

# Package ‘Opt5PL’

January 20, 2025

**Type** Package

**Title** Optimal Designs for the 5-Parameter Logistic Model

**Version** 0.1.1

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**Description** Obtain and evaluate various optimal designs for the 3, 4, and 5-parameter logistic models. The optimal designs are obtained based on the numerical algorithm in Hyun, Wong, Yang (2018) <[doi:10.18637/jss.v083.i05](https://doi.org/10.18637/jss.v083.i05)>.

**License** GPL-2

**Encoding** UTF-8

**LazyData** true

**Imports** Rcpp (>= 0.12.18), matrixcalc

**LinkingTo** Rcpp

**RoxygenNote** 6.1.0

**NeedsCompilation** yes

**Repository** CRAN

**Date/Publication** 2018-10-06 21:00:15 UTC

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**c\_weight***One iteration to run Newton Raphson to get c-optimal weights***Description**

This is a sub-part of running Newton Raphson method to search c-optimal weights for the given design points. This function provides the obtained c-optimal weights after one iteration of Newton Raphson method.

**Usage**

```
c_weight(W, T, X, d, p, order, UB, I)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values
X	A numeric vector. K dose levels for a given design
d	Numeric. Step adjustment
p	Numeric. Define EDp
order	Numeric. The number of model paraemters
UB	Numeric. The upper bound of the design points
I	A numeric matrix. Identity matrix.

c\_weight\_1

*The first derivative of the c-optimality criterion w.r.t the model parameters***Description**

The first derivative of the c-optimality criterion with respect to the model parameters. This is a sub-function of c\_weight to run one iteration of Newton-Raphson method.

**Usage**

```
c_weight_1(W, T, X, inv, p, order)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values
X	A numeric vector. K dose levels for a given design
inv	A numeric matrix. Inverse of the information matrix.
p	A numeric. Define EDp
order	numeric. The number of model paraemters

**c\_weight\_2**

*The second derivative of the c-optimality criterion with respect to the model parameters*

**Description**

The second derivative of the c-optimality criterion with respect to the model parameters. This is a sub-function of `c_weight` to run one iteration of Newton-Raphson method.

**Usage**

```
c_weight_2(W, T, X, inv, p, order)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values
X	A numeric vector. K dose levels for a given design
inv	A numeric matrix. Inverse of the information matrix.
p	A numeric. Define EDp
order	numeric. The number of model paraemters

**D1**

*Computing each element of the function c\_weight\_1*

**Description**

This function computes each element of the vector function `c_weight_1` which is the first derivative of the c-optimality criterion with respect to the model parameters.

**Usage**

```
D1(T, x, xl, inv, p, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x	A numeric. ith dose level for a given design
xl	A numeric. last dose level for a given design
inv	A numeric matrix. Inverse of the information matrix
p	A numeric. Define EDp
order	numeric. The number of model paraemters

d11

*Computing each element of the function DD\_weight\_1***Description**

This function computes each element of the vector function DD\_weight\_1 which is the first derivative of the Ds-optimality criterion with respect to the model parameters.

**Usage**

```
d11(T, x, x1, inv, inv1, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x	A numeric. ith dose level for a given design
x1	A numeric. last dose level for a given design
inv	A numeric matrix. Inverse of the information matrix for the 5PL model
inv1	A numeric matrix. Inverse of the information matrix for the 4PL model
order	numeric. The number of model paraemters

DD1

*Computing each element of the function c\_weight\_2***Description**

This function computes each element of the matrix function c\_weight\_2 which is the second derivative of the c-optimality criterion with respect to the model parameters.

**Usage**

```
DD1(T, x1, x2, x1, inv, p, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x1	A numeric. ith dose level for a given design
x2	A numeric. jth dose level for a given design
x1	A numeric. last dose level for a given design
inv	A numeric matrix. Inverse of the information matrix
p	A numeric. Define EDp
order	numeric. The number of model paraemters

dd11

*Computing each element of the function DD\_weight\_2***Description**

This function computes each element of the matrix function DD\_weight\_2 which is the second derivative of the Ds-optimality criterion with respect to the model parameters.

