Package 'MIDN'

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Title Nearly Exact Sample Size Calculation for Exact Powerful Nonrandomized Tests for Differences Between Binomial Proportions

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Description Implementation of the mid-n algorithms presented in Wellek S (2015) <DOI:10.1111/stan.12063> Statistica Neerlandica 69, 358-373 for exact sample size calculation for superiority trials with binary outcome.

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MIDN-package

Nearly exact sample size calculation for exact powerful nonrandomized tests for differences between binomial proportions

Description

Implementation of the mid-n algorithms presented in Wellek S (2015) Statistica Neerlandica 69, 358-373 for exact sample size calculation for superiority trials with binary outcome.

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References

Wellek S: Nearly exact sample size calculation for powerful nonrandomized tests for differences between binomial proportions. Statistica Neerlandica 69 (2015), 358-373.

Examples

```
result1 <- fisher_boschloo_midN(0.025,0.0001,0.95,0.8,0.8,2,1)
POWEX <- result1[5]
result1  # shows values of vector result1
POWEX    # shows value of POWEX

result2 <- McNem_Score_midn(0.025,0.0001,0.585,0.315,0.9)
POWEX <- result2[3]
result2  # shows values of vector result2
POWEX    # shows value of POWEX
```

fisher_boschloo_midN

Nearly exact sample size calculation for the Fisher-Boschloo test for differences between independent binomial proportions

Description

The function computes the exact sample sizes required in the randomized UMPU test and its conservative nonrandomized version for attaining prespecified power. In a final step, the mean of both quantities is output as an nearly exact value required in the Fisher-Boschloo test, a powerful nonrandomized version of the exact Fisher-type test.

Usage

```
fisher_boschloo_midN(alpha, SW, p1, p2, POWO, mton_a, mton_b)
```

Arguments

alpha	target significance level
SW	step width for increasing p2 in the search for the size of a given critical region in the sample space of (X,Y)
p1	true value of the responder rate for Population 1
p2	true value of the responder rate for Population 2
POWO	power to be obtained against the alternative (p1,p2)
mton_a	desired ratio of sample sizes: numerator
mton_b	desired ratio of sample sizes: denominator

Value

mstart	initial value of 1st sample size
nstart	initial value of 2nd sample size
Mex	size of Sample 1 for randomized UMPU test
Nex	size of Sample 2 for randomized UMPU test
POWEX	power of randomized UMPU test attained with m=Mex,n=Nex
Mnr	size of Sample 1 for conservative nonrandomized Fisher-type test
Nnr	size of Sample 2 for conservative nonrandomized Fisher-type test
POWNR	power of conservative nonrandomized Fisher-type test attained with m=Mnr,n=Nnr
midN_m	nearly exact size of Sample 1 for Boschloo-Fisher test
midN_n	nearly exact size of Sample 1 for Boschloo-Fisher test

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References

Wellek S: Nearly exact sample size calculation for powerful nonrandomized tests for differences between binomial proportions. Statistica Neerlandica 69 (2015), 358-373.

Examples

```
result1 <- fisher_boschloo_midN(0.025,0.0001,0.95,0.8,0.8,2,1)
POWEX <- result1[5]
result1  # shows values of vector result1
POWEX    # shows value of POWEX
```

`McNem_Score_midn` *Nearly exact sample size calculation for the level-corrected score test for differences between binomial proportions estimated from paired data*

Description

Again, the function computes the exact sample sizes required in the randomized UMPU test and its conservative nonrandomized counterepart for attaining prespecified power. However, in contrast to the parallel group setting, the midpoint of the interval between these two numbers shall now used as an nearly exact value of the number of pairs to be observed in the asymptotic test based on the score-statistic corrected for possible exceedances of the nominal significance level.

Usage

```
McNem_Score_midn(alpha, SW, ppl, pmi, POWO)
```

Arguments

`alpha`	target significance level, 1-sided
`SW`	width of search grid for determining the size of a given critical region in the sample space of N+ [= number of pairs with (Xi,Yi) = (1,0)] and N0 [= number of tied pairs]
`ppl`	true value of Pr[(X,Y) = (1,0)]
`pmi`	true value of Pr[(X,Y) = (0,1)]
`POWO`	power to be attained in the level-corrected score test against the alternative (ppl,pmi)

Value

`nstart`	initial value for the iteration algorithm
`Nex`	sample size required in the exact randomized McNemar test
`POWEX`	power of the exact randomized McNemar test performed with Nex pairs
`Nnr`	sample size required in the conservative nonrandomized McNemar test
`POWNR`	power of the nonrandomized McNemar test performed with Nnr pairs
`mid_n`	midpoint of the interval [Nex,Nnr], rounded to the next integer

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References

Wellek S: Nearly exact sample size calculation for powerful nonrandomized tests for differences between binomial proportions. Statistica Neerlandica 69 (2015), 358-373.

Examples

```
result2 <- McNem_Score_midn(0.025,0.0001,0.585,0.315,0.9)
POWEX <- result2[3]
result2  # shows values of vector result2
POWEX    # shows value of POWEX
```

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