

Package ‘lstat’

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Type Package

Title Power and Sample Size Calculation for Non-Proportional Hazards

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Description Performs power and sample size calculation for non-proportional hazards model using the Fleming-Harrington family of weighted log-rank tests. The sequentially calculated log-rank test score statistics are assumed to have independent increments as characterized in Anastasios A. Tsiatis (1982) <doi:10.1080/01621459.1982.10477898>. The mean and variance of log-rank test score statistics are calculated based on Kaifeng Lu (2021) <doi:10.1002/pst.2069>. The boundary crossing probabilities are calculated using the recursive integration algorithm described in Christopher Jennison and Bruce W. Turnbull (2000, ISBN:0849303168).

License GPL (>= 2)

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lrstat-package	<i>Power and Sample Size Calculation for Non-Proportional Hazards</i>
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Description

Performs power and sample size calculation for non-proportional hazards model using the Fleming-Harrington family of weighted log-rank tests.

Details

For proportional hazards, the power is determined by the total number of events and the constant hazard ratio along with information rates and spending functions. For non-proportional hazards, the hazard ratio varies over time and the calendar time plays a key role in determining the mean and variance of the log-rank test score statistic. It requires an iterative algorithm to find the calendar time at which the targeted number of events will be reached for each interim analysis. The lrstat package uses the analytic method in Lu (2021) to find the mean and variance of the weighted log-rank test score statistic at each interim analysis. In addition, the package approximates the variance and covariance matrix of the sequentially calculated log-rank test statistics under the alternative hypothesis with that under the null hypothesis to take advantage of the independent increments structure in Tsiatis (1982) applicable for the Fleming-Harrington family of weighted log-rank tests.

The most useful functions in the package are lrstat, lrpower, lrsamplesize, and lrsim, which calculate the mean and variance of log-rank test score statistic at a sequence of given calendar times, the power of the log-rank test, the sample size in terms of accrual duration and follow-up duration, and the log-rank test simulation, respectively. The accrual function calculates the number of patients accrued at given calendar times. The caltime function finds the calendar times to reach the targeted number of events. The exitprob function calculates the stagewise exit probabilities for specified boundaries with a varying mean parameter over time based on an adaptation of the recursive integration algorithm described in Chapter 19 of Jennison and Turnbull (2000).

The development of the lrstat package is heavily influenced by the rpact package. We find their function arguments to be self-explanatory. We have used the same names whenever appropriate so that users familiar with the rpact package can learn the lrstat package quickly. However, there are notable differences:

- lrstat uses direct approximation, while rpact uses the Schoenfeld method for log-rank test power and sample size calculation.
- lrstat uses accrualDuration to explicitly set the end of accrual period, while rpact incorporates the end of accrual period in accrualTime.
- lrstat considers the trial a failure at the last stage if the log-rank test cannot reject the null hypothesis up to this stage and cannot stop for futility at an earlier stage.
- the lrsm function uses the variance of the log-rank test score statistic as the information.

Author(s)

Kaifeng Lu, <kaifenglu@gmail.com>

References

Anastasios A. Tsiatis. Repeated significance testing for a general class of statistics used in censored survival analysis. *J Am Stat Assoc.* 1982;77:855-861.

Christopher Jennison, Bruce W. Turnbull. *Group Sequential Methods with Applications to Clinical Trials*. Chapman & Hall/CRC: Boca Raton, 2000, ISBN:0849303168

Kaifeng Lu. Sample size calculation for logrank test and prediction of number of events over time. *Pharm Stat.* 2021;20:229-244.

See Also

rpact, gsDesign

Examples

```
lrpower(kMax = 2, informationRates = c(0.8, 1),
       criticalValues = c(2.250, 2.025), accrualIntensity = 20,
       piecewiseSurvivalTime = c(0, 6),
       lambda1 = c(0.0533, 0.0309), lambda2 = c(0.0533, 0.0533),
       gamma1 = 0.00427, gamma2 = 0.00427,
       accrualDuration = 22, followupTime = 18)
```

accrual

Number of enrolled subjects

Description

Obtains the number of subjects enrolled by given calendar times.

Usage

```
accrual(
  time = NA_real_,
  accrualTime = 0L,
  accrualIntensity = NA_real_,
  accrualDuration = NA_real_
)
```

Arguments

time A vector of calendar times at which to calculate the number of enrolled subjects.

accrualTime Accrual time intervals, must start with 0, e.g., `c(0, 3)` breaks the time axis into 2 accrual intervals: `[0, 3)` and `[3, Inf)`.

accrualIntensity A vector of accrual intensities, one for each accrual time interval.

accrualDuration Duration of the enrollment period.