**Usage**

```
dd11(T, x1, x2, x1, inv, inv1, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x1	A numeric. ith dose level for a given design
x2	A numeric. jth dose level for a given design
x1	A numeric. last dose level for a given design
inv	A numeric matrix. Inverse of the information matrix for the 5PL model
inv1	A numeric matrix. Inverse of the information matrix for the 4PL model
order	numeric. The number of model paraemters

DD\_weight

*One iteration to run Newton Raphson to get Ds-optimal weights***Description**

This is a sub-part of running Newton Raphson method to search Ds-optimal weights for the given design points. This is function probvides the obtained Ds-optimal weights after one iteration of Newton Raphson method.

**Usage**

```
DD_weight(W, T, X, d, I4, I5, order)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values
X	A numeric vector. K dose levels for a given design
d	Numeric. Step adjustment
I4	A numeric matrix. Information matrix for the 4PL model
I5	A numeric matrix. Information matrix for the 5PL model
order	Numeric. The number of model paraemters

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DD_weight_1	<i>The first derivative of the Ds-optimality criterion with respect to the model parameters</i>
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**Description**

The first derivative of the Ds-optimality criterion with respect to the model parameters. This is a sub-function of D\_weight to run one iteration of Newton-Raphson method.

**Usage**

```
DD_weight_1(W, T, X, inv, inv1, order)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values for the 5PL model
X	A numeric vector. K dose levels for a given design
inv	A numeric matrix. Inverse information matrix for the 5PL model
inv1	A numeric matrix. Inverse information matrix for the 4PL model
order	numeric. The number of model paraemters

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DD_weight_2	<i>The second derivative of the Ds-optimality criterion with respect to the model parameters</i>
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**Description**

The second derivative of the Ds-optimality criterion with respect to the model parameters. This is a sub-function of D\_weight to run one iteration of Newton-Raphson method.

**Usage**

```
DD_weight_2(W, T, X, inv, inv1, order)
```

**Arguments**

W	A numeric vector. The first K-1 weights for a given design
T	A numeric vector. Model parameter values for the 5PL model
X	A numeric vector. K dose levels for a given design
inv	A numeric matrix. Inverse information matrix for the 5PL model
inv1	A numeric matrix. Inverse information matrix for the 4PL model
order	numeric. The number of model paraemters

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**D<sub>eff</sub>***Obtaining D-efficiency for estimating model parameters*

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**Description**

Obtaining D-efficiency for estimating model parameters for a given design under the 3, 4, 5-parameter logistic models. For the output, the function shows three things: 1. the obtained D-optimal design for the given parameter values; 2. the D-efficiency of the given design; and 3. the number of subjects required for the given design in order to provide the same performance as the D-optimal design does.

**Usage**

```
Deff(weight,dose,model,P,LB,UB,grid,N_dose)
```

**Arguments**

weight	A numeric vector. The first K-1 weights for a given design with K design points(K dose levels). The weights represent the proportional allocations of subjects to the dose levels in a given design.
dose	A numeric vector. K dose levels for a given design.
model	Numeric. The number of parameters in the model. This is used to specify the model to check the D-efficiency.
P	A numeric vector. Solicited information on nominal values for the vector for the 5PL model. P=( p <sub>1</sub> , p <sub>2</sub> , p <sub>3</sub> , p <sub>4</sub> , p <sub>5</sub> ), where p <sub>1</sub> is the upper limit of the response, p <sub>2</sub> is the slope that control the stiffness of the response curve, p <sub>3</sub> is the position of the transition region in dose(this is the ED50 under the 3PL and the 4PL model), p <sub>4</sub> is the lower limit of the response, and p <sub>5</sub> is the asymmetric factor. When a user computes the D-efficiency under the 3PL or 4PL model, P is the vector of model parameter values for each model with setting p <sub>4</sub> and p <sub>5</sub> differently. Details are given in the examples.
LB	Numeric. Predetermined lower bound of the dose range for the original dose level.
UB	Numeric. Predetermined lower bound of the dose range for the original dose level.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE

**Author(s)**

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

## References

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K, Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

## Examples

