

# Package ‘MinEDfind’

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**Type** Package

**Title** A Bayesian Design for Minimum Effective Dosing-Finding Trial

**Version** 0.1.3

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**Description** The nonparametric two-stage Bayesian adaptive design is a novel phase II clinical trial design for finding the minimum effective dose (MinED). This design is motivated by the top priority and concern of clinicians when testing a new drug, which is to effectively treat patients and minimize the chance of exposing them to subtherapeutic or overly toxic doses. It is used to design single-agent trials.

**License** GPL-2

**Encoding** UTF-8

**LazyData** true

**Depends** Iso, ggplot2, gridExtra

**NeedsCompilation** no

**Repository** CRAN

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get.OC.MinED	<i>Generate operating characteristics for finding the minimum effective dose (MinED)</i>
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### Description

Obtain the operating characteristics of the nonparametric two-stage Bayesian adaptive design for minimum effective dose (MinED)-based dosing-finding trials

### Usage

```
get.OC.MinED(ttox, teff, phi_t, phi_e, ct = 0.95, eps_t, eps_e, d0 = 1,
             cohortsize = 3, ncohort1, ncohort2, ntrial = 100, extrasafe = TRUE,
             cutoff.eli = 0.95, n.earlystop = 12)
```

### Arguments

ttox	a vector containing the true toxicity rates of the investigational dose levels
teff	a vector containing the true response rates of the investigational dose levels
phi_t	the target DLT rate
phi_e	the target response rate
ct	the cutoff used to eliminate the dose for too toxicity. The default value is ct = 0.95
eps_t	a small value such that $(\text{phi}_t - \text{eps}_t, \text{phi}_t + \text{eps}_t)$ is an indifference interval of $\text{phi}_t$ . The default value is $\text{eps}_t = 0.1 * \text{phi}_t$
eps_e	a small value such that $(\text{phi}_e - \text{eps}_e, \text{phi}_e + \text{eps}_e)$ is an indifference interval of $\text{phi}_e$ . The default value is $\text{eps}_e = 0.1 * \text{phi}_e$
d0	the starting dose level. The default value is $d_0 = 1$
cohortsize	the cohort size
ncohort1	the number of cohort used in stage I
ncohort2	the number of cohort used in stage II
ntrial	the number of simulated trial
extrasafe	extrasafe set extrasafe = TRUE to impose a more stringent stopping rule
cutoff.eli	the cutoff to eliminate an overly toxic dose for safety. The default value is cutoff.eli = 0.95
n.earlystop	the early stopping parameter. The default value is n.earlystop = 12

### Value

get.oc.MinED() returns the operating characteristics of nonparametric two-stage Bayesian adaptive design as a matrix object, including: (1) true DLT rate at each dose level, (2) true efficacy rate at each dose level, (3) selection percentage at each dose level, (4) the average number of patients treated at each dose level, (5) the average number of patients responded to toxicity at each dose level, (6) the average number of patients responded to efficacy at each dose level

**Author(s)**

Chia-Wei Hsu, Fang Wang, Rongji Mu, Haitao Pan, Guoying Xu

**References**

Rongji Mu, Guoying Xu, Haitao Pan (2020). A nonparametric two-stage Bayesian adaptive design for minimum effective dose (MinED)-based dosing-finding trials, (under review)

**Examples**

```
ttox = c(0.05, 0.15, 0.3, 0.45, 0.6)
teff = c(0.05, 0.15, 0.3, 0.45, 0.6)
phi_t = 0.3
phi_e = 0.3
eps_t = 0.1 * phi_t
eps_e = 0.1 * phi_e

oc = get.OC.MinED(ttox = ttox, teff = teff, phi_t = phi_t, phi_e = phi_e,
                 eps_t = eps_t, eps_e = eps_e, cohortsize = 3, ncohort1 = 6,
                 ncohort2 = 14, ntrial = 100)

print(oc)
```

---

next.MinED

*Determine the dose for the next cohort of new patients for single-agent trials that aim to find a minimum effective dose (MinED)*

---

**Description**

Determine the dose for the next cohort of new patients for single-agent trials that aim to find a MinED

**Usage**

```
next.MinED(n, y, z, d, phi_t, phi_e, eps_t, eps_e, ct = 0.95, N1 = 18)
```

**Arguments**

n	a vector of number of patients treated at each dose level
y	a vector of number of patients experiencing the toxicity at each dose level (with the same length as candidate doses)
z	a vector of number of patients showing response at each dose level (with the same length as candidate doses)
d	the starting dose level
phi_t	the target DLT rate
phi_e	the target response rate

eps_t	a small value such that $(\text{phi}_t - \text{eps}_t, \text{phi}_t + \text{eps}_t)$ is an indifference interval of $\text{phi}_t$ . The default value is $\text{eps}_t = 0.1 * \text{phi}_t$
eps_e	a small value such that $(\text{phi}_e - \text{eps}_e, \text{phi}_e + \text{eps}_e)$ is an indifference interval of $\text{phi}_e$ . The default value is $\text{eps}_e = 0.1 * \text{phi}_e$
ct	the cutoff used to eliminate the dose for too toxicity. The default value is $\text{ct} = 0.95$
N1	number of trials in the stage 1. The default value is $N1 = 18$

**Value**

next.MinED() returns recommended dose level for the next cohort as a list ( $\$nextdose$ )

