Package: survstan (via r-universe)

October 29, 2026

Title Fitting Survival Regression Models via 'Stan'

Version 0.0.7.1

Description Parametric survival regression models under the maximum likelihood approach via 'Stan'. Implemented regression models include accelerated failure time models, proportional hazards models, proportional odds models, accelerated hazard models, Yang and Prentice models, and extended hazard models. Available baseline survival distributions include exponential, Weibull, log-normal, log-logistic, gamma, generalized gamma, rayleigh, Gompertz and fatigue (Birnbaum-Saunders) distributions. References: Lawless (2002) <ISBN:9780471372158>; Bennett (1982) <doi:10.1002/sim.4780020223>; Chen and Wang(2000) <doi:10.1080/01621459.2000.10474236>; Demarqui and Mayrink (2021) <doi:10.1214/20-BJPS471>.

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.3.1

Biarch true

Depends R (>= 3.4.0), survival

Imports actuar (>= 3.0.0), broom, doFuture, dplyr, extraDistr, foreach, future, generics, ggplot2, gridExtra, MASS, methods, purrr, Rcpp (>= 0.12.0), RcppParallel (>= 5.0.1), Rdpack, rlang, rstan (>= 2.26.0), rstantools (>= 2.3.1), tibble, tidyr

RdMacros Rdpack

LinkingTo BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.0), StanHeaders (>= 2.26.0)

SystemRequirements GNU make

URL https://github.com/fndemarqui/survstan, https://fndemarqui.github.io/survstan/

BugReports https://github.com/fndemarqui/survstan/issues

Suggests emmeans (>= 1.4.2), estimability, GGally, knitr, rmarkdown, testthat (>= 3.0.0)

Config/testthat/edition 3

VignetteBuilder knitr

NeedsCompilation yes

Author Fabio Demarqui [aut, cre, cph] (<https://orcid.org/0000-0001-9236-1986>), Andrew Johnson [ctb]

Maintainer Fabio Demarqui <fndemarqui@est.ufmg.br>

Config/pak/sysreqs make

Repository https://cran.r-universe.dev

Date/Publication 2024-04-13 02:29:08 UTC

RemoteUrl https://github.com/cran/survstan

RemoteRef HEAD

RemoteSha 7c5aed0da4e22213fda1c9c45741841ea0cc31c0

Contents

survstan-package . . . 3
aftreg . . . 3
ahreg . . . 4
AIC.survstan . . . 5
anova.survstan . . . 6
coef.survstan . . . 7
confint.survstan . . . 7
cross_time . . . 8
cross_time.survstan . . . 9
ehreg . . . 10
emmeans-survstan-helpers . . . 11
estimates . . . 11
extractAIC.survstan . . . 12
gastric . . . 13
ggprentice . . . 13
ggresiduals . . . 15
ggresiduals.survstan . . . 15
ggstacy . . . 16
Gompertz . . . 17
ipass . . . 18
logLik.survstan . . . 19
model.matrix.survstan . . . 20
phreg . . . 20
poreg . . . 21
print.summary.survstan . . . 22
rank_models . . . 23
residuals.survstan . . . 24

se 24
se.survstan 25
summary.survstan 26
survfit.survstan 26
tidy.survstan 27
vcov.survstan 28
ypreg 28

Index **30**

`survstan-package` *The 'survstan' package.*

Description

The aim of the R package survstan is to provide a toolkit for fitting survival models using Stan. The R package survstan can be used to fit right-censored survival data under independent censoring. The implemented models allow the fitting of survival data in the presence/absence of covariates. All inferential procedures are currently based on the maximum likelihood (ML) approach.

_PACKAGE

References

Stan Development Team (2023). "RStan: the R interface to Stan." R package version 2.21.8, https://mc-stan.org/.

Lawless JF (2002). *Statistical Models and Methods for Lifetime Data*, Wiley Series in Probability and Statistics, 2nd Edition edition. John Wiley and Sons. ISBN 9780471372158.

Bennett S (1983). "Analysis of survival data by the proportional odds model." *Statistics in Medicine*, **2**(2), 273-277. doi:10.1002/sim.4780020223.

Chen YQ, Wang M (2000). "Analysis of Accelerated Hazards Models." *Journal of the American Statistical Association*, **95**(450), 608-618. doi:10.1080/01621459.2000.10474236.