```
##Under the 5PL model with the parameter values
##T5=(30000,0.5,800,0.5,2) and the dose range [1.95,32000],
##find the D-efficiency of the broad range design.

##The broad range design
dose=c(1.95,7.8,31.25,125,500,2000,8000,32000)
dlength=length(dose)
weight=rep(1/dlength,dlength-1)

##Parameter values
T5=c(30000,0.5,800,0.5,2)

##Compute D-efficiency of the broad range design
Deff(weight,dose,model=5,P=T5,LB=1.95,UB=32000,grid=.01)
```

Dp

*Target dose, ED<sub>p</sub>*

## Description

This finds the target dose level ED<sub>p</sub> which is the dose producing p percentage of the difference between the maximum and the minimum responses.

## Usage

```
Dp(T, p)
```

## Arguments

- |   |  |
|---|--|
| T | A numeric vector. Model parameter values |
| p | A numeric. Define ED <sub>p</sub>        |

DS1

*Sensitivity function of c-optimality criterion for the ED<sub>p</sub>***Description**

This evaluates the sensitivity function of the c-optimality criterion for the ED<sub>p</sub> at the given value of x.

**Usage**

```
DS1(T, x, inv, p, order)
```

**Arguments**

T	A numeric vector. Model parameter values for the 5PL model
x	A numeric. A given dose level
inv	A numeric matrix. Inverse information matrix for the 5PL model
p	A numeric. Define ED <sub>p</sub>
order	numeric. The number of model parameters

ds11

*Sensitivity function of Ds-optimality criterion***Description**

This evaluates the sensitivity function of the Ds-optimality criterion at the given value of x.

**Usage**

```
ds11(T, x, inv, inv1, order)
```

**Arguments**

T	A numeric vector. Model parameter values for the 5PL model
x	A numeric. A given dose level
inv	A numeric matrix. Inverse information matrix for the 5PL model
inv1	A numeric matrix. Inverse information matrix for the 4PL model
order	numeric. The number of model parameters

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Ds eff	<i>Obtaining Ds-efficiency for estimating the asymmetric factor under the 5-parameter logistic model.</i>
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## Description

Obtaining Ds-efficiency for estimating the asymmetric factor for a given design under the 5-parameter logistic models. For the output, the function shows three things: 1. the obtained Ds-optimal design for the given parameter values; 2. the Ds-efficiency of the given design; and 3. the number of subjects required for the given design in order to provide the same performance as the Ds-optimal design does.

## Usage

```
Ds eff(weight, dose, P, LB, UB, r, epsilon, grid, N_dose)
```

## Arguments

weight	A numeric vector. The first K-1 weights for a given design with K design points(K dose levels). The weights represent the proportional allocations of subjects to the dose levels in a given design.
dose	A numeric vector. K dose levels for a given design.
P	A numeric vector. Solicited information on nominal values for the vector for the SPL model. P=(p1, p2, p3, p4, p5), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, p3 is the position of the transition region in dose, p4 is the lower limit of the response, and p5 is the asymmetric factor.
LB	Numeric. Predetermined lower bound of the dose range for the original dose level.
UB	Numeric. Predetermined lower bound of the dose range for the original dose level.
r	Numeric. The number of iterations to set an initial design to search the Ds-optimal design to compute the efficiency. Default is 30 and needed to be increased (for example, r=50) if the searched the Ds-optimal design is not a true optimal. It can be verified by the verification plot.
epsilon	Numeric. Stopping criterion for the algorithm to search the Ds-optimal design. Default is 0.001.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE

## Author(s)

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

## References

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K., Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

## Examples

```
##Under the 5PL model with the parameter values
##T5=(30000,0.5,800,0.5,2) and the dose range [1.95,32000],
##find the Ds-efficiency of the broad range design.

