

R documentation

of all in ‘man’

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DesignEval	<i>Evaluate design in terms of criteria</i>
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Description

This function takes matrix X and crit to output the criterion value.

Usage

```
DesignEval(x, crit)
```

Arguments

x	an integer matrix object. Representing the design to be evaluated.
crit	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.

Value

criterion value.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
x = matrix(c(1,1,1,2,2,2,3,3,3),nrow=3,byrow=TRUE)
crit="MD2"
value = DesignEval(x,crit)
```

DesignPairPlot

Draw pair plot for design of experiments

Description

This function takes a design D and a boolean value Diag to draw pair plot.

Usage

```
DesignPairPlot(D,Diag)
```

Arguments

D a matrix object. Design of Experiment.
 Diag a boolean R object.

Value

A pair plot

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
##e.g.1
n=12 #(must be multiples of q)
s=3
q=4
crit = "MD2"#(Mixture L2 criteria)
D = DesignQuery(n=n,s=s,q=q,crit="MD2")
DesignPairPlot(D)

##e.g.2
n=12 #(must be multiples of q)
s=3
q=3
```

```
crit = "MD2"# (Mixture L2 criteria)
D = DesignQuery(n=n,s=s,q=q,crit="MD2")
DesignPairPlot(D,Diag=TRUE)
```

DesignQuery

Evaluate design in terms of criteria

Description

This function takes size of desired design,criterion crit. If the required design exists in database, then return the design, else return NULL.

Usage

```
DesignQuery(n,s,q,crit,ShowCrit)
```

Arguments

n	an integer R object. Run of Experiment
s	an integer R object. Factor of Experiment.
q	an integer R object. Level of Experiment.
crit	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
ShowCrit	If TRUE, print CD2,MD2,maximin value of required design. Default: TRUE

Value

Desired design

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
n = 9
s = 3
q = 3
crit="MD2"
D = DesignQuery(n,s,q,crit)
D
```

GenAUD

*Generating sequential Uniform Design of Experiments (Augmenting Runs)***Description**

This function takes n, s, q ; a unchanged initial design and other arguments to output a list (described below).

Usage

```
GenAUD(xp, n, s, q, initX, crit, maxiter, hits_ratio, levelpermt, rand_seed, vis)
```

Arguments

<code>xp</code>	an integer matrix R object. Representing the previous existing design matrix.
<code>n</code>	an integer R object. Run of Experiment (including the previous design <code>xp</code>).
<code>s</code>	an integer R object. Factor of Experiment.
<code>q</code>	an integer R object. Level of Experiment.
<code>init</code>	a string vector R object. Initialization method for the run-augmented design: "rand" – Randomly generate initial design (default); "input". – User specified.
<code>initX</code>	a user-defined integer matrix R object. This is the user-defined initial augmentation matrix, and will be used when <code>init="input"</code> .
<code>crit</code>	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
<code>maxiter</code>	a positive integer R object. Maximum iteration number in outer while loop of SOAT algorithm.
<code>hits_ratio</code>	a float R object. Default value is 0.1, which is the ratio to accept changes of design in inner for loop.
<code>levelpermt</code>	a boolean R object. It controls whether to use level permutation.
<code>rand_seed</code>	a integer R object. It controls the random seed used for optimization.
<code>vis</code>	a boolean variable. If true, plot the trace of criterion values.

Value

A list that contains Initial design matrix(`initial_design`), optimal design matrix(`final_design`), initial criterion value(`initial_criterion`), final criterion value(`final_criterion`) and criterion list(`criterion_history`) in update process.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
#Example.
#Set a fixed initial matrix:
n=12 # (must be multiples of q)
s=3
q=4
mat0 = matrix(c(1,1,1,2,2,2,3,3,3),ncol=3,byrow=TRUE)# nb. of columns=s
crit = "MD2" # (Mixture L2 criteria)
res = GenAUD(xp=mat0,n,s,q,crit=crit,maxiter=100,vis=TRUE)
```

GenAUD_COL

Generating sequential Uniform Design of Experiments (Augmenting Factors)

Description

This function takes n,s,q; a unchanged initial design and other arguments to output a list (described below).

