

Package ‘hctrial’

October 13, 2022

Type Package

Title Using Historical Controls for Designing Phase II Clinical Trials

Version 0.1.0

Description Provides functions for designing phase II clinical trials adjusting for the heterogeneity of the population using known subgroups or historical controls.

Depends R (>= 3.5.0)

License GPL-2 | GPL-3

LazyData TRUE

RoxygenNote 6.1.1

Imports clinfun, GenBinomApps, stats

Encoding UTF-8

NeedsCompilation no

Author Dominic Edelmann [aut, cre]

Maintainer Dominic Edelmann <dominic.edelmann@dkfz-heidelberg.de>

Repository CRAN

Date/Publication 2018-12-21 15:20:06 UTC

R topics documented:

hctrial	2
hist_end	2
hist_interim	3
hist_start	4
strat_end	5
strat_interim	5
strat_start	6
Index	8

hctrial	<i>hctrial: A package for designing phase 2 clinical trials adjusting for heterogeneous populations.</i>
---------	--

Description

The hctrial package provides functions for designing phase 2 clinical trials that adjust for the heterogeneity in the population.

Details

Two different ways are considered for designing a trial: based on known subgroups or based on historical data.

For initializing a stratified trial, use strat_start.

At interim, strat_interim should be used to adjust the trial.

At the end of the study, strat_end is used to adjust the trial again.

hist_start, hist_interim and hist_end work analogously, but are based on historical controls.

hist_end	<i>Adjust a design based on historical controls at the end of the study using the covariate data of the patients accrued in stage 2.</i>
----------	--

Description

Adjust a design based on historical controls at the end of the study using the covariate data of the patients accrued in stage 2.

Usage

```
hist_end(interim, stagetwo_data)
```

Arguments

interim An design based on historical controls and adjusted at interim as returned by hist_interim().

stagetwo_data A dataframe containing the relevant covariate data of the patients accrued in stage 2.

Value

A list returning the arguments of the function and the final design of the trial.

Examples

```

X <- abs(rnorm(1000, 0, 1))
Y <- rbinom(1000, 1, 1-exp(-X))
mydata <- data.frame("X" = X, "Y" = Y)
start <- hist_start(mydata, Y~X, c1 = 2)
n1 <- start$des_start[2]
X1 <- abs(rnorm(n1, 0, 1))
dataone <- data.frame("X" = X1)
interim <- hist_interim(start, dataone)
n2 <- interim$des_interim[4]
X2 <- abs(rnorm(n2, 0, 1))
datatwo <- data.frame("X" = X2)
hist_end(interim, datatwo)

```

hist_interim	<i>Adjust a design based on historical controls at interim using the covariate data of the patients accrued in stage 1.</i>
--------------	---

Description

Adjust a design based on historical controls at interim using the covariate data of the patients accrued in stage 1.

Usage

```
hist_interim(start, stageone_data)
```

Arguments

start	An initialized design based on historical controls as returned by hist_start().
stageone_data	A dataframe containing the relevant covariate data of the patients accrued in stage 1.

Value

A list returning the arguments of the function and the preliminary design of a trial based on historical controls adjusted at interim.

Examples

```

X <- abs(rnorm(1000, 0, 1))
Y <- rbinom(1000, 1, 1-exp(-X))
mydata <- data.frame("X" = X, "Y" = Y)
start <- hist_start(mydata, Y~X, c1 = 2)
n1 <- start$des_start[2]
X1 <- abs(rnorm(n1, 0, 1))
dataone <- data.frame("X" = X1)
hist_interim(start, dataone)

```

hist_start	<i>Initializes a design based on historical controls before the start of the study.</i>
------------	---

Description

Initializes a design based on historical controls before the start of the study.

