aipw
```

**Format**

A list of 14:

**call** a matched call

**varnames** a vector containing variable names

**varnames.aux** a vector containing auxiliary variable names

**coefficients** a vector of the regression coefficient estimates

**gamma** a vector of the estimated coefficients for the B-splines

**vcov** a variance-covariance matrix of the estimated regression coefficients

**alpha** a vector of the link function parameters

**k** a parameter that controls the number of knots in the B-spline

**loglikelihood** a loglikelihood of the fitted model  
**convergence** an indicator of convergence  
**tms** a vector of the minimum and maximum observation times  
**Bv** a list containing the B-splines basis functions evaluated at v  
**notconverged** a list of number of bootstrap samples not converged

### Examples

```
fit_aipw
```

---

fit_lt	<i>Output of ciregic_lt</i>
--------	-----------------------------

---

### Description

Object contains the output of the function `ciregic_lt`. Standard errors were estimated by the least-squares method.

### Usage

```
fit_lt
```

### Format

A list of components.

### Examples

```
fit_lt
```

---

longdata	<i>Simulated interval-censored competing risks data - long format</i>
----------	---

---

### Description

The data containing the subject id, series of time points, cause of failure, and covariates with 200 observations.

### Usage

```
longdata
```

### Format

A data frame with 868 rows and 5 variables.



**Examples**

```
library(intccr)
data(longdata)
```

---

longdata_lt	<i>Simulated left-truncated and interval-censored competing risks data - long format</i>
-------------	--

---

**Description**

Data containing observation time points, a left-truncation time, cause of failure, and baseline covariates with 275 observations.

**Usage**

```
longdata_lt
```

**Format**

A data frame with 275 unique individuals and 6 variables.

**Examples**

```
library(intccr)
data(longdata_lt)
```

---

naive_b	<i>Initial values for the sieve maximum likelihood estimation</i>
---------	---

---

**Description**

The function naive\_b provides a vector of initial values for the B-spline sieve maximum likelihood estimation.

**Usage**

```
naive_b(data, w = NULL, v, u, c, q, k = 1)
```

**Arguments**

data	a data frame that includes the variables named in each argument
w	a left-truncation time (default is w = NULL.)
v	the last observation time prior to the failure
u	the first observation time after the failure
c	an indicator of cause of failure, for example, if an observation is right-censored, event = 0; otherwise, event = 1 or event = 2, where 1 represents the first cause of failure, and 2 represents the second cause of failure. The current version of package only allows for two causes of failure.
q	a number of parameters in design matrix
k	a parameter that controls the number of knots in the B-spline with $0.5 \leq k \leq 1$

**Details**

The function naive\_b provides initial values for the optimization procedure.

**Value**

Initial values of B-spline estimation

b a vector of the initial values to be used in the optimization process

**Author(s)**

Giorgos Bakoyannis, <gbakogia@iu.edu>

Jun Park, <jun.park@alumni.iu.edu>

**Examples**

```
attach(simdata)
intccr::naive_b(data = simdata, v = v, u = u, c = c, q = 2)
```

---

predict.ciregic

*Covariate-Specific Cumulative Incidence Prediction*

---

**Description**

predict method for class ciregic. It provides the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Usage**

```
## S3 method for class 'ciregic'
predict(object, covp, times, ...)
```

**Arguments**

object	an object of class ciregic, which is a result of a call to ciregic
covp	a desired values for covariates
times	time points that user wants to predict value of cumulative incidence function
...	further arguments

**Details**

predict.ciregic returns the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Value**

The function predict.ciregic returns a list of predicted values of the model from object.

t	time points
cif1	the predicted value of cumulative incidence function for the event type 1
cif2	the predicted value of cumulative incidence function for the event type 2

**See Also**

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic](#) and summary of the fitted semiparametric regression model [summary.ciregic](#)

**Examples**

```
## Continuing the ciregic(...) example
pfit <- predict(object = fit, covp = c(1, 0.5), times = c(0.1, 0.15, 0.5, 0.7))
pfit
mint <- fit$tms[1]
maxt <- fit$tms[2]
pfit1 <- predict(object = fit, covp = c(1, 0.5),
                times = seq(mint, maxt, by = (maxt-mint)/99))
plot(pfit1$t, pfit1$cif1, ylim = c(0, 1), type = "l")
lines(pfit1$t, pfit1$cif2, ylim = c(0, 1), lty = 2, col = 2)
```

---

predict.ciregic\_aipw *Covariate-Specific Cumulative Incidence Prediction*

---

**Description**

predict method for class ciregic\_aipw. It provides the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Usage**