##The broad range design
dose=c(1.95,7.8,31.25,125,500,2000,8000,32000)
dlength=length(dose)
weight=rep(1/dlength,dlength-1)

##Parameter values for the 5PL model
T5=c(30000,0.5,800,0.5,2)

##Ds efficiency of the broad range design
Dseff(weight,dose,P=T5,LB=1.95,UB=32000,grid=.01)
```

DsOPT

*Search Ds-optimal design for estimating the asymmetric factor under the 5-parameter logistic model.*

## Description

Obtain Ds-optimal design (i.e., optimal dose levels and corresponding optimal weights) for estimating the asymmetric factor under the 5-parameter logistic model.

## Usage

```
DsOPT(LB,UB,P,grid,r,epsilon,N_dose,log_scale)
```

## Arguments

- |    |   |
|----|---|
| LB | Numeric. Predetermined lower bound of the dose range for the original dose level. |
| UB | Numeric. Predetermined lower bound of the dose range for the original dose level. |

P	A numeric vector. Solicited information on nominal values for the vector for the 5PL model. P=(p1, p2, p3, p4, p5), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, p3 is the position of the transition region in dose, p4 is the lower limit of the response, and p5 is the asymmetric factor.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
r	Numeric. The number of iterations to set an initial design to search the Ds-optimal design. Default is 30 and needed to be increased (for example, r=50) if the searched optimal design is not a true optimal. It can be verified by the verification plot.
epsilon	Numeric. Stopping criterion for the algorithm to search the Ds-optimal design. Default is 0.001.
N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE
log_scale	Set to TRUE if a user wants to set the X-axis(dose level) on log scale in the verification plot of the optimal design. Otherwise, set to FALSE. Default is TRUE. Note: this need to set FALSE when the original dose levels take negative values.

### Author(s)

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

### References

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K, Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

### Examples

```
##Find the Ds-optimal design for estimating theta_5 under the 5PL model
##with the parameter values T5=(30000,0.5,800,0.5,2) and the dose range [1.95,32000].

##Parameter values for the 5PL model
T5=c(30000,0.5,800,0.5,2)

##Find the Ds-optimal design
DsOPT(LB=1.95, UB=32000, P=T5,grid=.01,epsilon=.001)
```

**D\_weight***One iteration to run Newton Raphson to get D-optimal weights***Description**

This is a sub-part of running Newton Raphson method to search D-optimal weights for the given design points. This function provides the obtained D-optimal weights after one iteration of Newton Raphson method.

**Usage**

```
D_weight(W, T, X, d, q)
```

**Arguments**

<b>W</b>	A numeric vector. The first K-1 weights for a given design
<b>T</b>	A numeric vector. Model parameter values
<b>X</b>	A numeric vector. K dose levels for a given design
<b>d</b>	Numeric. Step adjustment
<b>q</b>	A numeric vector. User select weights defined in the function RDOPT

**D\_weight\_1***The first derivative of the D-optimality criterion w.r.t the model parameters***Description**

The first derivative of the D-optimality criterion with respect to the model parameters. This is a sub-function of D\_weight to run one iteration of Newton-Raphson method.

**Usage**