Demarqui FN, Mayrink VD (2021). "Yang and Prentice model with piecewise exponential baseline distribution for modeling lifetime data with crossing survival curves." *Brazilian Journal of Probability and Statistics*, **35**(1), 172 – 186. doi:10.1214/20BJPS471.

`aftreg` *Fitting Accelerated Failure Time Models*

Description

Function to fit accelerated failure time (AFT) models.

Usage

```
aftreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.
baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

aftreg returns an object of class "aftreg" containing the fitted model.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

---

ahreg *Fitting Accelerated Hazard Models*

Description

Function to fit accelerated hazard (AH) models.

Usage

```
ahreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.

baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

ahreg returns an object of class "ahreg" containing the fitted model.

Examples

```
library(survstan)
fit <- ahreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

AIC.survstan *Akaike information criterion*

Description

Akaike information criterion

Usage

```
## S3 method for class 'survstan'
AIC(object, ..., k = 2)
```

Arguments

object	an object of the class survstan.
...	further arguments passed to or from other methods.
k	numeric, the penalty per parameter to be used; the default k = 2 is the classical AIC.

Value

the Akaike information criterion value when a single model is passed to the function; otherwise, a data.frame with the Akaike information criterion values and the number of parameters is returned.

Examples

```
library(survstan)
fit1 <- aftreg(Surv(futime, fustat) ~ 1, data = ovarian, baseline = "weibull", init = 0)
fit2 <- aftreg(Surv(futime, fustat) ~ rx, data = ovarian, baseline = "weibull", init = 0)
fit3 <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
AIC(fit1, fit2, fit3)
```

---

anova.survstan *anova method for survstan models*

Description

Compute analysis of variance (or deviance) tables for one or more fitted model objects.

Usage

```
## S3 method for class 'survstan'
anova(...)
```

Arguments

...	further arguments passed to or from other methods.

Value

the ANOVA table.

Examples

```
library(survstan)
fit1 <- aftreg(Surv(futime, fustat) ~ 1, data = ovarian, baseline = "weibull", init = 0)
fit2 <- aftreg(Surv(futime, fustat) ~ rx, data = ovarian, baseline = "weibull", init = 0)
fit3 <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
anova(fit1, fit2, fit3)
```

coef.survstan *Estimated regression coefficients*

Description

Estimated regression coefficients

Usage

```
## S3 method for class 'survstan'
coef(object, ...)
```

Arguments

object	an object of the class survstan
...	further arguments passed to or from other methods

Value

the estimated regression coefficients

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
coef(fit)
```

confint.survstan *Confidence intervals for the regression coefficients*

Description

Confidence intervals for the regression coefficients

Usage

```
## S3 method for class 'survstan'
confint(object, parm = NULL, level = 0.95, ...)
```

Arguments

object	an object of the class survstan.
parm	a specification of which parameters are to be given confidence intervals, either a vector of numbers or a vector of names. If missing, all parameters are considered.
level	the confidence level required.
...	further arguments passed to or from other methods.

Value

100(1-alpha) confidence intervals for the regression coefficients.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
confint(fit)
```

---

cross_time *Generic S3 method cross_time*

Description

Generic S3 method cross_time

Usage

```
cross_time(object, ...)
```

Arguments

object	a fitted model object
...	further arguments passed to or from other methods.

Value

the crossing survival time

`cross_time.survstan` *Computes the crossing survival times*

Description

Computes the crossing survival times

Usage

```
## S3 method for class 'survstan'
cross_time(
  object,
  newdata1,
  newdata2,
  conf.level = 0.95,
  nboot = 1000,
  cores = 1,
  ...
)
```

Arguments

`object`	an object of class survstan
`newdata1`	a data frame containing the first set of explanatory variables
`newdata2`	a data frame containing the second set of explanatory variables
`conf.level`	level of the confidence/credible intervals
`nboot`	number of bootstrap samples (default nboot=1000).
`cores`	number of cores to be used in the bootstrap sampling; default is 1 core;
`...`	further arguments passed to or from other methods.

Value

the crossing survival time

Examples

```
library(survstan)
data(ipass)
fit <- ypreg(Surv(time, status)~arm, data=ipass, baseline = "weibull")
summary(fit)
newdata1 <- data.frame(arm=0)
newdata2 <- data.frame(arm=1)
tcross <- cross_time(fit, newdata1, newdata2, nboot = 10)
tcross
```

ehreg *Fitting Extended Hazard Models*

Description

Function to fit Extended Hazard (EH) models.

Usage

```
ehreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.
baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

ehreg returns an object of class "ehreg" containing the fitted model.