Value

A vector of total number of subjects enrolled by the specified calendar times.

Examples

```
# Example 1: Uniform enrollment with 20 patients per month for 12 months.
```

```
accrual(time = 3, accrualTime = 0, accrualIntensity = 20,
        accrualDuration = 12)
```

```
# Example 2: Piecewise accrual, 10 patients per month for the first
# 3 months, and 20 patients per month thereafter. Patient recruitment
# ends at 12 months for the study.
```

```
accrual(time = c(2, 9), accrualTime = c(0, 3),
        accrualIntensity = c(10, 20), accrualDuration = 12)
```

caltime

Calendar times for target number of events

Description

Obtains the calendar times to reach the target number of subjects having an event.

Usage

```

caltime(
  nevents = NA_real_,
  allocationRatioPlanned = 1,
  accrualTime = 0L,
  accrualIntensity = NA_real_,
  piecewiseSurvivalTime = 0L,
  stratumFraction = 1L,
  lambda1 = NA_real_,
  lambda2 = NA_real_,
  gamma1 = 0L,
  gamma2 = 0L,
  accrualDuration = NA_real_,
  followupTime = NA_real_,
  fixedFollowup = 0L,
  numSubintervals = 300L
)

```

Arguments

<code>nevents</code>	A vector of target number of events.
<code>allocationRatioPlanned</code>	Allocation ratio for the active treatment versus control. Defaults to 1 for equal randomization.
<code>accrualTime</code>	Accrual time intervals, must start with 0, e.g., $c(0, 3)$ breaks the time axis into 2 accrual intervals: $[0, 3)$ and $[3, \text{Inf})$.
<code>accrualIntensity</code>	A vector of accrual intensities, one for each accrual time interval.
<code>piecewiseSurvivalTime</code>	A vector that specifies the time intervals for the piecewise exponential survival distribution, must start with 0, e.g., $c(0, 6)$ breaks the time axis into 2 event intervals: $[0, 6)$ and $[6, \text{Inf})$. Defaults to 0 for exponential distribution.
<code>stratumFraction</code>	A vector of stratum fractions that sum to 1. Defaults to 1 for no stratification.
<code>lambda1</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the active treatment group.
<code>lambda2</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the control group.
<code>gamma1</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the active treatment group.
<code>gamma2</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the control group.

accrualDuration Duration of the enrollment period.
followupTime Follow-up time for the last enrolled subject.
fixedFollowup Whether a fixed follow-up design is used. Defaults to 0 for variable follow-up.
numSubintervals Number of sub-intervals to approximate the mean and variance of the weighted log-rank test score statistic. Defaults to 300. Specify a larger number for better approximation.

Value

A vector of calendar times expected to yield the target number of events.

Examples

```
# Piecewise accrual, piecewise exponential survivals, and 5% dropout by
# the end of 1 year.
```

```
caltime(nevents = c(24, 80), allocationRatioPlanned = 1,
        accrualTime = seq(0, 9),
        accrualIntensity = c(26/9*seq(1, 9), 26),
        piecewiseSurvivalTime = c(0, 6),
        stratumFraction = c(0.2, 0.8),
        lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
        lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
        gamma1 = -log(1-0.05)/12,
        gamma2 = -log(1-0.05)/12,
        accrualDuration = 22,
        followupTime = 18, fixedFollowup = FALSE)
```

exitprob

Stagewise exit probabilities

Description

Obtains the stagewise exit probabilities for both efficacy and futility stopping.

Usage

```
exitprob(b, a, theta, I)
```

Arguments

b Upper boundaries on the z-test statistic scale.
a Lower boundaries on the z-test statistic scale. Defaults to $c(\text{rep}(-6.0, \text{kMax}-1), \text{b}[\text{kMax}])$ if left unspecified, where $\text{kMax} = \text{length}(\text{b})$.

theta	Stagewise parameter of interest, e.g., $-U/V$ for weighted log-rank test, where U is the mean and V is the variance of the weighted log-rank test score statistic at each stage. For proportional hazards and conventional log-rank test, use the scalar input, $\text{theta} = -\log(\text{HR})$.
I	Stagewise cumulative information, e.g., V , the variance of the weighted log-rank test score statistic at each stage. For conventional log-rank test, information can be approximated by $\text{phi} \cdot (1 - \text{phi}) \cdot D$, where phi is the probability of being allocated to the active arm, and D is the total number of events at each stage.

Value

A list of stagewise exit probabilities: one vector for efficacy stopping probabilities, and the other vector for futility stopping probabilities.

