

# Package ‘ADCT’

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**Type** Package

**Title** Adaptive Design in Clinical Trials

**Version** 0.1.0

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**Description** Existing adaptive design methods in clinical trials. The package includes power, stopping boundaries (sample size) calculation functions for two-group group sequential designs, adaptive design with coprimary endpoints, biomarker-informed adaptive design, etc.

**Imports** stats, mvtnorm

**Suggests** clinfun, gsDesign

**License** GPL (>= 2)

**LazyData** TRUE

**NeedsCompilation** no

**Repository** CRAN

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BioInfo.Power      *Power calculation for Biomarker-Informed Design with Hierarchical Model*

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### Description

Given the Biomarker-Informed design information, returns the overall power and probability of the arm is selected as the winner.

### Usage

BioInfo.Power(uCtl, u0y, u0x, rhou, suy, sux, rho, sy, sx, Zalpha, N1, N, nArms, nSims)

### Arguments

uCtl	mean value for the control group.
u0y	mean parameter of the group 1 for the parent model.
u0x	mean parameter of the group 2 for the parent model.
rhou	correlation coefficient between two groups for the parent model.
suy	standard deviation of the group 1 for the parent model.
sux	standard deviation of the group 2 for the parent model.
rho	correlation coefficient between two groups for the lower level model.
sy	standard deviation of the group 1 for the lower level model.
sx	standard deviation of the group 2 for the lower level model.
Zalpha	critical point for rejection.
N1	sample size per group at interim analysis.
N	sample size per group at final analysis.
nArms	number of active groups.
nSims	number of simulation times.

### Value

The evaluated power and probability of selecting the arm as the winner.

### Author(s)

Yalin Zhu

### References

Chang, M. (2014). Adaptive design theory and implementation using SAS and R. *CRC Press*.

**Examples**

```
## Determine critical value Zalpha for alpha (power) =0.025
u0y=c(0,0,0); u0x=c(0,0,0)
BioInfo.Power(uCtl=0, u0y, u0x, rhou=1, suy=0, sux=0, rho=1, sy=4, sx=4,
  Zalpha=2.772, N1=100, N=300, nArms=3, nSims=1000)
## Power simulation
u0y=c(1,0.5,0.2)
u0x=c(2,1,0.5)
BioInfo.Power(uCtl=0, u0y, u0x, rhou=0.2, suy=0.2, sux=0.2, rho=0.2, sy=4, sx=4,
  Zalpha=2.772, N1=100, N=300, nArms=3, nSims=500)
```

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 CopriEndpt.Power

*Power Calculation for Two Coprimary Endpoints.*


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**Description**

Given the group sequential design information, returns the overall power.

**Usage**

```
CopriEndpt.Power(n, tau, mu1, mu2, rho, alpha1, alpha2, alternative)
```

**Arguments**

n	sample size for the design.
tau	information time for the interim analysis.
mu1	mean value for coprimary endpoint 1.
mu2	mean value for coprimary endpoint 2.
rho	correlation coefficient between two coprimary endpoints.
alpha1	significant level for the first stage.
alpha2	significant level for the second stage.
alternative	indicates the alternative hypothesis and must be one of "two.sided" or "two.sided".

**Value**

The evaluated power with attributes and computational error.

**Author(s)**

Yalin Zhu

**References**

Chang, M. (2014). Adaptive design theory and implementation using SAS and R. *CRC Press*.

**Examples**

```
# Example in Chang (2014) page 272
CopriEndpt.Power(n=197, tau=0.5, mu1=0.2, mu2=0.2, rho=0.5,
alpha1=0.0025, alpha2=0.024, alternative="one.sided")
sapply(c(-0.8,-0.5,-0.2,0,0.2,0.5,0.8),CopriEndpt.Power,
n=197, tau=0.5, mu1=0.2, mu2=0.2, alpha1=0.0025, alpha2=0.024, alternative="one.sided")
```

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OneArm.CondPower	<i>Conditional power for one-arm, two-stage design with two primary endpoints</i>
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**Description**

Given the group sequential design information, returns the conditional power.