Examples

```
library(survstan)
fit <- ehreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

emmeans-survstan-helpers *Support Functions for* **emmeans**

Description

Functions required for compatibility of **survstan** with **emmeans**. Users are not required to call these functions themselves. Instead, they will be called automatically by the `emmeans` function of the **emmeans** package.

Usage

```
recover_data.survstan(object, ...)

recover_data.ypreg(object, term = c("short", "long"), ...)

recover_data.ehreg(object, term = c("AF", "RH"), ...)
```

Arguments

`object`	An object of the same class as is supported by a new method.
`...`	Additional parameters that may be supported by the method.
`term`	character specifying whether AF or RH term regression coefficients are to be used.

estimates *Parameters estimates of a survstan model*

Description

Parameters estimates of a survstan model

Usage

```
estimates(object, ...)
```

Arguments

`object`	an object of the class survstan.
`...`	further arguments passed to or from other methods.

Value

the parameters estimates of a given survstan model.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
estimates(fit)
```

`extractAIC.survstan` *Extract AIC from a Fitted Model*

Description

Computes the (generalized) Akaike An Information Criterion for a fitted parametric model.

Usage

```
## S3 method for class 'survstan'
extractAIC(fit, scale, k = 2, ...)
```

Arguments

`fit`	a fitted model of the class survstan
`scale`	optional numeric specifying the scale parameter of the model. Currently only used in the "lm" method, where scale specifies the estimate of the error variance, and scale = 0 indicates that it is to be estimated by maximum likelihood.
`k`	numeric specifying the 'weight' of the equivalent degrees of freedom part in the AIC formula.
`...`	further arguments passed to or from other methods.

Value

the ANOVA table.

Examples

```
library(survstan)
fit1 <- aftreg(Surv(futime, fustat) ~ 1, data = ovarian, baseline = "weibull", init = 0)
fit2 <- aftreg(Surv(futime, fustat) ~ rx, data = ovarian, baseline = "weibull", init = 0)
fit3 <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
extractAIC(fit1)
extractAIC(fit2)
extractAIC(fit3)
```

gastric *Gastric cancer data set*

Description

Data set from a clinical trial conducted by the Gastrointestinal Tumor Study Group (GTSG) in 1982. The data set refers to the survival times of patients with locally nonresectable gastric cancer. Patients were either treated with chemotherapy combined with radiation or chemotherapy alone.

Format

A data frame with 90 rows and 3 variables:

- time: survival times (in days)
- status: failure indicator (1 - failure; 0 - otherwise)
- trt: treatments (1 - chemotherapy + radiation; 0 - chemotherapy alone)

Author(s)

Fabio N. Demarqui <fndemarqui@est.ufmg.br>

References

Gastrointestinal Tumor Study Group. (1982) A Comparison of Combination Chemotherapy and Combined Modality Therapy for Locally Advanced Gastric Carcinoma. Cancer 49:1771-7.

ggprentice *The Generalized Gamma Distribution (Prentice's alternative parametrization)*

Description

Probability function, distribution function, quantile function and random generation for the distribution with parameters mu, sigma and varphi.

Usage

```
dggprentice(x, mu, sigma, varphi, log = FALSE)

pggprentice(q, mu = 0, sigma = 1, varphi, lower.tail = TRUE, log.p = FALSE)

qggprentice(p, mu = 0, sigma = 1, varphi, lower.tail = TRUE, log.p = FALSE)

rggprentice(n, mu = 0, sigma = 1, varphi, ...)
```

Arguments

x	vector of (non-negative integer) quantiles.
mu	location parameter of the distribution.
sigma	scale parameter of the distribution (sigma > 0).
varphi	shape parameter of the distribution.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
q	vector of quantiles.
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$; otherwise, $P[X > x]$.
p	vector of probabilities.
n	number of random values to return.
...	further arguments passed to other methods.