Examples

```
exitprob(b = c(3.471, 2.454, 2.004), a = NA,
        theta = -log(0.6), I = c(50, 100, 150)/4)
```

getDesign

Get group sequential design

Description

Obtains the drift parameter and stopping boundaries for a generic group sequential design assuming a constant treatment effect.

Usage

```
getDesign(
  beta = 0.2,
  kMax = NA_integer_,
  informationRates = NA_real_,
  efficacyStopping = NA_integer_,
  futilityStopping = NA_integer_,
  criticalValues = NA_real_,
  alpha = 0.025,
  typeAlphaSpending = "sfOF",
  parameterAlphaSpending = NA_real_,
  userAlphaSpending = NA_real_,
  futilityBounds = NA_real_,
  typeBetaSpending = "none",
  parameterBetaSpending = NA_real_,
  userBetaSpending = NA_real_
)
```

Arguments

beta	Type II error. Defaults to 0.2.
kMax	The maximum number of stages.
informationRates	The information rates in terms of number of events. Fixed prior to the trial. Defaults to $(1:kMax) / kMax$ if left unspecified.
efficacyStopping	Indicators of whether efficacy stopping is allowed at each stage. Defaults to true if left unspecified.
futilityStopping	Indicators of whether futility stopping is allowed at each stage. Defaults to true if left unspecified.
criticalValues	Upper boundaries on the z-test statistic scale for stopping for efficacy.
alpha	The significance level. Defaults to 0.025.
typeAlphaSpending	The type of alpha spending. One of the following: "OF" for O'Brien-Fleming boundaries, "P" for Pocock boundaries, "WT" for Wang & Tsiatis boundaries, "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, "user" for user defined spending, and "none" for no early efficacy stopping. Defaults to "sfOF".
parameterAlphaSpending	The parameter value for the alpha spending. Corresponds to Delta for "WT", rho for "sfKD", and gamma for "sfHSD".
userAlphaSpending	The user defined alpha spending. Cumulative alpha spent up to each stage.
futilityBounds	Lower boundaries on the z-test statistic scale for stopping for futility at stages 1, ..., kMax-1. Defaults to rep(-6, kMax-1) if left unspecified.
typeBetaSpending	The type of beta spending. One of the following: "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, "user" for user defined spending, and "none" for no early futility stopping. Defaults to "none".
parameterBetaSpending	The parameter value for the beta spending. Corresponds to rho for "sfKD", and gamma for "sfHSD".
userBetaSpending	The user defined beta spending. Cumulative beta spent up to each stage.

Value

A list of S3 class design.

Examples

```

getDesign(beta = 0.2,
          kMax = 2,
          informationRates = c(0.5,1),
          alpha = 0.025,
          typeAlphaSpending = "sfOF",
          typeBetaSpending = "sfP")

```

Irpower

*Log-rank test power***Description**

Estimates the power, stopping probabilities, and expected sample size in a two-group survival design.

Usage

```

Irpower(
  kMax = NA_integer_,
  informationRates = NA_real_,
  efficacyStopping = NA_integer_,
  futilityStopping = NA_integer_,
  criticalValues = NA_real_,
  alpha = 0.025,
  typeAlphaSpending = "sfOF",
  parameterAlphaSpending = NA_real_,
  userAlphaSpending = NA_real_,
  futilityBounds = NA_real_,
  typeBetaSpending = "none",
  parameterBetaSpending = NA_real_,
  hazardRatioH0 = 1,
  allocationRatioPlanned = 1,
  accrualTime = 0L,
  accrualIntensity = 20L,
  piecewiseSurvivalTime = 0L,
  stratumFraction = 1L,
  lambda1 = 0.0309,
  lambda2 = 0.0533,
  gamma1 = 0L,
  gamma2 = 0L,
  accrualDuration = 11.6,
  followupTime = 18,
  fixedFollowup = 0L,
  rho1 = 0,

```

```

rho2 = 0,
numSubintervals = 300L,
estimateHazardRatio = 1L,
typeOfComputation = "direct"
)

```

Arguments

kMax The maximum number of stages.

informationRates The information rates in terms of number of events. Fixed prior to the trial. Defaults to $(1:kMax) / kMax$ if left unspecified.

efficacyStopping Indicators of whether efficacy stopping is allowed at each stage. Defaults to true if left unspecified.

futilityStopping Indicators of whether futility stopping is allowed at each stage. Defaults to true if left unspecified.

criticalValues Upper boundaries on the z-test statistic scale for stopping for efficacy.

alpha The significance level. Defaults to 0.025.

typeAlphaSpending The type of alpha spending. One of the following: "OF" for O'Brien-Fleming boundaries, "P" for Pocock boundaries, "WT" for Wang & Tsatis boundaries, "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, "user" for user defined spending, and "none" for no early efficacy stopping. Defaults to "sfOF".