```
D_weight_1(q, W, T1, T2, T3, X, inv1, inv2, inv3)
```

**Arguments**

<b>q</b>	A numeric vector. User select weights defined in the function RDOPT
<b>W</b>	A numeric vector. The first K-1 weights for a given design
<b>T1</b>	A numeric vector. Model parameter values for the 3PL model
<b>T2</b>	A numeric vector. Model parameter values for the 4PL model
<b>T3</b>	A numeric vector. Model parameter values for the 5PL model
<b>X</b>	A numeric vector. K dose levels for a given design
<b>inv1</b>	A numeric matrix. Inverse information matrix for the 3PL model
<b>inv2</b>	A numeric matrix. Inverse information matrix for the 4PL model
<b>inv3</b>	A numeric matrix. Inverse information matrix for the 5PL model

---

D_weight_2	<i>The second derivative of the D-optimality criterion w.r.t the model parameters</i>
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**Description**

The second derivative of the D-optimality criterion with respect to the model parameters. This is a sub-function of D\_weight to run one iteration of Newton-Raphson method.

**Usage**

```
D_weight_2(q, W, T1, T2, T3, X, inv1, inv2, inv3)
```

**Arguments**

q	A numeric vector. User select weights defined in the function RDOPT
W	A numeric vector. The first K-1 weights for a given design
T1	A numeric vector. Model parameter values for the 3PL model
T2	A numeric vector. Model parameter values for the 4PL model
T3	A numeric vector. Model parameter values for the 5PL model
X	A numeric vector. K dose levels for a given design
inv1	A numeric matrix. Inverse information matrix for the 3PL model
inv2	A numeric matrix. Inverse information matrix for the 4PL model
inv3	A numeric matrix. Inverse information matrix for the 5PL model

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EDpeff	<i>Obtaining c-efficiency for estimating the EDp under the 5-parameter logistic model.</i>
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**Description**

Obtaining c-efficiency for estimating the EDp for a given design under the 5-parameter logistic models. For the output, the function shows three things: 1. the obtained c-optimal design for the given parameter values; 2. the c-efficiency of the given design; and 3. the number of subjects required for the given design in order to provide the same performance as the c-optimal design does.

**Usage**

```
EDpeff(weight,dose,P,EDp,LB,UB,r,epsilon,grid,N_dose)
```

### Arguments

weight	A numeric vector. The first K-1 weights for a given design with K design points(K dose levels). The weights represent the proportional allocations of subjects to the dose levels in a given design.
dose	A numeric vector. K dose levels for a given design.
P	A numeric vector. Solicited information on nominal values for the vector for the SPL model. P=(p1, p2, p3, p4, p5), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, p3 is the position of the transition region in dose, p4 is the lower limit of the response, and p5 is the asymmetric factor.
EDp	Numeric. An user select value to define the EDp. For example, EDp=.05 to define the ED50.
LB	Numeric. Predetermined lower bound of the dose range for the original dose level.
UB	Numeric. Predetermined lower bound of the dose range for the original dose level.
r	Numeric. The number of iterations to set an initial design to search the c-optimal design for the EDp in order to compute the efficiency. Default is 30 and needed to be increased (for example, r=50) if the searched the c-optimal design is not a true optimal. It can be verified by the verification plot.
epsilon	Numeric. Stopping criterion for the algorithm to search the c-optimal design. Default is 0.001.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE

### Author(s)

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

### References

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K, Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

### Examples

```
##Under the 5PL model with the parameter values
##T5=(30000,0.5,800,0.5,2) and the dose range [1.95,32000],
##find the c-efficiency of the broad range design.

##The broad range design
```

```

dose=c(1.95,7.8,31.25,125,500,2000,8000,32000)
dlength=length(dose)
weight=rep(1/dlength,dlength-1)

##Parameter values for the 5PL model
T5=c(30000,0.5,800,0.5,2)

##Compute the c-efficiency of the broad range design
EDpeff(weight,dose,P=T5,EDp=.5,LB=1.95,UB=32000,grid=.01)

```

**EDpOPT**

*Search c-optimal designs for estimating the ED<sub>p</sub> under the 5-parameter logistic model*

**Description**

Obtain c-optimal design (i.e., optimal dose levels and corresponding optimal weights) for estimating the ED<sub>p</sub> under the 5-parameter logistic model. The ED<sub>p</sub> is the dose level that achieve the 100p% of the difference between the maximum and the minimum responses.

**Usage**