Details

Probability density function:

$$f(x|\mu,\sigma,\varphi) = \begin{cases} \frac{|\varphi|(\varphi^{-2})^{\varphi^{-2}}}{\sigma x \Gamma(\varphi^{-2})} \exp\{\varphi^{-2}[\varphi w - \exp(\varphi w)]\} I_{[0,\infty)}(x), & \varphi \neq 0 \\ \frac{1}{\sqrt{2\pi}x\sigma} \exp\left\{-\frac{1}{2}\left(\frac{log(x)-\mu}{\sigma}\right)^2\right\} I_{[0,\infty)}(x), & \varphi = 0 \end{cases}$$

where $w = \frac{\log(x)-\mu}{\sigma}$, for $-\infty < \mu < \infty$, $\sigma > 0$ and $-\infty < \varphi < \infty$.

Distribution function:

$$F(x|\mu,\sigma,\varphi) = \begin{cases} F_G(y|1/\varphi^2,1), & \varphi > 0 \\ 1 - F_G(y|1/\varphi^2,1), & \varphi < 0 \\ F_{LN}(x|\mu,\sigma), & \varphi = 0 \end{cases}$$

where $y = \left(\frac{x}{\sigma}\right)^{\varphi}$, $F_G(\cdot|\nu,1)$ is the distribution function of a gamma distribution with shape parameter $1/\varphi^2$ and scale parameter equals to 1, and $F_{LN}(x|\mu,\sigma)$ corresponds to the distribution function of a lognormal distribution with location parameter μ and scale parameter σ.

Value

dggprentice gives the (log) probability function, pggprentice gives the (log) distribution function, qggprentice gives the quantile function, and rggprentice generates random deviates.

ggresiduals *Generic S3 method ggresiduals*

Description

Generic S3 method ggresiduals

Usage

```
ggresiduals(object, ...)
```

Arguments

object	a fitted model object.
...	further arguments passed to or from other methods.

Details

Generic method to plot residuals of survival models.

Value

the desired residual plot.

ggresiduals.survstan *ggresiduals method for survstan models*

Description

ggresiduals method for survstan models

Usage

```
## S3 method for class 'survstan'
ggresiduals(object, type = c("coxsnell", "martingale", "deviance"), ...)
```

Arguments

object	a fitted model object of the class survstan.
type	type of residuals used in the plot: coxsnell (default), martingale and deviance.
...	further arguments passed to or from other methods.

Details

This function produces residuals plots of Cox-Snell residuals, martingale residuals and deviance residuals.

Value

the desired residual plot.

Examples

```
library(survstan)
ovarian$rx <- as.factor(ovarian$rx)
fit <- aftreg(Surv(futime, fustat) ~ age + rx, data = ovarian, baseline = "weibull", init = 0)
ggresiduals(fit, type = "coxsnell")
ggresiduals(fit, type = "martingale")
ggresiduals(fit, type = "deviance")
```

---

ggstacy *The Generalized Gamma Distribution (Stacy's original parametrization)*

Description

Probability function, distribution function, quantile function and random generation for the distribution with parameters alpha, gamma and kappa.

Usage

```
dggstacy(x, alpha, gamma, kappa, log = FALSE)

pggstacy(q, alpha, gamma, kappa, log.p = FALSE, lower.tail = TRUE)

qggstacy(
  p,
  alpha = 1,
  gamma = 1,
  kappa = 1,
  log.p = FALSE,
  lower.tail = TRUE,
  ...
)

rggstacy(n, alpha = 1, gamma = 1, kappa = 1, ...)
```

Arguments

x	vector of (non-negative integer) quantiles.
alpha	shape parameter of the distribution (alpha > 0).
gamma	scale parameter of the distribution (gamma > 0).
kappa	shape parameter of the distribution (kappa > 0).

log, log.p	logical; if TRUE, probabilities p are given as log(p).
q	vector of quantiles.
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$; otherwise, $P[X > x]$.
p	vector of probabilities.
...	further arguments passed to other methods.
n	number of random values to return.

Details

Probability density function:

$$f(x|\alpha,\gamma,\kappa) = \frac{\kappa}{\gamma^{\alpha}\Gamma(\alpha/\kappa)} x^{\alpha-1} \exp\left\{-\left(\frac{x}{\gamma}\right)^{\kappa}\right\} I_{[0,\infty)}(x),$$

for $\alpha > 0$, $\gamma > 0$ and $\kappa > 0$.

Distribution function:

$$F(t|\alpha,\gamma,\kappa) = F_G(x|\nu,1),$$

where $x = \left(\frac{t}{\gamma}\right)^{\kappa}$, and $F_G(\cdot|\nu,1)$ corresponds to the distribution function of a gamma distribution with shape parameter $\nu = \alpha/\gamma$ and scale parameter equals to 1.