parameterAlphaSpending The parameter value for the alpha spending. Corresponds to Delta for "WT", rho for "sfKD", and gamma for "sfHSD".

userAlphaSpending The user defined alpha spending. Cumulative alpha spent up to each stage.

futilityBounds Lower boundaries on the z-test statistic scale for stopping for futility at stages 1, ..., kMax-1. Defaults to $\text{rep}(-6, kMax-1)$ if left unspecified.

typeBetaSpending The type of beta spending. One of the following: "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, and "none" for no early futility stopping. Defaults to "none".

parameterBetaSpending The parameter value for the beta spending. Corresponds to rho for "sfKD", and gamma for "sfHSD".

hazardRatioH0 Hazard ratio under the null hypothesis for the active treatment versus control. Defaults to 1 for superiority test.

allocationRatioPlanned Allocation ratio for the active treatment versus control. Defaults to 1 for equal randomization.

accrualTime	Accrual time intervals, must start with 0, e.g., $c(\theta, 3)$ breaks the time axis into 2 accrual intervals: $[0, 3)$ and $[3, \text{Inf})$.
accrualIntensity	A vector of accrual intensities, one for each accrual time interval.
piecewiseSurvivalTime	A vector that specifies the time intervals for the piecewise exponential survival distribution, must start with 0, e.g., $c(\theta, 6)$ breaks the time axis into 2 event intervals: $[0, 6)$ and $[6, \text{Inf})$. Defaults to 0 for exponential distribution.
stratumFraction	A vector of stratum fractions that sum to 1. Defaults to 1 for no stratification.
lambda1	A vector of hazard rates for the event in each analysis time interval by stratum for the active treatment group.
lambda2	A vector of hazard rates for the event in each analysis time interval by stratum for the control group.
gamma1	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the active treatment group.
gamma2	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the control group.
accrualDuration	Duration of the enrollment period.
followupTime	Follow-up time for the last enrolled subject.
fixedFollowup	Whether a fixed follow-up design is used. Defaults to 0 for variable follow-up.
rho1	First parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
rho2	Second parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
numSubintervals	Number of sub-intervals to approximate the mean and variance of the weighted log-rank test score statistic. Defaults to 300. Specify a larger number for better approximation.
estimateHazardRatio	Whether to estimate the hazard ratio from weighted Cox regression model and report the stopping boundaries on the hazard ratio scale.
typeOfComputation	Whether to use the direct approximation method or the Schoenfeld method. Defaults to "direct". Can use "Schoenfeld" under proportional hazards and conventional log-rank test.

Value

A list of S3 class `Irpower` with 3 components:

* overallResults containing the overall rejection probability, overall significance level, maximum and expected number of events, maximum and expected number of dropouts, total and expected number of subjects, maximum and expected study duration, along with input parameters including accrual duration, followup duration, whether a fixed follow-up is used, parameters for the FH weights, allocation ratio, number of stages, and hazard ratio under H0.

* byStageResults containing information rates, efficacy and futility boundaries on the Z-scale, probability for efficacy and futility stopping at the stage, cumulative probability for efficacy and futility stopping by the stage, cumulative alpha spent, expected number of events, number of dropouts, number of subjects, and expected study time, efficacy and futility boundaries on the HR scale and on the p-value scale, information for weighted log-rank test, hazard ratio from weighted Cox regression, and where efficacy and futility stopping are allowed by stage.

* settings containing input parameters such as alpha and beta spending function and parameter values, accrual time, accrual intensity, piecewise survival time, stratum fraction, and hazard rates for survival and dropout by group.

Examples

```
# Piecewise accrual, piecewise exponential survivals, and 5% dropout by
# the end of 1 year.
```

```
lrpower(kMax = 2, informationRates = c(0.8, 1),
        alpha = 0.025, typeAlphaSpending = "sfOF",
        allocationRatioPlanned = 1, accrualTime = seq(0, 9),
        accrualIntensity = c(26/9*seq(1, 9), 26),
        piecewiseSurvivalTime = c(0, 6),
        stratumFraction = c(0.2, 0.8),
        lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
        lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
        gamma1 = -log(1-0.05)/12,
        gamma2 = -log(1-0.05)/12, accrualDuration = 22,
        followupTime = 18, fixedFollowup = FALSE)
```

Irsamplesize

Log-rank test sample size

Description

Obtains the needed accrual duration given power and follow-up time, the needed follow-up time given power and accrual duration, or the needed absolute accrual rates given power, accrual duration, follow-up duration, and relative accrual rates in a two-group survival design.