**Author(s)**

Chia-Wei Hsu, Fang Wang, Rongji Mu, Haitao Pan, Guoying Xu

**References**

Rongji Mu, Guoying Xu, Haitao Pan (2020). A nonparametric two-stage Bayesian adaptive design for minimum effective dose (MinED)-based dosing-finding trials, (under review)

**Examples**

```
n = c(3, 6, 0, 0, 0)
y = c(0, 1, 0, 0, 0)
z = c(0, 1, 0, 0, 0)
d = 2
phi_t = 0.3
phi_e = 0.3
eps_t = 0.1 * phi_t
eps_e = 0.1 * phi_e
next.dose <- next.MinED(n = n, y = y, z = z, d = d, phi_t = phi_t,
                        phi_e = phi_e, eps_t = eps_t, eps_e = eps_e)
print(next.dose)
```

---

plot.MinED

*Plot the simulation results for nonparametric two-stage Bayesian adaptive designs*

---

**Description**

Plot the objects returned by other functions, including (1) operating characteristics of the design, including selection percentage and the number of patients treated at each dose; (2) the estimates of toxicity and response probability for each dose in the admissible set and corresponding 95% credible interval

**Usage**

```
## S3 method for class 'MinED'  
plot(x, name, ...)
```

**Arguments**

x	the object returned by other functions
name	the name in the object to be plotted
...	ignored arguments

**Value**

plot.MinED() returns a figure

**Author(s)**

Chia-Wei Hsu, Fang Wang, Rongji Mu, Haitao Pan, Guoying Xu

**References**

Rongji Mu, Guoying Xu, Haitao Pan (2020). A nonparametric two-stage Bayesian adaptive design for minimum effective dose (MinED)-based dosing-finding trials, (under review)

**Examples**

```
## select the MinED based on the trial data  
n = c(3, 6, 0, 0, 0)  
y = c(0, 1, 0, 0, 0)  
z = c(0, 1, 0, 0, 0)  
phi_t = 0.3  
phi_e = 0.3  
eps_t = 0.1 * phi_t  
eps_e = 0.1 * phi_e  
select.dose <- select.MinED(n, y, z, phi_t, phi_e, eps_t, eps_e, ct = 0.95)  
plot.MinED(select.dose)  
  
## get the operating characteristics for nonparametric two-stage Bayesian adaptive designs  
ttox = c(0.05, 0.15, 0.3, 0.45, 0.6)  
teff = c(0.05, 0.15, 0.3, 0.45, 0.6)  
phi_t = 0.3  
phi_e = 0.3  
eps_t = 0.1 * phi_t  
eps_e = 0.1 * phi_e  
  
oc = get.OC.MinED(ttox = ttox, teff = teff, phi_t = phi_t, phi_e = phi_e,  
                 eps_t = eps_t, eps_e = eps_e, cohortsize=3, ncohort1 = 6,  
                 ncohort2 = 14, ntrial = 100)  
  
plot.MinED(oc, "Sel%")  
plot.MinED(oc, "#Pts.treated")  
plot.MinED(oc, "#Pts.response.to.tox")
```

```
plot.MinED(oc, "#Pts.response.to.eff")
```

---

```
select.MinED
```

*Select the minimum effective dose (MinED) for single agent trials*

---

### Description

Select the minimum effective dose (MinED) when the trial is completed

### Usage

```
select.MinED(n, y, z, phi_t, phi_e, eps_t, eps_e, ct = 0.95)
```

### Arguments

n	a vector of number of patients treated at each dose level
y	a vector of number of patients experiencing the toxicity at each dose level (with the same length as candidate doses)
z	a vector of number of patients showing response at each dose level (with the same length as candidate doses)
phi_t	the target DLT rate
phi_e	the target response rate
eps_t	a small value such that $(\text{phi}_t - \text{eps}_t, \text{phi}_t + \text{eps}_t)$ is an indifference interval of $\text{phi}_t$ . The default value is $\text{eps}_t = 0.1 * \text{phi}_t$
eps_e	a small value such that $(\text{phi}_e - \text{eps}_e, \text{phi}_e + \text{eps}_e)$ is an indifference interval of $\text{phi}_e$ . The default value is $\text{eps}_e = 0.1 * \text{phi}_e$
ct	the cutoff used to eliminate the dose for too toxicity. The default value is $\text{ct} = 0.95$

### Value

select.MinED() returns the selected dose with detailed information as a list, including: (1) selected dose level (`$Selected_Dose`), (2) target level for efficacy and toxicity rate (`$Target_Level`), (3) posterior estimate of efficacy and toxicity with its corresponding lower and upper bound etc. (`$Info`)

### Author(s)

Chia-Wei Hsu, Fang Wang, Rongji Mu, Haitao Pan, Guoying Xu

### References

Rongji Mu, Guoying Xu, Haitao Pan (2020). A nonparametric two-stage Bayesian adaptive design for minimum effective dose (MinED)-based dosing-finding trials, (under review)

**Examples**

```
n = c(3, 6, 0, 0, 0)
y = c(0, 1, 0, 0, 0)
z = c(0, 1, 0, 0, 0)
phi_t = 0.3
phi_e = 0.3
eps_t = 0.1 * phi_t
eps_e = 0.1 * phi_e
select.dose <- select.MinED(n, y, z, phi_t, phi_e, eps_t, eps_e)
print(select.dose)
```

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