

# Package ‘WAreg’

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**Type** Package

**Title** While-Alive Regression for Composite Endpoints with  
Cluster-Robust Inference

**Version** 0.1.0

**Description** Provides estimation and inference for while-alive regression models targeting the while-alive loss rate for composite endpoints that include recurrent events and a terminal event. The implementation supports flexible time-varying covariate effects through user-selected time bases, including B-splines, natural splines, M-splines, step functions, truncated linear bases, interval-local bases, and piecewise polynomials. Inference can be performed using cluster-robust variance estimators for cluster-randomized trials, with subject-level (IID) variance as a special case. The package includes prediction and plotting utilities and K-fold cross-validation for selecting basis and tuning parameters. Methodology is based on Fang et al. (2025) <[doi:10.1093/biostatistics/kxaf047](https://doi.org/10.1093/biostatistics/kxaf047)>.

**License** GPL-3

**Encoding** UTF-8

**Depends** R (>= 4.1)

**Imports** dplyr, tidyr, tibble, ggplot2, survival, nleqslv, splines,  
MASS, magrittr, rlang

**Suggests** splines2, testthat (>= 3.0.0), knitr, rmarkdown

**Config/testthat/edition** 3

**URL** <https://github.com/fancy575/WAreg>

**BugReports** <https://github.com/fancy575/WAreg/issues>

**RoxygenNote** 7.3.3

**LazyData** true

**NeedsCompilation** no

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**Repository** CRAN

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crt_dt	<i>Clustered Recurrent-Time Dataset: crt_dt</i>
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### Description

A simulated dataset of clustered recurrent events with terminal/censoring outcomes and covariates, suitable for examples and tests.

### Usage

```
data(crt_dt)
```

### Format

A data frame with the following columns:

**id** Integer subject ID (within the whole sample).

**cluster** Integer cluster ID.

**time** Numeric event/censoring time.

**status** Integer event type indicator: 0 = censored, 1 = recurrent type 1, 2 = recurrent type 2, 3 = death (terminal).

**trt** Cluster-level treatment indicator carried to subjects (e.g., 0/1).

**Z1** Numeric covariate.

**Z2** Numeric covariate.

### Details

Rows represent observed events (including censoring and death) for each subject. Multiple rows per id indicate multiple recurrent events; terminal/censoring rows mark the end of observation for that subject.

### Source

Generated by the package's simulation utilities.

**Examples**

```
data(crt_dt)
head(crt_dt)
```

---

`irt_dt`*Individual Recurrent-Time Dataset: irt\_dt*

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**Description**

A simulated dataset of recurrent events with terminal/censoring outcomes and covariates, organized in long format.

**Usage**

```
data(irt_dt)
```

**Format**

A data frame with the following columns:

**id** Integer subject ID (within the whole sample).

**time** Numeric event/censoring time.

**status** Integer event type indicator: 0 = censored, 1 = recurrent type 1, 2 = recurrent type 2, 3 = death (terminal).

**trt** Cluster-level treatment indicator carried to subjects (e.g., 0/1).

**Z1** Numeric covariate.

**Z2** Numeric covariate.

**Details**

Long-format events: each row is an event (or censoring/death) for a subject.

**Source**

Generated by the package's simulation utilities.

**Examples**

```
data(irt_dt)
head(irt_dt)
```

plot.WA

*Plot while-alive trajectory or a covariate's time-varying effect***Description**

Plot while-alive trajectory or a covariate's time-varying effect

**Usage**

```
## S3 method for class 'WA'
plot(
  x,
  newdata,
  t_seq,
  id = 1,
  mode = c("wa", "cov"),
  covariate = NULL,
  ylab_wa = "While-alive loss rate",
  ylab_cov = NULL,
  xlab = "Time",
  level = 0.95,
  smooth = FALSE,
  span = 0.3,
  ...
)
```

**Arguments**

x	A "WA" object.
newdata	Data used to rebuild the RHS design (same columns as in the model).
t_seq	Times to plot over (numeric vector).
id	Row index of newdata to use for the while-alive trajectory (mode = "wa").
mode	"wa" to plot the while-alive loss rate, or "cov" to plot a specific covariate's time-varying effect.
covariate	Character; covariate name (must appear on RHS) when mode="cov".
ylab_wa	Y-axis label for while-alive plot.
ylab_cov	Y-axis label for covariate-effect plot; default "Effect of <covariate> on $\beta(t)$ ".
xlab	X-axis label.
level	Confidence level for ribbons (default 0.95).
smooth	Logical; if TRUE, apply LOESS smoothing to the displayed curve/CI.
span	LOESS span used when smooth=TRUE.
...	Unused.

**Value**

A **ggplot2** object.