```
## S3 method for class 'ciregic_aipw'
predict(object, covp, times, ...)
```

**Arguments**

object	an object of class <code>ciregic_aipw</code> , which is a result of a call to <code>ciregic_aipw</code>
covp	a desired values for covariates
times	time points that user wants to predict value of cumulative incidence function
...	further arguments

**Details**

`predict.ciregic_aipw` returns the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Value**

The function `predict.ciregic_aipw` returns a list of predicted values of the model from object.

t	time points
cif1	the predicted value of cumulative incidence function for the event type 1
cif2	the predicted value of cumulative incidence function for the event type 2

**See Also**

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic\\_aipw](#) and summary of the fitted semiparametric regression model [summary.ciregic\\_aipw](#)

**Examples**

```
## Continuing the ciregic_aipw(...) example
pfit <- predict(object = fit_aipw, covp = c(1, 0.5), times = c(0.1, 0.15, 0.5, 0.7))
pfit
mint <- fit_aipw$tms[1]
maxt <- fit_aipw$tms[2]
pfit1 <- predict(object = fit_aipw, covp = c(1, 0.5),
                times = seq(mint, maxt, by = (maxt - mint) / 99))
plot(pfit1$t, pfit1$cif1, ylim = c(0, 1), type = "l")
lines(pfit1$t, pfit1$cif2, ylim = c(0, 1), lty = 2, col = 2)
```

---

predict.ciregic\_lt      *Covariate-Specific Cumulative Incidence Prediction*

---

**Description**

predict method for class `ciregic_lt`. It provides the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Usage**

```
## S3 method for class 'ciregic_lt'
predict(object, covp, times, ...)
```

**Arguments**

object	an object of class <code>ciregic_lt</code> , which is a result of a call to <code>ciregic_lt</code>
covp	a desired values for covariates
times	time points that user wants to predict value of cumulative incidence function
...	further arguments

**Details**

`predict.ciregic_lt` returns the predicted cumulative incidence function for a given covariate pattern and timepoint(s).

**Value**

The function `predict.ciregic_lt` returns a list of predicted values of the model from object.

t	time points
cif1	the predicted value of cumulative incidence function for the event type 1
cif2	the predicted value of cumulative incidence function for the event type 2

**See Also**

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic\\_lt](#) and summary of the fitted semiparametric regression model [summary.ciregic\\_lt](#)

**Examples**

```
## Continuing the ciregic_lt(...) example
pfit <- predict(object = fit_lt, covp = c(1, 0.5), times = c(0.1, 0.15, 0.5, 0.7))
pfit
mint <- fit_lt$tms[1]
maxt <- fit_lt$tms[2]
pfit1 <- predict(object = fit_lt, covp = c(1, 0.5),
                times = seq(mint, maxt, by = (maxt - mint) / 99))
plot(pfit1$t, pfit1$cif1, ylim = c(0, 1), type = "l")
lines(pfit1$t, pfit1$cif2, ylim = c(0, 1), lty = 2, col = 2)
```

---

predict.dbs

*Prediction of derivative of B-spline*

---

**Description**

Evaluates the derivative of the B-splines basis matrix at given values.

**Usage**

```
## S3 method for class 'dbs'
predict(object, newx)
```

**Arguments**

object	returned object of B-splines
newx	a vector of points

**Details**

The function `predict` is a generic function of `bs.derivs`

**Value**

The function `predict` returns a predicted B-splines.

**Author(s)**

Giorgos Bakoyannis, <gbakogia@iu.edu>

Jun Park, <jp84@alumni.iu.edu>

---

<code>pseudo.HIV.long</code>	<i>Artificial HIV dataset</i>
------------------------------	-------------------------------

---

**Description**

Artificial dataset that was simulated to resemble the HIV study on loss to HIV care and death in sub-Saharan Africa, that was presented in Bakoyannis, Yu, & Yiannoutsos (2017). It contains subject id, observation times, cause of failure, and covariates.

**Usage**