```
EDpOPT(LB,UB,P,EDp,grid,r,epsilon,N_dose,log_scale)
```

**Arguments**

LB	Numeric. Predetermined lower bound of the dose range for the original dose level.
UB	Numeric. Predetermined lower bound of the dose range for the original dose level.
P	A numeric vector. Solicited information on nominal values for the vector for the 5PL model. P=(p <sub>1</sub> , p <sub>2</sub> , p <sub>3</sub> , p <sub>4</sub> , p <sub>5</sub> ), where p <sub>1</sub> is the upper limit of the response, p <sub>2</sub> is the slope that control the stiffness of the response curve, p <sub>3</sub> is the position of the transition region in dose, p <sub>4</sub> is the lower limit of the response, and p <sub>5</sub> is the asymmetric factor.
EDp	Numeric. An user select value to define the ED <sub>p</sub> . For example, EDp=.05 to define the ED50.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
r	Numeric. The number of iterations to set an initial design to search the c-optimal design. Default is 30 and needed to be increased (for example, r=50) if the searched optimal design is not a true optimal. It can be verified by the verification plot.
epsilon	Numeric. Stopping criterion for the algorithm to search the c-optimal design. Default is 0.001.

N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE
log_scale	Set to TRUE if a user wants to set the X-axis(dose level) on log scale in the verification plot of the optimal design. Otherwise, set to FALSE. Default is TRUE. Note: this need to set FALSE when the original dose levels take negative values.

**Author(s)**

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

**References**

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K., Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

**Examples**

```
##Search the c-optimal design for estimating the ED50 under the 5PL model

##Parameter values for the 5PL model
T5=c(30000,0.5,800,0.5,2)

##Dose range=[1.95,32000]

EDpOPT(LB=1.95,UB=32000, P=T5,EDp=.5,grid=.01, r =30,epsilon=.001)
```

**Description**

This gives the first derivative of the mean response function (3, 4, or 5PL) with respect to the model parameters.

**Usage**

f(T, x, order)

**Arguments**

T	A numeric vector. Model parameter values
x	A numeric. A given dose level
order	numeric. The number of model paraemters

---

g

*Partial derivative of the EDp with respect to the model parameters*

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### Description

Partial derivative of the EDp with respect to the model parameters

### Usage

g(T, p)

### Arguments

T	A numeric vector. Model parameter values
p	A numeric. Define EDp

---

ginv

*Generalized Inverse Matrix*

---

### Description

Obtain generalized inverse of the given matrix X.

### Usage

ginv(X, tol = sqrt(.Machine\$double.eps))

### Arguments

X	A numeric matrix
tol	Default

**infor***Obtain a information matrix at a single design point***Description**

Obtain a information matrix at a single given design point x.

**Usage**

```
infor(T, x, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x	numeric. A single design point(dose level)
order	numeric. The number of model paraemters

**Inv***Adjusting inverse information matrix being not singular***Description**

When the inverse matrix is a singular, it is adjusted to become nonsingular matrix.

**Usage**

```
Inv(M, I)
```

**Arguments**

M	A numeric matrix
I	Identity matrix

---

Minus

*Matrix subtraction*

---

**Description**

Matrix subtraction: M1-M2.

**Usage**

Minus(M1, M2)

**Arguments**

M1	A numeric matrix
M2	A numeric matrix

---

Multiple

*Matrix multiplication*

---

**Description**

Matrix multiplication: M1\*M2.

**Usage**

Multiple(M1, M2)

**Arguments**

M1	A numeric matrix
M2	A numeric matrix

---

Plus

*Matrix addition*

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**Description**

Matrix addition: M1+M2.

