## Package 'Rwtdttt'

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```
Type Package
Title Parametric Waiting Time Distribution estimation
Version 0.1.0
Author Sabrina Giometto, Malcolm Gillies, Henrik Støvring
Maintainer Henrik Støvring <hstov@health.sdu.dk>
Description Estimation of prescription durations and treatment
      probability based on the parametric Waiting Time Distribution.
      Pharmacoepidemiologic databases contains information on medication
      dispensings at pharmacies. Studies using such data typically require
      some estimate of duration of treatment after a dispensing (known as
      the prescription duration), which can be estimated using the
      parametric Waiting Time Distribution.
License What license is it under?
Roxygen list(markdown = TRUE)
Imports bbmle,
      class,
      data.table,
      dplyr,
      haven,
      methods,
      graphics,
      stats
Suggests testthat (>= 3.0.0)
Config/testthat/edition 3
Encoding UTF-8
LazyData true
RoxygenNote 7.2.3
Collate 'Rwtdttt-package.R'
      'dfunctions.R'
      'wtd-class.R'
      'plot.R'
      'pred_dur_prob.R'
      'ranwtdttt.R'
      'wtdttt.R'
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dexp

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The Exponential distribution

## Description

The Exponential distribution

## Usage

```
dexp(x, logitp, lnbeta, delta = 1, log = FALSE)
```

## **Arguments**

X	vector of prescription redemption times
logitp	log-odds of being a prevalent user
lnbeta	log of beta (scale)
dol+a	VVVV

delta XXXX

logical; if TRUE, probabilities p are given as log(p)

dlnorm

The Lognormal distribution

## Description

The Lognormal distribution

## Usage

```
dlnorm(x, logitp, mu, lnsigma, delta = 1, log = FALSE)
```

## Arguments

X	vector of prescription redemption times
logitp	log-odds of being a prevalent user
mu	mean on log-scale
lnsigma	log of standard deviation on log-scale

delta XXXX

logical; if TRUE, probabilities p are given as log(p).

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dweib	The Weibull distribution	

## Description

The Weibull distribution

## Usage

```
dweib(x, logitp, lnalpha, lnbeta, delta = 1, log = FALSE)
```

## **Arguments**

log

X	vector of prescription redemption times
logitp	log-odds of being a prevalent user
lnalpha	log of alpha (shape)
lnbeta	log of beta (scale)
delta	XXXX

plot, wtd, ANY-method Make WTD diagnostic plots

## Description

Make diagnostic plots showing the fit of an estimated parametric Waiting Time Distribution (WTD) with respect to the observed histogram of prescription redemptions. HS 240108: I think this needs to incorporate information on time period of observations (start and end specified in call to wtdttt)

logical; if TRUE, probabilities p are given as log(p)

## Usage

```
## S4 method for signature 'wtd,ANY'
plot(object, x, y, ...)
```

## Arguments

wtd object, typically result of wtdttt

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predict,wtd-method

Make WTD predictions

## **Description**

Make predictions based on an estimated parametric Waiting Time Distribution (WTD) model, either the probability of a person still being in treatment or the duration of observed prescription redemptions.

## Usage

```
## S4 method for signature 'wtd'
predict(
  object,
  newdata = NULL,
  type = "dur",
  distrx = NULL,
  quantile = 0.8,
  se.fit = FALSE,
  na.action = na.pass,
  ...
)
```

## **Arguments**

wtd

a fitted object of class inheriting from "wtd"

## Value

A vector of predictions

ranwtdttt

Fit Waiting Time Distribution with random index times

## Description

ranwtdttt() estimates maximum likelihood estimates for parametric Waiting Time Distribution (WTD) based on observed prescription redemptions with adjustment for covariates using one or more random index times for each individual. Reports estimates of prevalence fraction and specified percentile of inter-arrival density together with regression coefficients.

## Usage

```
ranwtdttt(
  form,
  parameters = NULL,
  data,
  id,
  start,
  end,
```

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```
reverse = F,
nsamp = 1,
subset,
na.action = na.pass,
init,
control = NULL,
...
)
```

#### **Arguments**

form an object of class "formula" (or one that can be coered to that class): a symbolic

description of the model to be fitted. The details of the model specification are

given under 'Details'

parameters model formulae for distribution parameters

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 wtdttt is called.

id the name of the variable that identifies distinct individuals

start start of observation window end end of observation window

reverse logical; Fit the reverse waiting time distribution.

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

process.

na.action a function which indicates what should happen when the data contain NAs. The

default is set by the na.action setting of options, and is na.fail if that is unset. The 'factory-fresh' default is na.omit. Another possible value is NULL, no action.

Value na.exclude can be useful.

init starting values for the parameters.

control a list of parameters for controlling the fitting process.

... further arguments passed to other methods.

## Value

wtdttt returns an object of class "wtd" inheriting from "mle".

wtdttt Fit Waiting Time Distribution

#### **Description**

Estimates the maximum likelihood estimate for a parametric Waiting Time Distribution (WTD) based on observed prescription redemptions with adjustment for covariates. Reports estimates of prevalence fraction and specified percentile of inter-arrival density together with regression coefficients.

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#### **Usage**

```
wtdttt(
  data,
  form,
  parameters = NULL,
  start = NA,
  end = NA,
  reverse = F,
  id = NA,
  subset = NA,
  na.action = na.pass,
  init = NULL,
  control = NULL,
  ...
)
```

## **Arguments**

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 wtdttt is called.

form an object of class "formula" (or one that can be coered to that class): a symbolic

description of the model to be fitted. The details of the model specification are

given under 'Details'

parameters optional model formulae for distribution parameters start start of observation window (date or real number) end end of observation window (date or real number)

reverse logical; Fit the reverse waiting time distribution (default F).

id name of the id variable (optional)

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

process.

na.action a function which indicates what should happen when the data contain NAs. The

default is set by the na.action setting of options, and is na.fail if that is unset. The 'factory-fresh' default is na.omit. Another possible value is NULL, no action.

Value na.exclude can be useful.

init starting values for the parameters.

control a list of parameters for controlling the fitting process.

.. further arguments passed to other methods.

## Value

wtdttt returns an object of class "wtd" inheriting from "mle".

#### Model formula

The model formula form follows the pattern obstime ~ dist(alpha, beta, gamma) with

• obstime: the redemption time variable (date or real number)

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• dist: the parametric distribution for the forward or backward recurrence density (FRD/BRD), which must be dexp(), dweib() or dlnorm() i.e named after their corresponding interarrival density (IAD).

## Data format

The WTD is fit to the first prescription redemption of each individual within an observation window (ordinary WTD), or the last (reverse WTD), respectively.

You may prepare the data to this format, or optionally specify the name of an id variable to select the first or last redemption automatically.

If the redemption time data are of type date, a continuity correction will be applied automatically.

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