```
pseudo.HIV.long
```

**Format**

A data frame with 22710 rows and 6 variables.

**References**

Bakoyannis, G., Yu, M., and Yiannoutsos C. T. (2017). Semiparametric regression on cumulative incidence function with interval-censored competing risks data. *Statistics in Medicine*, **36**:3683-3707.

**Examples**

```
head(pseudo.HIV.long, n = 20)
```

---

simdata	<i>Simulated interval-censored competing risks data with 2 covariates - wide format</i>
---------	---

---

**Description**

The data containing the individual identification number, the last time point prior to the event, the first time point after the event, cause of failure, and covariates with 200 observations.

**Usage**

```
simdata
```

**Format**

A data frame with 200 rows and 6 variables.

**id** subject id

**v** the last observation time prior to the event

**u** the first observation time after the event

**c** cause of failure with missing

**z1** binary variable

**z2** continuous variable

**Examples**

```
library(intccr)
data(simdata)
```

---

simdata_aipw	<i>Simulated interval censored data with 2 covariates in the presence of 50% of missing cause of failure - wide format</i>
--------------	--

---

**Description**

The dataset containing the individual identification number, the last time prior to the event, the first time after the event, cause of failure with 50% of missingness, and covariates.

**Usage**

```
simdata_aipw
```

**Format**

A data frame with 200 rows and 7 variables:

**id** subject id  
**v** the last observation time prior to the event  
**u** the first observation time after the event  
**c** cause of failure with missing  
**z1** binary variable  
**z2** continuous variable  
**a** auxiliary variable

**Examples**

```
library(intccr)
data(simdata_aipw)
```

---

simdata_lt	<i>Simulated left-truncated and interval-censored competing risks data with 2 covariates - wide format</i>
------------	--

---

**Description**

The data containing the individual identification number, the left-truncated time, the last and first observation time prior to the event and after the event, cause of failure, and baseline covariates with 275 observations.

**Usage**

```
simdata_lt
```

**Format**

A data frame with 275 unique individuals and 7 variables.

**id** subject id  
**w** the left truncation time  
**v** the last observation time prior to the event  
**u** the first observation time after the event  
**c** cause of failure with missing  
**z1** binary variable  
**z2** continuous variable

**Examples**

```
library(intccr)
data(simdata_lt)
```



---

summary.ciregic	<i>Summary of ciregic</i>
-----------------	---------------------------

---

## Description

summary method for class ciregic

## Usage

```
## S3 method for class 'ciregic'  
summary(object, ...)
```

## Arguments

object	an object of class ciregic, which is a result of a call to ciregic
...	further arguments

## Details

The function `summary.ciregic` returns the coefficients, bootstrap standard errors, and etc. Additionally, 'significance star' is included.

## Value

The function `summary.ciregic` returns a list of summary statistics of the model from object.

varnames	a vector containing variable names
coefficients	a vector of the regression coefficient estimates
se	a bootstrap standard error of the coefficients
z	z value of the estimated coefficients
p	p value of the estimated coefficients
call	a matched call

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data `ciregic` and values of the predicted cumulative incidence functions `predict.ciregic`

## Examples

```
## Continuing the ciregic(...) example  
sfit <- summary(fit)  
sfit
```

---

summary.ciregic\_aipw *Summary of ciregic\_aipw*

---

## Description

summary method for class ciregic\_aipw

## Usage

```
## S3 method for class 'ciregic_aipw'  
summary(object, ...)
```

## Arguments

object            an object of class ciregic\_aipw, which is a result of a call to ciregic\_aipw  
...               further arguments

## Details

The function `summary.ciregic_aipw` returns the coefficients, bootstrap standard errors, and etc. Additionally, 'significance star' is included.

## Value

The function `summary.ciregic_aipw` returns a list of summary statistics of the model from object.

varnames	a vector containing variable names
coefficients	a vector of the regression coefficient estimates
se	a bootstrap standard error of the coefficients
z	z value of the estimated coefficients
p	p value of the estimated coefficients
call	a matched call

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data `ciregic_aipw` and values of the predicted cumulative incidence functions `predict.ciregic_aipw`

## Examples

```
## Continuing the ciregic_aipw(...) example  
sfit <- summary(fit_aipw)  
sfit
```

---

summary.ciregic\_lt      *Summary of ciregic\_lt*

---

### Description

summary method for class ciregic\_lt

### Usage