**Examples**

```
ex_dt <- crt_dt[crt_dt$cluster %in% c(1,2,3,4,7,10), ]
fit <- WA_fit(survival::Surv(time, status) ~ trt + Z1 + Z2,
             data = ex_dt, id="id", cluster="cluster",
             knots=seq(0, max(ex_dt$time), length.out=6),
             tau_grid=seq(0, max(ex_dt$time), length.out=6),
             basis="bz", degree=1, link="log",
             w_recur=c(1,1), w_term=2, ipcw="km")
nd <- unique(ex_dt[, c("trt", "Z1", "Z2")])
plot(fit, newdata = nd,
     t_seq = seq(0, max(fit$tau_grid), length.out = 200),
     id = 1, mode = "wa", smooth = TRUE)
```

---

predict.WA

*Predict while-alive loss rates*

---

**Description**

Predict while-alive loss rates

**Usage**

```
## S3 method for class 'WA'
predict(object, newdata, t_seq, level = 0.95, ...)
```

**Arguments**

object	A "WA" object.
newdata	Data frame with columns matching the RHS of the fitted model. Predictions are computed for the rows of newdata.
t_seq	Numeric vector of times at which to evaluate predictions.
level	Confidence level for pointwise intervals (default 0.95).
...	Unused.

**Value**

A data frame with columns `id` (row index in `newdata`), `t`, `mu` (predicted while-alive rate), and CI columns `lb`, `ub`.

**Examples**

```

ex_dt <- crt_dt[crt_dt$cluster %in% c(1,2,3,4,7,10), ]
fit <- WA_fit(survival::Surv(time, status) ~ trt + Z1 + Z2,
             data = ex_dt, id="id", cluster="cluster",
             knots=seq(0, max(ex_dt$time), length.out=6),
             tau_grid=seq(0, max(ex_dt$time), length.out=6),
             basis="bz", degree=1, link="log",
             w_recur=c(1,1), w_term=2, ipcw="km")
nd <- unique(ex_dt[, c("trt","Z1","Z2")])
pred <- predict(fit, newdata = nd, t_seq = seq(0, max(fit$tau_grid), by = 0.2))
head(pred)

```

---

summary.WA

*Summarize a WA object*


---

**Description**

Summarize a WA object

**Usage**

```

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

```

**Arguments**

object	A "WA" object from <a href="#">WA_fit</a> .
...	Unused.

**Value**

An object of class "summary.WA" containing configuration and a coefficient table with estimates, standard errors, and z-scores.

**Examples**

```

ex_dt <- crt_dt[crt_dt$cluster %in% c(1,2,3,4,7,10), ]
fit <- WA_fit(survival::Surv(time, status) ~ trt + Z1 + Z2,
             data = ex_dt, id="id", cluster="cluster",
             knots=seq(0, max(ex_dt$time), length.out=6),
             tau_grid=seq(0, max(ex_dt$time), length.out=6),
             basis="bz", degree=1, link="log",
             w_recur=c(1,1), w_term=2, ipcw="km")
summary(fit)

```

**Description**

Runs K-fold CV over a grid of basis types, degrees, interior-knot counts, and link functions. For each configuration, fits the model on K-1 folds and accumulates the prediction error (PE) on the held-out fold using `WA_PE()` (IPCW computed on the training subjects).

**Usage**

```
WA_cv(
  formula,
  data,
  id,
  cluster = NULL,
  basis_set = c("il", "pl", "bz"),
  degree_vec = 1:2,
  n_int_vec = c(0, 2, 4),
  knot_scheme = c("equidist", "quantile"),
  link_set = c("log"),
  time_range = NULL,
  tau_grid = NULL,
  w_recur,
  w_term,
  ipcw = c("cox", "km"),
  ipcw_formula = ~1,
  K = 5,
  seed = 1L,
  verbose = TRUE
)
```

**Arguments**

<code>formula</code>	A <code>Surv(time, status) ~ RHS formula</code> ; see <a href="#">WA_fit</a> .
<code>data</code>	Long-format data frame; see <a href="#">WA_fit</a> .
<code>id</code>	Character scalar; subject ID column name; see <a href="#">WA_fit</a> .
<code>cluster</code>	Optional character scalar; cluster column name; see <a href="#">WA_fit</a> .
<code>basis_set</code>	Character vector of candidate bases.
<code>degree_vec</code>	Integer vector of candidate degrees.
<code>n_int_vec</code>	Integer vector of interior-knot counts; 0 means boundaries only.
<code>knot_scheme</code>	"equidist" or "quantile" to construct interior knots.
<code>link_set</code>	Character vector of candidate links (subset of <code>c("log", "identity")</code> ).
<code>time_range</code>	Optional numeric length-2 vector <code>c(tmin, tmax)</code> . If NULL, inferred from data.

tau_grid	Optional numeric vector; if NULL, a default dense grid over time_range is created.
w_recur	recurrent-event weights
w_term	Numeric scalar; terminal-event weight; see <a href="#">WA_fit</a> .
ipcw	IPCW method ("cox" or "km") for PE computation.
ipcw_formula	One-sided RHS formula for IPCW Cox model (if ipcw="cox").
K	Number of folds.
seed	RNG seed for fold assignment.
verbose	Logical; show a text progress bar and per-fold messages.