**Usage**

```
OneArm.CondPower(mu1, mu2, n1, n2, rho, tau, alpha2, alternative)
```

**Arguments**

mu1	mean value for the first stage (endpoint 1).
mu2	mean value for the second stage (endpoint 2).
n1	sample size for the first stage.
n2	sample size for the second stage.
rho	correlation coefficient between two coprimary endpoints.
tau	information time for the interim analysis.
alpha2	significant level for the second stage.
alternative	indicates the alternative hypothesis and must be one of "two.sided" or "two.sided".

**Value**

The evaluated power with attributes and computational error.

**Author(s)**

Yalin Zhu

**References**

Chang, M. (2014). Adaptive design theory and implementation using SAS and R. *CRC Press*.

**Examples**

```
# Example in Chang (2014) page 277
OneArm.CondPower(mu1=0.1333, mu2=0.1605, n1=130, n2=130, rho=0.35,
  tau=0.5, alpha2=0.024, alternative = "one.sided")
OneArm.CondPower(mu1=0.1333, mu2=0.1605, n1=130, n2=414, rho=0.35,
  tau=0.5, alpha2=0.024, alternative = "one.sided")
```

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TwoArms.CondPower	<i>Conditional power for two-group design, two-stage design with two primary endpoints</i>
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**Description**

Given the group sequential design information, returns the conditional power.

**Usage**

```
TwoArms.CondPower(mu1, mu2, sigma1, sigma2, n1, n2, rho, tau, alpha2, alternative)
```

**Arguments**

mu1	mean value for the first stage (endpoint 1).
mu2	mean value for the second stage (endpoint 2).
sigma1	standard deviation for the first stage.
sigma2	standard deviation for the second stage.
n1	sample size for the first stage.
n2	sample size for the second stage.
rho	correlation coefficient between two coprimary endpoints.
tau	information time for the interim analysis.
alpha2	significant level for the second stage.
alternative	indicates the alternative hypothesis and must be one of "two.sided" or "two.sided".

**Value**

The evaluated power with attributes and computational error.

**Author(s)**

Yalin Zhu

**References**

Chang, M. (2014). Adaptive design theory and implementation using SAS and R. *CRC Press*.

**Examples**

```
# Example in Chang (2014) page 278
TwoArms.CondPower(mu1=0.28, sigma1=1.9, mu2=0.35, sigma2=2.2, n1=340, n2=340,
rho=0.3, tau=0.5, alpha2=0.024, alternative = "one.sided")
TwoArms.CondPower(mu1=0.28, sigma1=1.9, mu2=0.35, sigma2=2.2, n1=340, n2=482,
rho=0.3, tau=0.5, alpha2=0.024, alternative = "one.sided")
TwoArms.CondPower(mu1=0.32, sigma1=2, mu2=0.4, sigma2=1.8, n1=340, n2=340,
rho=0.3, tau=0.5, alpha2=0.024, alternative = "one.sided")
```

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TwoGrpCopriEndpt.SimPower

*Power Simulation for Two Group Two Coprimary Endpoints Group Sequential Design.*

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**Description**

Given the group sequential design information, returns the simulated overall power.

**Usage**

```
TwoGrpCopriEndpt.SimPower(mu11,mu12, mu21, mu22, rho, tau,
alpha1, alpha2, alternative , Nmax, B)
```

**Arguments**

mu11	standardized mean value for coprimary endpoint 1 in group 1.
mu12	standardized mean value for coprimary endpoint 2 in group 1.
mu21	standardized mean value for coprimary endpoint 1 in group 2.
mu22	standardized mean value for coprimary endpoint 2 in group 2.
rho	correlation coefficient between two coprimary endpoints.
tau	information time for the interim analysis.
alpha1	significant level for the first stage.
alpha2	significant level for the second stage.
alternative	indicates the alternative hypothesis and must be one of "two.sided" or "two.sided".
Nmax	maximum sample size per group.
B	the simulation iterative time.

**Value**

The evaluated power with attributes and computational error.

**Author(s)**

Yalin Zhu

## References

Chang, M. (2014). Adaptive design theory and implementation using SAS and R. *CRC Press*.

## Examples

```
# Example in Chang (2014) page 275
TwoGrpCopriEndpt.SimPower(mu11=0.2,mu12=0.25, mu21=0.005, mu22=0.015, rho=0.25,
tau=0.5, alpha1=0.0025, alpha2=0.024, alternative = "two.sided",Nmax=584, B=10000)
```

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