Value

dggstacy gives the (log) probability function, pggstacy gives the (log) distribution function, qggstacy gives the quantile function, and rggstacy generates random deviates.

Gompertz *The Gompertz Distribution*

Description

Probability function, distribution function, quantile function and random generation for the distribution with parameters alpha and gamma.

Usage

```
dgompertz(x, alpha = 1, gamma = 1, log = FALSE, ...)

pgompertz(q, alpha = 1, gamma = 1, lower.tail = TRUE, log.p = FALSE, ...)

qgompertz(p, alpha = 1, gamma = 1, lower.tail = FALSE, log.p = FALSE, ...)

rgompertz(n, alpha = 1, gamma = 1, ...)
```

Arguments

x	vector of (non-negative integer) quantiles.
alpha	shape parameter of the distribution (alpha > 0).
gamma	scale parameter of the distribution (gamma > 0).
log, log.p	logical; if TRUE, probabilities p are given as log(p).
...	further arguments passed to other methods.
q	vector of quantiles.
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$; otherwise, $P[X > x]$.
p	vector of probabilities.
n	number of random values to return.

Details

Probability density function:

$$f(x|\alpha,\gamma) = \alpha\gamma \exp\{\gamma x - \alpha(e^{\gamma x} - 1)\} I_{[0,\infty)}(x),$$

for $\alpha > 0$ and $\gamma > 0$.

Distribution function:

$$F(x|\alpha,\gamma) = 1 - \exp\{-\alpha(e^{\gamma x} - 1)\},$$

for $x > 0$, $\alpha > 0$ and $\gamma > 0$.

Value

dgompertz gives the (log) probability function, pgompertz gives the (log) distribution function, qgompertz gives the quantile function, and rgompertz generates random deviates.

---

ipass *IRESSA Pan-Asia Study (IPASS) data set*

Description

Reconstructed IPASS clinical trial data reported in Argyropoulos and Unruh (2015). Although reconstructed, this data set preserves all features exhibited in references with full access to the observations from this clinical trial. The data base is related to the period of March 2006 to April 2008. The main purpose of the study is to compare the drug gefitinib against carboplatin/paclitaxel doublet chemotherapy as first line treatment, in terms of progression free survival (in months), to be applied to selected non-small-cell lung cancer (NSCLC) patients.

Format

A data frame with 1217 rows and 3 variables:

- time: progression free survival (in months)
- status: failure indicator (1 - failure; 0 - otherwise)
- arm: (1 - gefitinib; 0 - carboplatin/paclitaxel doublet chemotherapy)

Author(s)

Fabio N. Demarqui <fndemarqui@est.ufmg.br>

References

Argyropoulos, C. and Unruh, M. L. (2015). Analysis of time to event outcomes in randomized controlled trials by generalized additive models. PLOS One 10, 1-33.

---

logLik.survstan *Extract Log-Likelihood from a Fitted Model*

Description

Extracts the log-likelihood function for a fitted parametric model.

Usage

```
## S3 method for class 'survstan'
logLik(object, ...)
```

Arguments

object	a fitted model of the class survstan
...	further arguments passed to or from other methods.

Value

the log-likelihood value when a single model is passed to the function; otherwise, a data.frame with the log-likelihood values and the number of parameters is returned.

Examples

```
library(survstan)
fit1 <- aftreg(Surv(futime, fustat) ~ 1, data = ovarian, baseline = "weibull", init = 0)
fit2 <- aftreg(Surv(futime, fustat) ~ rx, data = ovarian, baseline = "weibull", init = 0)
fit3 <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
logLik(fit1, fit2, fit3)
```

model.matrix.survstan *Model.matrix method for survstan models*

Description

Reconstruct the model matrix for a survstan model.

Usage

```
## S3 method for class 'survstan'
model.matrix(object, ...)
```

Arguments

object	an object of the class survstan.
...	further arguments passed to or from other methods.

Value

The model matrix (or matrices) for the fit.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
model.matrix(fit)
```

phreg *Fitting Proportional Hazards Models*

Description

Function to fit proportional hazards (PH) models.

Usage

```
phreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.
baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

phreg returns an object of class "phreg" containing the fitted model.