**Value**

A data frame with columns: basis, degree, n\_int, link, and aggregated PE. Lower PE is better.

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WA_fit	<i>While-Alive Regression (WA) for Composite Endpoints</i>
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---

**Description**

Fits the while-alive regression model targeting the while-alive loss rate for composite endpoints with recurrent and terminal events. Time-varying covariate effects are represented via user-chosen time bases (e.g., B-spline, piecewise polynomial, interval-local). Robust inference supports cluster-randomized trials (CRTs) via cluster-robust variance; if cluster = NULL, IID (subject-as-cluster) variance is used.

**Usage**

```
WA_fit(
  formula,
  data,
  id,
  cluster = NULL,
  knots,
  tau_grid,
  basis = c("il", "pl", "bz", "ns", "ms", "st", "tl", "tf"),
  degree = 1,
  link = c("log", "identity"),
  w_recur,
  w_term,
  ipcw = c("km", "cox"),
  ipcw_formula = ~1
)
```

**Arguments**

formula	A Surv(time, status) ~ RHS formula. time and status must exist in data. The RHS contains baseline covariates (no explicit time-varying covariates here; time-variation is induced via the chosen basis).
data	Long-format data frame with one row per <i>event/checkpoint</i> per subject, containing time, status, id, optional cluster, and RHS covariates.
id	Character scalar; subject ID column name.
cluster	Optional character scalar; cluster column name for CRT-robust inference. If NULL, IID inference treats each subject as its own cluster.
knots	Numeric vector (length $\geq 2$ ) specifying the basis boundaries and optional interior knots that define the time basis shape.
tau_grid	Numeric vector of evaluation times used to stack the estimating equations. Independent of knots.
basis	One of "il", "pl", "bz", "ns", "ms", "st", "tl", "tf": interval-local ("il"), piecewise polynomial ("pl"), B-spline ("bz"), natural spline ("ns"), M-spline ("ms", requires <b>splines2</b> ), step ("st"), truncated linear ("tl"), or time-fixed ("tf").
degree	Integer degree for bases that use it (e.g., "bz", "pl", "ns", "ms").
link	Link function: "log" (default) or "identity".
w_recur	Numeric vector of weights for each recurrent event type. Its length must match the number of recurrent status codes in data (i.e., excluding 0 for censoring and the max code for terminal).
w_term	Numeric scalar; weight for the terminal event.
ipcw	IPCW method: "km" or "cox".
ipcw_formula	A one-sided formula specifying RHS covariates for the IPCW Cox model when ipcw = "cox" (e.g., ~ x1 + x2). Ignored for ipcw = "km".

**Details**

The estimating equations solve  $E[Z(t)\{L(t) - \mu_\beta(t)X_{\min}(t)\}V/G] = 0$  over tau\_grid, where  $L(t)$  is the weighted composite loss (recurrent+terminal),  $\mu_\beta(t)$  the while-alive loss rate under the chosen link,  $X_{\min}(t) = \min(T, t)$ ,  $V$  the at-risk/terminal indicator, and  $G$  the censoring survival modeled via ipcw.

**Value**

An object of class "WA" with elements:

- est: named coefficient vector.
- vcov: cluster-robust variance matrix.
- se: standard errors.
- converged: logical.
- basis, degree, link, Z\_cols, knots, tau\_grid, id\_var, cluster\_var, w\_recur, w\_term, status\_codes, formula.

**Examples**

```
ex_dt <- crt_dt[crt_dt$cluster %in% c(1,2,3,4,7,10), ]
fit <- WA_fit(
  survival::Surv(time, status) ~ trt + Z1 + Z2,
  data      = ex_dt,
  id        = "id",
  cluster   = "cluster",
  knots     = seq(0, max(ex_dt$time, na.rm = TRUE), length.out = 6),
  tau_grid  = seq(0, max(ex_dt$time, na.rm = TRUE), length.out = 6),
  basis     = "bz", degree = 1, link = "log",
  w_recur   = c(1, 1), w_term = 2,
  ipcw      = "km"
)
s <- summary(fit)
nd <- unique(ex_dt[, c("trt", "Z1", "Z2")])
plot(fit, newdata = nd,
     t_seq = seq(0, max(fit$tau_grid), length.out = 200),
     id = 1, mode = "wa", smooth = TRUE)
```

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