Usage

```
lrsamplesize(
  beta = 0.2,
  kMax = NA_integer_,
  informationRates = NA_real_,
```

```

efficacyStopping = NA_integer_,
futilityStopping = NA_integer_,
criticalValues = NA_real_,
alpha = 0.025,
typeAlphaSpending = "sfOF",
parameterAlphaSpending = NA_real_,
userAlphaSpending = NA_real_,
futilityBounds = NA_real_,
typeBetaSpending = "none",
parameterBetaSpending = NA_real_,
userBetaSpending = NA_real_,
hazardRatioH0 = 1,
allocationRatioPlanned = 1,
accrualTime = 0L,
accrualIntensity = 20L,
piecewiseSurvivalTime = 0L,
stratumFraction = 1L,
lambda1 = 0.0309,
lambda2 = 0.0533,
gamma1 = 0L,
gamma2 = 0L,
accrualDuration = NA_real_,
followupTime = 18,
fixedFollowup = 0L,
rho1 = 0,
rho2 = 0,
numSubintervals = 300L,
estimateHazardRatio = 1L,
typeOfComputation = "direct",
interval = as.numeric(c(0.001, 240))
)

```

Arguments

beta	Type II error. Defaults to 0.2.
kMax	The maximum number of stages.
informationRates	The information rates in terms of number of events. Fixed prior to the trial. Defaults to (1:kMax) / kMax if left unspecified.
efficacyStopping	Indicators of whether efficacy stopping is allowed at each stage. Defaults to true if left unspecified.
futilityStopping	Indicators of whether futility stopping is allowed at each stage. Defaults to true if left unspecified.
criticalValues	Upper boundaries on the z-test statistic scale for stopping for efficacy.
alpha	The significance level. Defaults to 0.025.

<code>typeAlphaSpending</code>	The type of alpha spending. One of the following: "OF" for O'Brien-Fleming boundaries, "P" for Pocock boundaries, "WT" for Wang & Tsatis boundaries, "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, "user" for user defined spending, and "none" for no early efficacy stopping. Defaults to "sfOF".
<code>parameterAlphaSpending</code>	The parameter value for the alpha spending. Corresponds to Delta for "WT", rho for "sfKD", and gamma for "sfHSD".
<code>userAlphaSpending</code>	The user defined alpha spending. Cumulative alpha spent up to each stage.
<code>futilityBounds</code>	Lower boundaries on the z-test statistic scale for stopping for futility at stages 1, ..., $k_{\text{Max}}-1$. Defaults to $\text{rep}(-6, k_{\text{Max}}-1)$ if left unspecified.
<code>typeBetaSpending</code>	The type of beta spending. One of the following: "sfOF" for O'Brien-Fleming type spending function, "sfP" for Pocock type spending function, "sfKD" for Kim & DeMets spending function, "sfHSD" for Hwang, Shi & DeCani spending function, "user" for user defined spending, and "none" for no early futility stopping. Defaults to "none".
<code>parameterBetaSpending</code>	The parameter value for the beta spending. Corresponds to rho for "sfKD", and gamma for "sfHSD".
<code>userBetaSpending</code>	The user defined beta spending. Cumulative beta spent up to each stage.
<code>hazardRatioH0</code>	Hazard ratio under the null hypothesis for the active treatment versus control. Defaults to 1 for superiority test.
<code>allocationRatioPlanned</code>	Allocation ratio for the active treatment versus control. Defaults to 1 for equal randomization.
<code>accrualTime</code>	Accrual time intervals, must start with 0, e.g., $c(0, 3)$ breaks the time axis into 2 accrual intervals: $[0, 3)$ and $[3, \text{Inf})$.
<code>accrualIntensity</code>	A vector of accrual intensities, one for each accrual time interval.
<code>piecewiseSurvivalTime</code>	A vector that specifies the time intervals for the piecewise exponential survival distribution, must start with 0, e.g., $c(0, 6)$ breaks the time axis into 2 event intervals: $[0, 6)$ and $[6, \text{Inf})$. Defaults to 0 for exponential distribution.
<code>stratumFraction</code>	A vector of stratum fractions that sum to 1. Defaults to 1 for no stratification.
<code>lambda1</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the active treatment group.
<code>lambda2</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the control group.

gamma1	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the active treatment group.
gamma2	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the control group.
accrualDuration	Duration of the enrollment period.
followupTime	Follow-up time for the last enrolled subject.
fixedFollowup	Whether a fixed follow-up design is used. Defaults to 0 for variable follow-up.
rho1	First parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
rho2	Second parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
numSubintervals	Number of sub-intervals to approximate the mean and variance of the weighted log-rank test score statistic. Defaults to 300. Specify a larger number for better approximation.
estimateHazardRatio	Whether to estimate the hazard ratio from weighted Cox regression model and report the stopping boundaries on the hazard ratio scale.
typeOfComputation	Whether to use the direct approximation method or the Schoenfeld method. Defaults to "direct". Can use "Schoenfeld" under proportional hazards and conventional log-rank test.
interval	The interval to search for the solution of accrualDuration or followupDuration. Defaults to $c(0.001, 240)$. Adjustment may be needed for non-monotone relationship with study power.