**Usage**

Plus(M1, M2)

**Arguments**

M1	A numeric matrix
M2	A numeric matrix

---

RDOPT*Search the robust D-optimal designs for estimating model parameters*

---

### Description

Obtain robust D-optimal design (i.e., optimal dose levels and corresponding optimal weights) for estimating the model parameters that works well under the 3, 4, 5-parameter logistic models. In addition, the function can be used to obtain D-optimal design for each of the three models.

### Usage

```
RDOPT(LB,UB, P3,P4,P5,q,grid,r,epsilon,N_dose,log_scale)
```

### Arguments

LB	Numeric. Predetermined lower bound of the dose range for the original dose level.
UB	Numeric. Predetermined lower bound of the dose range for the original dose level.
P3	A numeric vector. Solicited information on nominal values for the vectors for the 3PL model. P3=(p1, p2, p3), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, and p3 is the ED50.
P4	A numeric vector. Solicited information on nominal values for the vectors for the 4PL model. P4=(p1, p2, p3, p4), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, p3 is the ED50, and p4 is the lower limit of the response.
P5	A numeric vector. Solicited information on nominal values for the vectors for the 5PL model. P5=(p1, p2, p3, p4, p5), where p1 is the upper limit of the response, p2 is the slope that control the stiffness of the response curve, p3 is the position of the transition region in dose, p4 is the lower limit of the response, and p5 is the asymmetric factor. For searching the robust D-optimal design, the nominal values for P3, P4, P5 need to be specified. However, when a user searches the D-optimal design for the 3PL, 4PL, or 5PL model, only P5 needs to be specified. Details are given in the examples.
q	A numeric vector. User select weights. q=c(q1, q2, q3), where q1, q2, q3 represent the relative importance of the 3, 4, 5PL models respectively. They are non-negative and q1+q2<=1. For example, q=c(0,0,1) is used to find D-optimal design for the 5PL model only and q=c(1/3,1/3,1/3) is used to find robust D-optimal design for the three models under the assumption that the three models are equally important.
grid	Numeric. The grid density to discretize the predetermined dose interval. Default is 0.01.
r	Numeric. The number of iterations to set an initial design to search the D-optimal design. Default is 30 and needed to be increased (for example, r=50) if the searched optimal design is not a true optimal. It can be verified by the verification plot.

epsilon	Numeric. Stopping criterion for the algorithm to search the D-optimal design. Default is 0.001.
N_dose	Set to TRUE if the original dose levels take negative values. Set to FALSE if the original dose levels take positive values. Default is FALSE
log_scale	Set to TRUE if a user wants to set the X-axis(dose level) on log scale in the verification plot of the optimal design. Otherwise, set to FALSE. Default is TRUE. Note: this need to set FALSE when the original dose levels take negative values.

### Author(s)

Seung Won Hyun, Weng Kee Wong, and Yarong Yang

### References

- Hyun, S. W., Wong, W.K., Yang, Y. Optimal designs for asymmetric sigmoidal response curves in bioassays and immunoassays. (submitted to Statistical Methods in Medical Research)
- Hyun, S.W., Wong, W.K, Yang, Y. VNM: An R Package for Finding Multiple-Objective Optimal Designs for the 4-Parameter Logistic Model. (Journal of Statistical Software, 83, 1-19, 2018, doi:10.18637/jss.v083.i05.)