```
## S3 method for class 'ciregic_lt'  
summary(object, ...)
```

### Arguments

object            an object of class ciregic\_lt, which is a result of a call to ciregic\_lt  
...               further arguments

### Details

The function `summary.ciregic_lt` returns the coefficients, bootstrap standard errors, and etc. Additionally, 'significance star' is included.

### Value

The function `summary.ciregic_lt` returns a list of summary statistics of the model from object.

varnames        a vector containing variable names  
coefficients    a vector of the regression coefficient estimates  
se               a bootstrap standard error of the coefficients  
z                z value of the estimated coefficients  
p                p value of the estimated coefficients  
call            a matched call

### See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data `ciregic_lt` and values of the predicted cumulative incidence functions `predict.ciregic_lt`

### Examples

```
## Continuing the ciregic_lt(...) example  
sfit_lt <- summary(fit_lt)  
sfit_lt
```

---

`Surv2`*Creating data frame*

---

**Description**

The function `Surv2` generates the survival object to be treated as the response from `ciregic`.

**Usage**

```
Surv2(v, u, w = NULL, sub = NULL, event)
```

**Arguments**

<code>v</code>	the last observation time prior to the failure; $0 \leq v \leq u$
<code>u</code>	the first observation time after the failure; $u \geq 0$
<code>w</code>	a left truncation time or delayed entry time. The default setting is <code>w = NULL</code> for non left-truncated data.
<code>sub</code>	an indicator variable in the data set. It is an optional argument for interval-censored competing risks data and missing cause of failure, and the default is <code>NULL</code> . <code>sub = 1</code> for the observations that are subject to missingness and <code>sub = 0</code> elsewhere.
<code>event</code>	an indicator of cause of failure. If an observation is right-censored, <code>event = 0</code> ; otherwise, <code>event = 1</code> or <code>event = 2</code> , where 1 represents the first cause of failure, and 2 represents the second cause of failure. The current version of package only allows for two causes of failure.

**Details**

The function `Surv2` provides a response data frame which is used in the function `ciregic` and `ciregic_lt`. For interval-censored competing risks data, the function `Surv2` must use three parameters (`v`, `u`, `c`). For left-truncated and interval censored competing risks data, the function `Surv2` must use four parameters (`v`, `u`, `w`, `c`). If data are partially left-truncated, but all interval-censored, `w = 0` for only interval-censored competing risks data.

**Value**

data frame

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## Examples

```
attach(simdata)
Surv2(v = v, u = u, event = c)
attach(simdata_lt)
Surv2(v = v, u = u, w = w, event = c)
```

---

vcov.ciregic	<i>Variance-covariance matrix of ciregic</i>
--------------	--

---

## Description

vcov method for class ciregic

## Usage

```
## S3 method for class 'ciregic'
vcov(object, ...)
```

## Arguments

object	an object of class ciregic, which is a result of a call to ciregic
...	further arguments

## Details

The function vcov returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic](#), summary of the fitted semiparametric regression model [summary.ciregic](#), and values of predicted cumulative incidence functions [predict.ciregic](#)

## Examples

```
## Continuing the ciregic(...) example
vcov(fit)
```

---

vcov.ciregic_aipw	<i>Variance-covariance matrix of ciregic_aipw</i>
-------------------	---

---

## Description

vcov method for class ciregic\_aipw

## Usage

```
## S3 method for class 'ciregic_aipw'  
vcov(object, ...)
```

## Arguments

object	an object of class ciregic_aipw, which is a result of a call to ciregic_aipw
...	further arguments

## Details

The function vcov returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic\\_aipw](#), summary of the fitted semiparametric regression model [summary.ciregic\\_aipw](#), and values of predicted cumulative incidence functions [predict.ciregic\\_aipw](#)

## Examples

```
## Continuing the ciregic_aipw(...) example  
vcov(fit_aipw)
```

---

vcov.ciregic_lt	<i>Variance-covariance matrix of ciregic_lt</i>
-----------------	---

---

## Description

vcov method for class ciregic\_lt

## Usage

```
## S3 method for class 'ciregic_lt'  
vcov(object, ...)
```

## Arguments

object	an object of class ciregic_lt, which is a result of a call to ciregic_lt
...	further arguments

## Details

The function vcov returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic\\_lt](#), summary of the fitted semiparametric regression model [summary.ciregic\\_lt](#), and values of predicted cumulative incidence functions [predict.ciregic\\_lt](#)

## Examples