Usage

```
GenAUD_COL(xp,n,s,q,init,initX,crit,maxiter,hits_ratio,levelpermt,rand_seed,vis)
```

Arguments

xp	an integer matrix R object, representing the previous existing design matrix.
n	an integer R object. Run of Experiment
s	an integer R object. Factor of Experiment (including the number of factors in previous design xp).
q	an integer R object. Level of Experiment.
init	a string vector R object. Initialization method for the factor-augmented design: "rand" –Randomly generate initial design (default); "input". –User specified.
initX	a user-defined integer matrix R object. This is the user-defined initial augmentation matrix, and will be used when init="input".
crit	a character R object. Type of criterion to use: "CD2" –Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" –Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimize Coherence ; "A2" – Minimize Average Chi-Square.
maxiter	a positive integer R object. Maximum iteration number in outer while loop of SOAT algorithm.

hits_ratio	a float R object. Default value is 0.1, which is the ratio to accept changes of design in inner for loop.
levelpermt	a boolean R object. It controls whether to use level permutation.
rand_seed	a integer R object. It controls the random seed used for optimization.
vis	a boolean variable. If true, plot the trace of criterion values.

Value

A list that contains Initial design matrix(initial_design), optimal design matrix(final_design), initial criterion value(initial_criterion), final criterion value(final_criterion) and criterion list(criterion_history) in update process.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
#Example.
#Set a fixed initial matrix:
n=6 # (must be multiples of q)
s=4
q=3
mat0 = matrix(c(1,2,2,1,1,3,3,1,2,3,3,2), ncol=2, byrow=TRUE)
crit = "MD2" # (Mixture L2 criteria)
res = GenAUD_COL(xp=mat0, n, s, q, crit=crit, maxiter=100, vis=TRUE)
```

GenAUD_COL_MS

Generating sequential Uniform Design of Experiments (Augmenting Factors) using different initial designs

Description

This function takes n,s,q and other arguments to output a list(described below).

Usage

```
GenAUD_COL_MS(xp, n, s, q, crit, maxiter, nshoot, rand_seed, vis=FALSE)
```

Arguments

xp	an integer matrix R object. Representing the previous existing design matrix.
n	an integer R object.
s	an integer R object. Factor of Experiment (including the number of factors in previous design xp).
q	an integer R object. Level of Experiment.

<code>crit</code>	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
<code>maxiter</code>	a positive integer R object. Maximum iteration number in outer while loop of SATA algorithm in each shoot.
<code>nshoot</code>	Total counts to try different initial designs.
<code>rand_seed</code>	a integer R object. It controls the random seed used for optimization.
<code>vis</code>	a boolean R object. If true, plot the criterion value sequence for all shoots.

Value

Best design over all shoots.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
D1 = DesignQuery(n=18, s=3, q=3, crit="MD2")
D2 = GenAUD_COL_MS(xp=D1, n=18, s=5, q=3, crit="MD2",
                   maxiter=100, nshoot=5)
```

GenAUD_MS

Generating sequential Uniform Design of Experiments (Augmenting Runs) using different initial designs

Description

This function takes n,s,q and other arguments to output a list(described below).

Usage

```
GenAUD_MS(xp, n, s, q, crit, maxiter, nshoot, rand_seed, vis=FALSE)
```

Arguments

<code>xp</code>	an integer matrix R object. Representing the previous existing design matrix.
<code>n</code>	an integer R object. Run of Experiment (including the previous design xp).
<code>s</code>	an integer R object. Factor of Experiment.
<code>q</code>	an integer R object. Level of Experiment.

<code>crit</code>	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
<code>maxiter</code>	a positive integer R object. Maximum iteration number in outer while loop of SATA algorithm in each shoot.
<code>nshoot</code>	Total counts to try different initial designs.
<code>rand_seed</code>	a integer R object. It controls the random seed used for optimization.
<code>vis</code>	a boolean R object. If true, plot the criterion value sequence for all shoots.

