

Package ‘bbssr’

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Type Package

Title Blinded Sample Size Re-Estimation for Binary Endpoints

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Description Provides comprehensive tools for blinded sample size re-estimation (BSSR) in two-arm clinical trials with binary endpoints. Unlike traditional fixed-sample designs, BSSR allows adaptive sample size adjustments during trials while maintaining statistical integrity and study blinding. Implements five exact statistical tests: Pearson chi-squared, Fisher exact, Fisher mid-p, Z-pooled exact unconditional, and Boschloo exact unconditional tests. Supports restricted, unrestricted, and weighted BSSR approaches with exact Type I error control. Statistical methods based on Mehrotra et al. (2003) <[doi:10.1111/1541-0420.00051](https://doi.org/10.1111/1541-0420.00051)> and Kieser (2020) <[doi:10.1007/978-3-030-49528-2_21](https://doi.org/10.1007/978-3-030-49528-2_21)>.

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URL <https://github.com/gosukehommaEX/bbssr>

BugReports <https://github.com/gosukehommaEX/bbssr/issues>

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BinaryPower	<i>Power Calculation for Two-Arm Trials with Binary Endpoints</i>
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Description

Calculates power for two-arm trials with binary endpoints using exact statistical tests. The function supports five different one-sided tests and can handle vectors of probabilities.

Usage

```
BinaryPower(p1, p2, N1, N2, alpha, Test)
```

Arguments

p1	True probability of responders for group 1 (can be a vector with different values)
p2	True probability of responders for group 2 (can be a vector with different values)
N1	Sample size for group 1
N2	Sample size for group 2
alpha	One-sided level of significance
Test	Type of statistical test. Options: 'Chisq', 'Fisher', 'Fisher-midP', 'Z-pool', or 'Boschloo'

Details

The function supports the following five one-sided tests:

- The one-sided Pearson chi-squared test (Chisq)
- The Fisher exact test (Fisher)
- The Fisher mid-p test (Fisher-midP)
- The Z-pooled exact unconditional test (Z-pool)
- The Boschloo exact unconditional test (Boschloo)

The power calculation is based on the exact distribution of the test statistic under the specified alternative hypothesis.

Value

A numeric value or vector of power values. If vectors are provided for p1 and p2, a vector of powers corresponding to each combination will be returned.

Author(s)

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Examples

```
# Simple power calculation with fast Chi-squared test
power1 <- BinaryPower(p1 = 0.5, p2 = 0.2, N1 = 5, N2 = 5,
                      alpha = 0.025, Test = 'Chisq')
print(power1)

# More computationally intensive examples
# Single power calculation with larger sample size
power2 <- BinaryPower(p1 = 0.5, p2 = 0.2, N1 = 10, N2 = 40,
                      alpha = 0.025, Test = 'Boschloo')
print(power2)

# Multiple power calculations
p1_vec <- c(0.5, 0.6, 0.7, 0.8)
p2_vec <- c(0.2, 0.2, 0.2, 0.2)
powers <- BinaryPower(p1 = p1_vec, p2 = p2_vec, N1 = 10, N2 = 40,
                      alpha = 0.025, Test = 'Fisher')
print(powers)
```

BinaryPowerBSSR

*Power Calculation for Two-Arm Trials with Binary Endpoints Using
Blinded Sample Size Re-estimation (BSSR)*

Description

Calculates the power for two-arm trials with binary endpoints when blinded sample size re-estimation (BSSR) is implemented. The function supports five different statistical tests and allows for both restricted and unrestricted designs with optional weighted approaches.

Usage

```
BinaryPowerBSSR(
  asmd.p1,
  asmd.p2,
  p,
  Delta.A,
  Delta.T,
  N1,
  N2,
  omega,
  r,
```

```

    alpha,
    tar.power,
    Test,
    restricted,
    weighted
  )

```

Arguments

<code>asmd.p1</code>	Assumed proportion of responders for group 1
<code>asmd.p2</code>	Assumed proportion of responders for group 2
<code>p</code>	Vector of pooled proportions of responders from both groups (can specify multiple values)
<code>Delta.A</code>	Assumed treatment effect (risk difference)
<code>Delta.T</code>	True treatment effect (risk difference)
<code>N1</code>	Initial sample size of group 1
<code>N2</code>	Initial sample size of group 2
<code>omega</code>	Fraction of sample size used for interim analysis (i.e., for BSSR)
<code>r</code>	Allocation ratio to group 1
<code>alpha</code>	One-sided level of significance
<code>tar.power</code>	Target power
<code>Test</code>	Type of statistical test. Options: 'Chisq', 'Fisher', 'Fisher-midP', 'Z-pool', or 'Boschloo'
<code>restricted</code>	Logical. If TRUE, restricted design is chosen
<code>weighted</code>	Logical. If TRUE, weighted approach is chosen

Details

The function supports the following five one-sided tests:

- The one-sided Pearson chi-squared test (Chisq)
- The Fisher exact test (Fisher)
- The Fisher mid-p test (Fisher-midP)
- The Z-pooled exact unconditional test (Z-pool)
- The Boschloo exact unconditional test (Boschloo)