Value

A list of S3 class `lrpower`.

Examples

```
# Piecewise accrual, piecewise exponential survivals, and 5% dropout by
# the end of 1 year.

# Example 1: Obtains accrual duration given power and follow-up duration

Irsamplesize(beta = 0.2, kMax = 2,
              informationRates = c(0.8, 1),
              alpha = 0.025, typeAlphaSpending = "sfOF",
              accrualTime = seq(0, 9),
              accrualIntensity = c(26/9*seq(1, 9), 26),
              piecewiseSurvivalTime = c(0, 6),
```

```

stratumFraction = c(0.2, 0.8),
lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
gamma1 = -log(1-0.05)/12,
gamma2 = -log(1-0.05)/12,
accrualDuration = NA,
followupTime = 18, fixedFollowup = FALSE)

```

Example 2: Obtains follow-up duration given power and accrual duration

```

Irsamplesize(beta = 0.2, kMax = 2,
  informationRates = c(0.8, 1),
  alpha = 0.025, typeAlphaSpending = "sfOF",
  accrualTime = seq(0, 9),
  accrualIntensity = c(26/9*seq(1, 9), 26),
  piecewiseSurvivalTime = c(0, 6),
  stratumFraction = c(0.2, 0.8),
  lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
  lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
  gamma1 = -log(1-0.05)/12,
  gamma2 = -log(1-0.05)/12,
  accrualDuration = 22,
  followupTime = NA, fixedFollowup = FALSE)

```

Example 3: Obtains absolute accrual intensity given power,
accrual duration, follow-up duration, and relative accrual intensity

```

Irsamplesize(beta = 0.2, kMax = 2,
  informationRates = c(0.8, 1),
  alpha = 0.025, typeAlphaSpending = "sfOF",
  accrualTime = seq(0, 9),
  accrualIntensity = c(26/9*seq(1, 9), 26),
  piecewiseSurvivalTime = c(0, 6),
  stratumFraction = c(0.2, 0.8),
  lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
  lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
  gamma1 = -log(1-0.05)/12,
  gamma2 = -log(1-0.05)/12,
  accrualDuration = 22,
  followupTime = 18, fixedFollowup = FALSE)

```

Example 4: Non-inferiority trial with fixed follow-up and
superiority alternative

```

lr <- Irsamplesize(beta = 0.1,
  kMax = 3,
  alpha = 0.025,
  typeAlphaSpending = "sfOF",
  hazardRatioH0 = 1.1,
  accrualTime = c(0, 6),

```



```

accrualIntensity = c(1000, 1500),
lambda1 = log(2)/48*0.95,
lambda2 = log(2)/48,
gamma1 = -log(1-0.08)/12,
gamma2 = -log(1-0.08)/12,
accrualDuration = NA,
followupTime = 18,
fixedFollowup = 1)

```

```
lr
```

```
lrsim
```

```
Log-rank test simulation
```

Description

Performs simulation for two-arm group sequential trials based on weighted log-rank test.

Usage

```

lrsim(
  kMax = NA_integer_,
  informationTime = NA_real_,
  criticalValues = NA_real_,
  futilityBounds = NA_real_,
  hazardRatioH0 = 1,
  allocation1 = 1L,
  allocation2 = 1L,
  accrualTime = 0L,
  accrualIntensity = NA_real_,
  piecewiseSurvivalTime = 0L,
  stratumFraction = 1L,
  lambda1 = NA_real_,
  lambda2 = NA_real_,
  gamma1 = 0L,
  gamma2 = 0L,
  accrualDuration = NA_real_,
  followupTime = NA_real_,
  fixedFollowup = 0L,
  rho1 = 0,
  rho2 = 0,
  plannedEvents = NA_integer_,
  plannedTime = NA_real_,
  maxNumberOfIterations = 1000L,
  maxNumberOfRawDatasetsPerStage = 0L,
  seed = NA_integer_
)