Examples

```
library(survstan)
fit <- phreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

---

poreg *Fitting Proportional Odds Models*

Description

Function to fit proportional odds (PO) models.

Usage

```
poreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.

baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

poreg returns an object of class "poreg" containing the fitted model.

Examples

```
library(survstan)
fit <- poreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

---

print.summary.survstan

Print the summary.survstan output

Description

Produces a printed summary of a fitted survstan model.

Usage

```
## S3 method for class 'summary.survstan'
print(x, ...)
```

Arguments

x	an object of the class summary.survstan.
...	further arguments passed to or from other methods.

Value

No return value, called for side effects.

rank_models *Rank a collection of survstan models*

Description

Rank a collection of survstan models

Usage

```
rank_models(formula, data, survreg, baseline, dist = NULL, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.
survreg	survival regression models to be fitted (AFT, AH, PH, PO, YP and EH).
baseline	baseline distributions to be fitted; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distributions (for compability with the survreg function); default is NULL.
...	further arguments passed to other methods.

Value

a tibble containing the fitted models ranked according to their AICs.

Examples

```
library(survstan)
library(dplyr)

veteran <- veteran %>%
  mutate(across(c(trt, prior, celltype), as.factor))
fits <- rank_models(
  formula = Surv(time, status) ~ celltype+karno,
  data = veteran,
  survreg = c("aftreg", "ahreg", "phreg", "poreg", "ypreg", "ehreg"),
 baseline = c("exponential", "weibull", "lognormal", "loglogistic", "fatigue", "gamma", "rayleigh")
)
```

residuals.survstan *residuals method for survstan models*

Description

residuals method for survstan models

Usage

```
## S3 method for class 'survstan'
residuals(object, type = c("coxsnell", "martingale", "deviance"), ...)
```

Arguments

object	a fitted model object of the class survstan.
type	type of residuals desired: coxsnell (default), martingale and deviance.
...	further arguments passed to or from other methods.

Details

This function extracts the residuals, martingale residuals and deviance residuals of a survstan object.

Value

a vector containing the desired residuals.

Examples

```
library(survstan)
ovarian$rx <- as.factor(ovarian$rx)
fit <- aftreg(Surv(futime, fustat) ~ age + rx, data = ovarian, baseline = "weibull", init = 0)
residuals(fit, type = "coxsnell")
residuals(fit, type = "martingale")
residuals(fit, type = "deviance")
```

se *Generic S3 method se*

Description

Generic S3 method se

Usage

```
se(object, ...)
```

Arguments

object	a fitted model object.
...	further arguments passed to or from other methods.

Value

the standard errors associated with a set of parameter estimators for a given model.

---

se.survstan *Estimated standard errors*

Description

Estimated standard errors

Usage

```
## S3 method for class 'survstan'
se(object, ...)
```

Arguments

object	an object of the class survstan.
...	further arguments passed to or from other methods.

Value

a vector with the standard errors.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
se(fit)
```

summary.survstan *Summary for a survstan object*

Description

Summary for a survstan object

Usage

```
## S3 method for class 'survstan'
summary(object, conf.level = 0.95, ...)
```

Arguments

object	the result of a call to summary.survstan
conf.level	the confidence level required.
...	further arguments passed to or from other methods.

Value

an object of the class summary.survstan containing a summary of the fitted model.

survfit.survstan *survfit method for survstan models*

Description

Computes the predicted survivor function for a phpe model.

Usage

```
## S3 method for class 'survstan'
survfit(formula, newdata = NULL, ...)
```

Arguments

formula	an object of the class survstan
newdata	a data frame containing the set of explanatory variables; if NULL, a data.frame with the observed failure times and their corresponding estimated baseline survivals is returned.
...	further arguments passed to or from other methods.

Value

a data.frame containing the estimated survival probabilities.

Examples

```
library(survstan)
library(ggplot2)
data(ipass)
ipass$arm <- as.factor(ipass$arm)
fit <- ypreg(Surv(time, status)~arm, data=ipass, baseline = "weibull")
summary(fit)
newdata <- data.frame(arm=as.factor(0:1))
surv <- survfit(fit, newdata)
ggplot(surv, aes(x=time, y=surv, color = arm)) +
  geom_line()
```

---

tidy.survstan *Tidy a survstan object*

Description

Tidy a survstan object

Usage

```
## S3 method for class 'survstan'
tidy(x, conf.int = FALSE, conf.level = 0.95, ...)
```

Arguments

x	a fitted model object.
conf.int	Logical indicating whether or not to include a confidence interval in the tidied output. Defaults to FALSE.
conf.level	the confidence level required.
...	further arguments passed to or from other methods.

Details

Convert a fitted model into a tibble.

Value

a tibble with a summary of the fit.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
tidy(fit)
```

vcov.survstan *Variance-covariance matrix*

Description

This function extracts and returns the variance-covariance matrix associated with the regression coefficients when the maximum likelihood estimation approach is used in the model fitting.