Value

Best design over all shoots.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
D1 = DesignQuery(n=18, s=7, q=3, crit="MD2")
D2 = GenAUD_MS(xp=D1, n=21, s=7, q=3, crit="MD2",
               maxiter=100, nshoot=5)
```

GenUD

Generating Uniform Design of Experiments

Description

This function takes `n,s,q` and other arguments to output a list(described below).

Usage

```
GenUD(n, s, q, init, initX, crit, maxiter, hits_ratio, levelpermt, rand_seed, vis)
```

Arguments

<code>n</code>	an integer R object. Run of Experiment.
<code>s</code>	an integer R object. Factor of Experiment.
<code>q</code>	an integer R object. Level of Experiment.
<code>init</code>	a string vector R object. Initialization method for the design: "rand" –Randomly generate initial design (default); "input". –User specified.
<code>initX</code>	a user-defined integer matrix R object. This is the user-defined initial design matrix, and will be used when <code>init="input"</code> .

<code>crit</code>	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
<code>maxiter</code>	a positive integer R object. Maximum iteration number in outer while loop of SATA algorithm.
<code>levelpermt</code>	a boolean R object. It controls whether to use level permutation.
<code>rand_seed</code>	a integer R object. It controls the random seed used for optimization.
<code>hits_ratio</code>	a float R object. Default value is 0.1, which is the ratio to accept changes of design in inner for loop.
<code>vis</code>	a boolean R object. If true, plot the criterion value sequence.

Value

A list that contains Initial design matrix(`initial_design`), optimal design matrix(`final_design`), initial criterion value(`initial_criterion`), final criterion value(`final_criterion`) and criterion list(`criterion_history`) in update process.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
##Example 1
n=12 # (must be multiples of q)
s=3
q=4
crit = "CD2" # (Centered L2 criteria)
res = GenUD(n,s,q,crit=crit,maxiter=100)

##Example 2
n=10 # (must be multiples of q)
s=3
q=5
init = "rand"
crit = "MD2" # (Mixture L2 criteria)
vis=TRUE
res=GenUD(n,s,q,init=init,crit=crit,maxiter=100,vis=vis)

##Example 3
#If init="input", algorithm will search for better a better design with same size as in
init = "input"
n=6
s=2
q=3
initX = matrix(c(1,1,2,2,3,3,3,3,1,1,2,2),ncol=2)
res = GenUD(n,s,q,init=init,initX=initX,maxiter=100)
```

GenUD_MS

*Generating Uniform Design of Experiments using different initial designs***Description**

This function takes `n,s,q,crit` and `nshoot` to return a design. `nshoot` number of random initial designs are used in each shoot. The design returned is the best design over all shoots.

Usage

```
GenUD_MS(n, s, q, crit, maxiter, nshoot, rand_seed, vis)
```

Arguments

<code>n</code>	an integer R object. Run of Experiment
<code>s</code>	an integer R object. Factor of Experiment.
<code>q</code>	an integer R object. Level of Experiment.
<code>crit</code>	a character R object. Type of criterion to use: "CD2" – Centered L2 Discrepancy (default) ; "WD2" – Wrap-around L2 Discrepancy ; "MD2" – Mixture L2 Discrepancy ; "maximin" – Maximin Discrepancy ; "MC" – Minimum Coherence ; "A2" – Mean Squared Correlation.
<code>maxiter</code>	a positive integer R object. Maximum iteration number in outer while loop of SATA algorithm in each shoot.
<code>nshoot</code>	Total counts to try different initial designs.
<code>rand_seed</code>	a integer R object. It controls the random seed used for optimization.
<code>vis</code>	a boolean R object. If true, plot the criterion value sequence for all shoots.

Value

Best design over all shoots.

Author(s)

Aijun Zhang, Zebin Yang, Haoyu Li, Shijie Quan

Examples

```
D = GenUD_MS(36, 4, 6, crit="CD2",
              maxiter=50, nshoot=6,
              vis=TRUE)
```

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