```
## Continuing the ciregic_lt(...) example  
vcov(fit_lt)
```

---

vcov.summary.ciregic *Variance-covariance matrix of summary.ciregic*

---

## Description

vcov method for class `summary.ciregic`

## Usage

```
## S3 method for class 'summary.ciregic'  
vcov(object, ...)
```

## Arguments

`object` an object of class `summary.ciregic`, which is a result of a call to `ciregic`  
`...` further arguments

## Details

The `vcov` returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data `ciregic`, summary of the fitted semiparametric regression model `summary.ciregic`, and values of the predicted cumulative incidence functions `predict.ciregic`

## Examples

```
## Continuing the ciregic(...) example  
vcov(summary(fit))
```



---

```
vcov.summary.ciregic_aipw
```

*Variance-covariance matrix of summary.ciregic\_aipw*

---

## Description

vcov method for class `summary.ciregic_aipw`

## Usage

```
## S3 method for class 'summary.ciregic_aipw'  
vcov(object, ...)
```

## Arguments

<code>object</code>	an object of class <code>summary.ciregic_aipw</code> , which is a result of a call to <code>ciregic_aipw</code>
<code>...</code>	further arguments

## Details

The `vcov` returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic\\_aipw](#), summary of the fitted semiparametric regression model [summary.ciregic\\_aipw](#), and values of the predicted cumulative incidence functions [predict.ciregic\\_aipw](#)

## Examples

```
## Continuing the ciregic_aipw(...) example  
vcov(summary(fit_aipw))
```

---

`vcov.summary.ciregic_lt`*Variance-covariance matrix of summary.ciregic\_lt*

---

## Description

vcov method for class `summary.ciregic_lt`

## Usage

```
## S3 method for class 'summary.ciregic_lt'  
vcov(object, ...)
```

## Arguments

`object` an object of class `summary.ciregic_lt`, which is a result of a call to `ciregic_lt`  
`...` further arguments

## Details

The `vcov` returns the variance-covariance matrix of the fitted semiparametric regression model.

## Value

The estimated bootstrap variance-covariance matrix

## See Also

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data `ciregic_lt`, summary of the fitted semiparametric regression model `summary.ciregic_lt`, and values of the predicted cumulative incidence functions `predict.ciregic_lt`

## Examples

```
## Continuing the ciregic_lt(...) example  
vcov(summary(fit_lt))
```

---

waldtest	<i>Wald test for ciregic and ciregic_lt</i>
----------	---

---

**Description**

waldtest for class ciregic or ciregic\_lt. This provides the result of Wald test for the fitted model from the function ciregic or ciregic\_lt.

**Usage**

```
waldtest(obj1, obj2 = NULL, ...)
```

**Arguments**

obj1	an object of the fitted model in ciregic or ciregic_lt
obj2	an object of the fitted model in ciregic or ciregic_lt, the default is NULL
...	further arguments

**Details**

The function waldtest.ciregic returns a result of Wald test.

**Value**

The function waldtest returns an output table of Wald test of the model from object.

varnames.full	a variable name of a vector of variables names in the full model
varnames.nested	a variable name of a vector of variables names in the nested model
vcov	the estimated bootstrap variance-covariance matrix for overall Wald test
vcov.event1	the estimated bootstrap variance-covariance matrix for cause-specific Wald test (event type 1)
vcov.event2	the estimated bootstrap variance-covariance matrix for cause-specific Wald test (event type 2)
table	a table including test statistic, degrees of freedom, and p-value

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**See Also**

The fitted semiparametric regression on cumulative incidence function with interval-censored competing risks data [ciregic](#) and left-truncated and interval-censored competing risks data [ciregic\\_lt](#)

**Examples**

```
## Continuing the ciregic(...) example
library(intccr)
waldtest(obj1 = fit)
set.seed(12345)
newdata <- dataprep(data = longdata, ID = id, time = t,
                    event = c, Z = c(z1, z2))
fit.nested <- ciregic(formula = Surv2(v = v, u = u, event = c) ~ z2, data = newdata,
                    alpha = c(1, 1), nboot = 0, do.par = FALSE)
waldtest(obj1 = fit, obj2 = fit.nested)
```

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