Usage

```
hist_start(hist_data, formula, phi = "odds_ratio", c1, modelfit = NULL,
           mean0 = NULL, mean1 = NULL, alpha = 0.05, beta = 0.2)
```

Arguments

hist_data	A data frame containing covariates and binary responses for historical controls.
formula	A formula which is used for fitting a logistic regression model on the historical data.
phi	The relation between the response rate under the null and the response rate under the interesting alternative. "odds_ratio" assumes that the odds ratio (OR) between these response rates is constant with $OR = c1+1$. "difference" assumes that the response rate under the alternative is $c1$ higher than under the null. Can also be specified by the user by providing a function with arguments c and x .
c1	parameter for obtaining the response rate under the alternative, see description of phi.
modelfit	Can be used instead of formula and hist_data to provide an arbitrary fitted model that is compatible with <code>predict(modelfit, type="response")</code> . formula and hist_data are ignored if modelfit is specified.
mean0	Optional: Can be used to overwrite the estimated average response rate under the null of the fitted model.
mean1	Optional: Can be used to overwrite the estimated average response rate under the alternative of the fitted model.
alpha	Specified type I error of the trial.
beta	Specified type II error of the trial.

Value

A list returning the arguments of the function and the preliminary design for starting the stratified trial.

Examples

```
X <- abs(rnorm(1000, 0, 1))
Y <- rbinom(1000, 1, 1-exp(-X))
mydata <- data.frame("X" = X, "Y" = Y)
hist_start(mydata, Y~X, c1 = 2)
```

strat_end	<i>Adjust a subspace stratified design at the end of the study.</i>
-----------	---

Description

Adjust a subspace stratified design at the end of the study.

Usage

```
strat_end(interim, sub_stagetwo)
```

Arguments

interim A preliminary stratified design adjusted at interim as returned by strat_interim().
sub_stagetwo The subtypes observed for the patients accrued in stage 2.

Value

A list returning the arguments of the function and the final design of the stratified trial.

Examples

```
p0_sub <- c(0.1, 0.3, 0.5)
p1_sub <- c(0.3, 0.5, 0.7)
distr_sub <- c(1/3, 1/3, 1/3)
start <- strat_start(p0_sub, p1_sub, distr_sub)
n1 <- start$des_start[2]
subone <- sample(c(1,2,3), n1, TRUE)
interim <- strat_interim(start, subone)
n2 <- interim$des_interim[4]
subtwo <- sample(c(1,2,3), n2, TRUE)
strat_end(interim, subtwo)
```

strat_interim	<i>Adjust a subspace stratified design at interim.</i>
---------------	--

Description

Adjust a subspace stratified design at interim.

Usage

```
strat_interim(start, sub_stageone)
```

Arguments

start An initialized stratified design as returned by strat_start().
 sub_stageone The subtypes observed for the patients accrued in stage 1.

Value

A list returning the arguments of the function and the preliminary design of a stratified trial adjusted at interim.

Examples

```
p0_sub <- c(0.1, 0.3, 0.5)
p1_sub <- c(0.3, 0.5, 0.7)
distr_sub <- c(1/3, 1/3, 1/3)
start <- strat_start(p0_sub, p1_sub, distr_sub)
n1 <- start$des_start[2]
subone <- sample(c(1,2,3), n1, TRUE)
strat_interim(start, subone)
```

 strat_start

Initializes a subspace stratified design before the start of the study.

Description

Initializes a subspace stratified design before the start of the study.

Usage

```
strat_start(p0_sub, p1_sub, distr_sub, alpha = 0.05, beta = 0.2)
```

Arguments

p0_sub A vector, where the i -th entry corresponds to the response rate under the null for the i -th subtype.
 p1_sub A vector, where the i -th entry corresponds to the response rate under the alternative for the i -th subtype.
 distr_sub A vector, where the i -th entry corresponds to the prevalence of the i -th subtype in the population.
 alpha Specified type I error of the trial.
 beta Specified type II error of the trial.

Value

A list returning the arguments of the function and the preliminary design for starting the stratified trial.

Examples

```
p0_sub <- c(0.1, 0.3, 0.5)
p1_sub <- c(0.3, 0.5, 0.7)
distr_sub <- c(1/3, 1/3, 1/3)
strat_start(p0_sub, p1_sub, distr_sub)
```

Index

hctrtrial, [2](#)
hctrtrial-package (hctrtrial), [2](#)
hist_end, [2](#)
hist_interim, [3](#)
hist_start, [4](#)

strat_end, [5](#)
strat_interim, [5](#)
strat_start, [6](#)