Value

A data frame containing:

p1 True probability of responders for group 1
p2 True probability of responders for group 2
p True probability of pooled responders from both groups
power.BSSR Power for BSSR design
power.TRAD Power for traditional design

Author(s)

Gosuke Homma (<my.name.is.gosuke@gmail.com>)

Examples

```
# Simple BSSR calculation with fast Chi-squared test
result1 <- BinaryPowerBSSR(
  asmd.p1 = 0.6, asmd.p2 = 0.3,
  p = 0.45,
  Delta.A = 0.3, Delta.T = 0.3,
  N1 = 5, N2 = 5, omega = 0.5, r = 1,
  alpha = 0.025, tar.power = 0.8,
  Test = 'Chisq',
  restricted = FALSE, weighted = FALSE
)
print(result1)

# More computationally intensive BSSR examples
result2 <- BinaryPowerBSSR(
  asmd.p1 = 0.45,
  asmd.p2 = 0.09,
  p = seq(0.14, 0.23, by = 0.01),
  Delta.A = 0.36,
  Delta.T = 0.36,
  N1 = 24,
  N2 = 24,
  omega = 0.5,
  r = 1,
  alpha = 0.025,
  tar.power = 0.8,
  Test = 'Z-pool',
  restricted = FALSE,
  weighted = TRUE
)
print(result2)
```

BinaryRR

Rejection Region for Two-Arm Trials with Binary Endpoints

Description

Provides a rejection region (RR) for two-arm trials with binary endpoints using various exact statistical tests. The function supports five different one-sided tests.

Usage

```
BinaryRR(N1, N2, alpha, Test)
```

Arguments

N1	Sample size for group 1
N2	Sample size for group 2
alpha	One-sided level of significance
Test	Type of statistical test. Options: 'Chisq', 'Fisher', 'Fisher-midP', 'Z-pool', or 'Boschloo'

Details

The function supports the following five one-sided tests:

- The one-sided Pearson chi-squared test (Chisq)
- The Fisher exact test (Fisher)
- The Fisher mid-p test (Fisher-midP)
- The Z-pooled exact unconditional test (Z-pool)
- The Boschloo exact unconditional test (Boschloo)

Value

A logical matrix representing the rejection region (RR). Matrix dimensions are (N1+1) x (N2+1), where TRUE indicates rejection of the null hypothesis.

Author(s)

Gosuke Homma (<my.name.is.gosuke@gmail.com>)

Examples

```
# Simple example with small sample sizes (runs quickly)
N1 <- 5
N2 <- 5
alpha <- 0.025
Test <- 'Chisq'
RR <- BinaryRR(N1, N2, alpha, Test)
print(dim(RR)) # Should be (6, 6)

# More computationally intensive example
N1 <- 20
N2 <- 10
alpha <- 0.025
Test <- 'Boschloo'
RR <- BinaryRR(N1, N2, alpha, Test)
print(RR)
```

Description

Calculates the required sample size for two-arm trials with binary endpoints using various exact statistical tests. The function supports five different one-sided tests.

Usage

```
BinarySampleSize(p1, p2, r, alpha, tar.power, Test)
```

Arguments

p1	True probability of responders for group 1
p2	True probability of responders for group 2
r	Allocation ratio to group 1 (i.e., allocation ratio of group 1:group 2 = r:1, $r > 0$)
alpha	One-sided level of significance
tar.power	Target power
Test	Type of statistical test. Options: 'Chisq', 'Fisher', 'Fisher-midP', 'Z-pool', or 'Boschloo'

Details

The function supports the following five one-sided tests:

- The one-sided Pearson chi-squared test (Chisq)
- The Fisher exact test (Fisher)
- The Fisher mid-p test (Fisher-midP)
- The Z-pooled exact unconditional test (Z-pool)
- The Boschloo exact unconditional test (Boschloo)

The calculation uses a three-step approach:

1. Calculate initial sample size using normal approximation for chi-squared test
2. Perform power calculation with the initial sample size
3. Use grid search algorithm to find the optimal sample size

Value

A data frame containing:

p1 True probability of responders for group 1

p2 True probability of responders for group 2

r Allocation ratio to group 1

alpha One-sided level of significance

tar.power Target power

Test Name of the statistical test

Power Calculated power

N1 Required sample size of group 1

N2 Required sample size of group 2

N Total required sample size

Author(s)

Gosuke Homma (<my.name.is.gosuke@gmail.com>)

Examples

```
# Simple sample size calculation with fast Chi-squared test
result1 <- BinarySampleSize(p1 = 0.4, p2 = 0.2, r = 1, alpha = 0.025,
                           tar.power = 0.8, Test = 'Chisq')
print(result1)

# More computationally intensive examples
# Sample size for Fisher exact test
result2 <- BinarySampleSize(p1 = 0.5, p2 = 0.2, r = 2, alpha = 0.025,
                           tar.power = 0.9, Test = 'Fisher')
print(result2)

# Sample size for Boschloo test
result3 <- BinarySampleSize(p1 = 0.6, p2 = 0.3, r = 1, alpha = 0.025,
                           tar.power = 0.8, Test = 'Boschloo')
print(result3)
```


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