### Examples

```
##1. Search the robust D-optimal design for estimating the model parameters
##under the 3,4,5PL models

##Parameter values for the 5PL model
T5=c(30000,0.5,800,0.5,2)
##Parameter values for the 4PL model
T4=c(27264.92,0.67,3340.95,-225.55)
##Parameter values for the 3PL model
T3=c(26715.52,0.70,3204.92)

##Dose range=[1.95,32000]
##Assuming the three models are equally important
RDOPT(LB=1.95,UB=32000,P3=T3,P4=T4,P5=T5,q=c(1/3,1/3,1/3))

##2. Search the D-optimal design for the 3PL model using
##the parameter values T3=(26715.52,0.70,3204.92)
##and the dose range [1.95,32000].
```

```
#Set the parameter values for the 3PL model in the form of P5
P5=c(26715.52,0.70,3204.92,0,1)
RDOPT(LB=1.95, UB=32000,P5=P5,q=c(1,0,0))

##3. Search the D-optimal design for the 4PL model using
##the parameter values T4=(27264.92,0.67,3340.95,-225.55)
##and the dose range [1.95,32000].
```

```
#Set the parameter values for the 4PL model in the form of P5.
P5=c(27264.92,0.67,3340.95,-225.55,1)
```

---

```
RDOPT(LB=1.95, UB=32000,P5=P5,q=c(0,1,0))
```

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SDM	<i>Summation of diagonal elements in a matrix</i>
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### Description

Summation of diagonal elements in a matrix M.

### Usage

```
SDM(M)
```

### Arguments

M	A numeric matrix
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smalld1	<i>Sub-function of the function D_weight_I</i>
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### Description

This computes the first derivative of the D-optimality criterion with respect to the model parameters for a specified model.

### Usage

```
smalld1(T, x, x1, inv, order)
```

### Arguments

T	A numeric vector. Model parameter values
x	A numeric. ith dose level
x1	A numeric. last dose level
inv	A numeric matrix. Inverse information matrix
order	numeric. The number of model parameters: 3, 4, 5 for the 3, 4, 5PL models respectively

smalldd1

*Sub-function of the function D\_weight\_2***Description**

This computes the second derivative of the D-optimality criterion with respect to the model parameters for a specified model.

**Usage**

```
smalldd1(T, x1, x2, x1, inv, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x1	A numeric. ith dose level
x2	A numeric. jth dose level
x1	A numeric. last dose level
inv	A numeric matrix. Inverse information matrix
order	numeric. The number of model paraemters: 3, 4, 5 for the 3, 4, 5PL models respectively

smallds1

*Sensitivity function of D-optimality criterion***Description**

This evaluates the sensitivity function of the D-optimality criterion at the given value of x for a specified model.

**Usage**

```
smallds1(T, x, inv, order)
```

**Arguments**

T	A numeric vector. Model parameter values
x	A numeric. A given dose level
inv	A numeric matrix. Inverse information matrix
order	numeric. The number of model paraemters: 3, 4, 5 for the 3, 4, 5PL models respectively

**sMultiple**                    *Multiply a constant to a matrix*

### Description

Multiply a constant to a matrix: s\*M.

### Usage

```
sMultiple(s, M)
```

### Arguments

<b>s</b>	Numeric
<b>M</b>	A numeric matrix

**S\_weight**                    *Newton Raphson method to get optimal weights*

### Description

Newton Raphson method to get optimal weights for given design points "X".

### Usage

```
S_weight(X, T, e1, f, ...)
```

### Arguments

<b>X</b>	A numeric vector. K dose levels for a given design
<b>T</b>	A numeric vector. Model parameter values
<b>e1</b>	Numeric. Threshold to stop the iteration.
<b>f</b>	A character. D_weight for D-optimal weight; c_weight for c-optimal weight; DD_weight for Ds-optimal weight
<b>...</b>	Default.

---

Trans	<i>Transpose of a matrix</i>
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**Description**

Transpose of a matrix M.

**Usage**

Trans(M)

**Arguments**

M                   A numeric matrix

---

upinfor	<i>Obtain normalized Fisher information matrix</i>
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---

**Description**

Obtain normalized Fisher information matrix for the 3, 4, 5PL models.

**Usage**

upinfor(W, T, X, order)

**Arguments**

W                   A numeric vector. The first K-1 weights for a given design  
T                   A numeric vector. Model parameter values  
X                   A numeric vector. K design points for a given design  
order              numeric. The number of model paraemters

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