```

Arguments

<code>kMax</code>	The maximum number of stages.
<code>informationTime</code>	Information time in terms of variance of weighted log-rank test score statistic under the null hypothesis. Same as <code>informationRates</code> in terms of number of events for the conventional log-rank test. Use <code>calTime</code> and <code>lrstat</code> to derive the information time for weighted log-rank tests. Fixed prior to the trial. Defaults to <code>plannedEvents / sum(plannedEvents)</code> if left unspecified.
<code>criticalValues</code>	Upper boundaries on the z-test statistic scale for stopping for efficacy.
<code>futilityBounds</code>	Lower boundaries on the z-test statistic scale for stopping for futility at stages 1, ..., <code>kMax-1</code> . Defaults to <code>rep(-6, kMax-1)</code> if left unspecified.
<code>hazardRatioH0</code>	Hazard ratio under the null hypothesis for the active treatment versus control. Defaults to 1 for superiority test.
<code>allocation1</code>	Number of subjects in the active treatment group in a randomization block. Defaults to 1 for equal randomization.
<code>allocation2</code>	Number of subjects in the control group in a randomization block. Defaults to 1 for equal randomization.
<code>accrualTime</code>	Accrual time intervals, must start with 0, e.g., <code>c(0, 3)</code> breaks the time axis into 2 accrual intervals: <code>[0, 3)</code> and <code>[3, Inf)</code> .
<code>accrualIntensity</code>	A vector of accrual intensities, one for each accrual time interval.
<code>piecewiseSurvivalTime</code>	A vector that specifies the time intervals for the piecewise exponential survival distribution, must start with 0, e.g., <code>c(0, 6)</code> breaks the time axis into 2 event intervals: <code>[0, 6)</code> and <code>[6, Inf)</code> . Defaults to 0 for exponential distribution.
<code>stratumFraction</code>	A vector of stratum fractions that sum to 1. Defaults to 1 for no stratification.
<code>lambda1</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the active treatment group.
<code>lambda2</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the control group.
<code>gamma1</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the active treatment group.
<code>gamma2</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the control group.
<code>accrualDuration</code>	Duration of the enrollment period.
<code>followupTime</code>	Follow-up time for the last enrolled subject.
<code>fixedFollowup</code>	Whether a fixed follow-up design is used. Defaults to 0 for variable follow-up.
<code>rho1</code>	First parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.

rho2	Second parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
plannedEvents	The planned cumulative total number of events at each stage.
plannedTime	The planned analysis time for each stage needed for analyses planned at calendar times, in which case, plannedEvents should be missing.
maxNumberOfIterations	The number of simulation iterations. Defaults to 1000.
maxNumberOfRawDatasetsPerStage	The number of raw datasets per stage to extract. Defaults to 1.
seed	The seed to reproduce the simulation results. The computer clock will be used if left unspecified,

Value

A list of S3 class `lrsim` with 3 components:

* `overview` is a list of containing incremental and cumulative efficacy and futility stopping probabilities by stage, expected number of events, number of dropouts, number of subjects, and analysis time by stage, overall rejection probability, overall expected number of events, number of dropouts, number of subjects, and study duration, the hazard ratio under H_0 , and whether the analyses are planned based on the number of events or calendar time.

* `sumdata` is a data frame of summary data by stage for each iteration, containing at which stage the trial stops, whether the target number of events is achieved, the analysis time, number of accrued subjects overall and by treatment group, number of events overall and by treatment group, number of dropouts overall and by treatment group, numerator and variance of weighted log-rank score statistic, log-rank test Z-statistic, and whether the trial stops for efficacy or futility at the stage.

* `rawdata` (exists if `maxNumberOfRawDatasetsPerStage` is a positive integer) is a data frame for subject-level data for selected replications, containing the subject number, arrival time, stratum, treatment group, survival time, dropout time, observation time when the trial stops, time under observation, and event and dropout indicators.

Examples

```
# Example 1: analyses based on number of events

sim1 = lrsim(kMax = 2, informationTime = c(0.5, 1),
            criticalValues = c(2.797, 1.977),
            accrualIntensity = 11,
            lambda1 = 0.018, lambda2 = 0.030,
            accrualDuration = 12,
            plannedEvents = c(60, 120),
            maxNumberOfIterations = 1000,
            maxNumberOfRawDatasetsPerStage = 1,
            seed = 314159)

# summary statistics
sim1

# summary for each simulated data set
```

```

head(sim1$sumdata)

# raw data for selected replication
head(sim1$rawdata)

# Example 2: analyses based on calendar time have similar power

sim2 = lrstat(kMax = 2, informationTime = c(0.5, 1),
              criticalValues = c(2.797, 1.977),
              accrualIntensity = 11,
              lambda1 = 0.018, lambda2 = 0.030,
              accrualDuration = 12,
              plannedTime = c(31.9, 113.2),
              maxNumberOfIterations = 1000,
              maxNumberOfRawDatasetsPerStage = 1,
              seed = 314159)

# summary statistics
sim2

# summary for each simulated data set
head(sim2$sumdata)

```

Irstat

Number of subjects having an event and log-rank statistics

Description

Obtains the number of subjects accrued, number of events and number of dropouts in each group, mean and variance of weighted log-rank score statistic, estimated hazard ratio from weighted Cox regression and variance of log hazard ratio estimate at given calendar times.