Usage

```
## S3 method for class 'survstan'
vcov(object, all = FALSE, ...)
```

Arguments

object	an object of the class survstan.
all	logical; if FALSE (default), only covariance matrix associated with regression coefficients is returned; if TRUE, the full covariance matrix is returned.
...	further arguments passed to or from other methods.

Value

the variance-covariance matrix associated with the parameters estimators.

Examples

```
library(survstan)
fit <- aftreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull", init = 0)
vcov(fit)
```

ypreg *Fitting Yang and Prentice Models*

Description

Function to fit Yang and Prentice (YP) models.

Usage

```
ypreg(formula, data, baseline = "weibull", dist = NULL, init = 0, ...)
```

Arguments

formula	an object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted.
data	data an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment from which function is called.
baseline	the chosen baseline distribution; options currently available are: exponential, weibull, lognormal, loglogistic and Birnbaum-Saunders (fatigue) distributions.
dist	alternative way to specify the baseline distribution (for compatibility with the survreg function); default is NULL.
init	initial values specification (default value is 0); see the detailed documentation for init in optimizing.
...	further arguments passed to other methods.

Value

ypreg returns an object of class "ypreg" containing the fitted model.

Examples

```
library(survstan)
fit <- ypreg(Surv(futime, fustat) ~ ecog.ps + rx, data = ovarian, baseline = "weibull")
summary(fit)
```

Index

∗ **Continuous**
  ggprentice, 13
  ggstacy, 16
  Gompertz, 17
∗ **Univariate**
  ggprentice, 13
  ggstacy, 16
  Gompertz, 17
∗ **datasets**
  gastric, 13
  ipass, 18
∗ **distribution**
  ggprentice, 13
  ggstacy, 16
  Gompertz, 17

aftreg, 3
ahreg, 4
AIC.survstan, 5
anova.survstan, 6

coef.survstan, 7
confint.survstan, 7
cross_time, 8
cross_time.survstan, 9

dggprentice *(*ggprentice*)*, 13
dggstacy *(*ggstacy*)*, 16
dgompertz *(*Gompertz*)*, 17

ehreg, 10
emmeans-survstan-helpers, 11
estimates, 11
extractAIC.survstan, 12

gastric, 13
ggprentice, 13
ggresiduals, 15
ggresiduals.survstan, 15
ggstacy, 16
Gompertz, 17
gompertz *(*Gompertz*)*, 17

Hgompertz *(*Gompertz*)*, 17
hgompertz *(*Gompertz*)*, 17

ipass, 18

logLik.survstan, 19

model.matrix.survstan, 20

optimizing, *4, 5*, *10*, *21, 22*, *29*

pggprentice *(*ggprentice*)*, 13
pggstacy *(*ggstacy*)*, 16
pgompertz *(*Gompertz*)*, 17
phreg, 20
poreg, 21
print.summary.survstan, 22

qggprentice *(*ggprentice*)*, 13
qggstacy *(*ggstacy*)*, 16
qgompertz *(*Gompertz*)*, 17

rank_models, 23
recover_data.ehreg
        *(*emmeans-survstan-helpers*)*, 11
recover_data.survstan
        *(*emmeans-survstan-helpers*)*, 11
recover_data.ypreg
        *(*emmeans-survstan-helpers*)*, 11
residuals.survstan, 24
rggprentice *(*ggprentice*)*, 13
rggstacy *(*ggstacy*)*, 16
rgompertz *(*Gompertz*)*, 17

se, 24
se.survstan, 25
summary.survstan, 26
survfit.survstan, 26
survreg, *4, 5*, *10*, *21–23*, *29*

survstan (survstan-package), 3
survstan-package, 3

tidy.survstan, 27

vcov.survstan, 28

ypreg, 28