Usage

```

lrstat(
  time = NA_real_,
  hazardRatioH0 = 1,
  allocationRatioPlanned = 1,
  accrualTime = 0L,
  accrualIntensity = NA_real_,
  piecewiseSurvivalTime = 0L,
  stratumFraction = 1L,
  lambda1 = NA_real_,
  lambda2 = NA_real_,
  gamma1 = 0L,
  gamma2 = 0L,
  accrualDuration = NA_real_,

```

```

followupTime = NA_real_,
fixedFollowup = 0L,
rho1 = 0,
rho2 = 0,
numSubintervals = 300L,
predictEventOnly = 0L
)

```

Arguments

<code>time</code>	A vector of calendar times at which to calculate the number of events and the mean and variance of log-rank test score statistic.
<code>hazardRatioH0</code>	Hazard ratio under the null hypothesis for the active treatment versus control. Defaults to 1 for superiority test.
<code>allocationRatioPlanned</code>	Allocation ratio for the active treatment versus control. Defaults to 1 for equal randomization.
<code>accrualTime</code>	Accrual time intervals, must start with 0, e.g., $c(0, 3)$ breaks the time axis into 2 accrual intervals: $[0, 3)$ and $[3, \text{Inf})$.
<code>accrualIntensity</code>	A vector of accrual intensities, one for each accrual time interval.
<code>piecewiseSurvivalTime</code>	A vector that specifies the time intervals for the piecewise exponential survival distribution, must start with 0, e.g., $c(0, 6)$ breaks the time axis into 2 event intervals: $[0, 6)$ and $[6, \text{Inf})$. Defaults to 0 for exponential distribution.
<code>stratumFraction</code>	A vector of stratum fractions that sum to 1. Defaults to 1 for no stratification.
<code>lambda1</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the active treatment group.
<code>lambda2</code>	A vector of hazard rates for the event in each analysis time interval by stratum for the control group.
<code>gamma1</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the active treatment group.
<code>gamma2</code>	The hazard rate for exponential dropout, a vector of hazard rates for piecewise exponential dropout applicable for all strata, or a vector of hazard rates for dropout in each analysis time interval by stratum for the control group.
<code>accrualDuration</code>	Duration of the enrollment period.
<code>followupTime</code>	Follow-up time for the last enrolled subject.
<code>fixedFollowup</code>	Whether a fixed follow-up design is used. Defaults to 0 for variable follow-up.
<code>rho1</code>	First parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.

<code>rho2</code>	Second parameter of the Fleming-Harrington family of weighted log-rank test. Defaults to 0 for conventional log-rank test.
<code>numSubintervals</code>	Number of sub-intervals to approximate the mean and variance of the weighted log-rank test score statistic. Defaults to 300. Specify a larger number for better approximation.
<code>predictEventOnly</code>	Whether to predict the number of events only. Defaults to 0 for obtaining log-rank score statistic mean and variance. Set <code>predictEventOnly = 1</code> for predicting the number of events only. Set <code>predictEventOnly = 2</code> for predicting the number of events, calculating the mean and variance of log-rank score statistic, and calculating the estimated hazard ratio and variance of log hazard ratio.

Value

A data frame of the number of subjects enrolled, the number of subjects having an event overall and in each group, the number of subjects who drop out overall and in each group, the mean and variance of weighted log-rank score statistic, the estimated hazard ratio from weighted Cox regression, and variance of the log hazard ratio estimate at the specified calendar times.

Examples

```
# Piecewise accrual, piecewise exponential survivals, and 5% dropout by
# the end of 1 year.
```

```
lrstat(time = c(22, 40), allocationRatioPlanned = 1,
        accrualTime = seq(0, 9),
        accrualIntensity = c(26/9*seq(1, 9), 26),
        piecewiseSurvivalTime = c(0, 6),
        stratumFraction = c(0.2, 0.8),
        lambda1 = c(0.0533, 0.0309, 1.5*0.0533, 1.5*0.0309),
        lambda2 = c(0.0533, 0.0533, 1.5*0.0533, 1.5*0.0533),
        gamma1 = -log(1-0.05)/12,
        gamma2 = -log(1-0.05)/12,
        accrualDuration = 22,
        followupTime = 18, fixedFollowup = FALSE)
```

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