

# Package ‘optDesignSlopeInt’

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**Type** Package

**Title** Optimal Designs for Estimating the Slope Divided by the Intercept

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**Description** Aids practitioners to optimally design experiments that measure the slope divided by the intercept and provides confidence intervals for the ratio.

**Encoding** UTF-8

**License** GPL-3

**Depends** R (>= 4.0.0)

**Imports** stats, graphics, MCMCpack

**RoxygenNote** 7.2.3

**NeedsCompilation** no

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design_bakeoff	<i>A visualiation for comparing slope-divided-by-intercept estimates for a number of designs</i>
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## Description

A visualiation for comparing slope-divided-by-intercept estimates for a number of designs

## Usage

```
design_bakeoff(
  xmin,
  xmax,
  designs,
  gen_resp = function(xs) {
    1 + 2 * xs + rnorm(length(xs), 0, 1)
  },
  Nsim = 1000,
  l_quantile_display = 0.01,
  u_quantile_display = 0.99,
  error_est = function(est) {
    quantile(est, 0.99) - quantile(est, 0.01)
  },
  num_digits_round = 3,
  draw_theta_at = NULL,
  xlab_names = NULL,
  ...
)
```

## Arguments

xmin	The minimum value of the independent variable.
xmax	The maximum value of the independent variable.
designs	A $d \times n$ matrix where each of the $d$ rows is a design (the $x$ values used to run the experiment).
gen_resp	A model for the response which takes the design as its parameter.
Nsim	The number of estimates per design. Default is 1000.
l_quantile_display	The lowest quantile of the simulation estimates displayed. Default is 0.025.
u_quantile_display	The highest quantile of the simulation estimates displayed. Default is 0.975.
error_est	The error metric for the estimates. The sample standard deviation (i.e. sd) is unstable at low sample sizes. The default is the 90 percentile minus the 10 percentile.

num_digits_round	The number of digits to round the error results. Default is 2.
draw_theta_at	If the user wishes to draw a horizontal line marking theta (to checked biasedness) it is specified here. The default is NULL with no line being drawn.
xlab_names	Text for the x-grid labels. This vector's size should equal length(designs).
...	Additional arguments passed to the boxplot function.

**Value**

A list with the simulated estimates and error estimates for each design.

**Author(s)**

Adam Kapelner

**Examples**

```
xmin = 5 / 15
xmax = 19 / 1
n = 10 #must be even for this demo
designs = rbind(
  c(rep(xmin, n / 2), rep(xmax, n / 2)),      #design A
  seq(from = xmin, to = xmax, length.out = n) #design B
)
design_bakeoff_info = design_bakeoff(xmin, xmax, designs) #design A wins
```

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err\_vs\_theta0\_plot\_for\_homo\_design

*Plots a standard error estimate of thetihat (slope over intercept) over a range of possible theta0 values in order to investigate robustness of the the initial theta0 guess.*

---

**Description**

Plots a standard error estimate of thetihat (slope over intercept) over a range of possible theta0 values in order to investigate robustness of the the initial theta0 guess.

**Usage**

```
err_vs_theta0_plot_for_homo_design(
  n,
  xmin,
  xmax,
  theta,
  theta0_min,
  theta0_max,
  theta0 = NULL,
```

```

beta0 = 1,
sigma = 1,
RES = 500,
Nsim = 5000,
error_est = function(est) {
  quantile(est, 0.99) - quantile(est, 0.01)
},
theta_logged = TRUE,
error_pct = TRUE,
plot_rhos = FALSE,
...
)

```

### Arguments

<code>n</code>	The number of experimental runs.
<code>xmin</code>	The minimum value of the independent variable.
<code>xmax</code>	The maximum value of the independent variable.
<code>theta</code>	The putative true value. This is used to see how much efficiency given up by designing it for $\theta_0$ .
<code>theta0_min</code>	Simulating over different guesses of $\theta_0$ , this is the minimum guess.
<code>theta0_max</code>	Simulating over different guesses of $\theta_0$ , this is the maximum guess.
<code>theta0</code>	The guess used to construct the experimental design. Specify only if you wish to see this value plotted. Default is NULL.
<code>beta0</code>	A guess to be used for the intercept. Defaults to 1.
<code>sigma</code>	A guess to be used for the homoskedastic variance of the measurement errors. If known accurately, then the standard errors (i.e. the y-axis on the plot) will be accurate. Otherwise, the standard errors are useful only when compared to each other in a relative sense. Defaults to 1.
<code>RES</code>	The number of points on the x-axis to simulate. Higher numbers will give smoother results. Default is 20.
<code>Nsim</code>	The number of models to be simulated for estimating the standard error at each value on the x-axis. Default is 1000.
<code>error_est</code>	The error metric for the estimates. The sample standard deviation (i.e. <code>sd</code> ) is unstable at low sample sizes. The default is the 90 percentile minus the 10 percentile.
<code>theta_logged</code>	Should the values of $\theta$ be logged? Default is TRUE.
<code>error_pct</code>	Plot error as a percentage increase from minimum. Default is TRUE.
<code>plot_rhos</code>	Plot an additional graph of $\rho$ by $\theta_0$ . Default is FALSE.
<code>...</code>	Additional arguments passed to the <code>plot</code> function.

### Value

A list with original parameters as well as data from the simulation

**Author(s)**

Adam Kapelner

**Examples**

```
xmin = 5 / 15
xmax = 19 / 1
n = 10
theta0 = 0.053
plot_info = err_vs_theta_plot_for_homo_design(
  n, xmin, xmax, theta0, theta0_min = 0.001, theta0_max = 1
)
```

---

experimental\_results *Report the results of the experiment as well as confidence intervals.*

---

**Description**

Report the results of the experiment as well as confidence intervals.

**Usage**

```
experimental_results(xs, ys, alpha = 0.05, B = 1000)
```

**Arguments**

xs	The design
ys	The measurements of the response
alpha	1 - alpha is the confidence of the computed intervals. Default is 0.05.
B	For the confidence interval methods with an embedded bootstrap (or resampling), the number of resamples (defaults to 1000).

**Value**

A list object containing the estimate as well as confidence intervals and parameters.

**Author(s)**

Adam Kapelner

**Examples**

```
n = 10
xmin = 5 / 15
xmax = 19 / 1
xs = runif(n, xmin, xmax)
ys = 2 + 3 * xs + rnorm(n)
experimental_results_info = experimental_results(xs, ys)
```

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naph	<i>This is data for the PRV measurement of the <math>k_H</math> of Napthalene in water. See Section 3 of our paper below for more information.</i>
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**Description**

This is data for the PRV measurement of the  $k_H$  of Napthalene in water. See Section 3 of our paper below for more information.

**Usage**

```
data(naph)
```

**Format**

A data frame with 100 rows and 2 variables

**Author(s)**

Adam Kapelner <kapelner@qc.cuny.edu>

**References**

<https://arxiv.org/abs/1604.03480>

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oed_for_slope_over_intercept	<i>Create an optimal design for measuring the slope divided by the intercept</i>
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**Description**

Create an optimal design for measuring the slope divided by the intercept

**Usage**

```
oed_for_slope_over_intercept(  
  n,  
  xmin,  
  xmax,  
  theta0,  
  f_hetero = NULL,  
  MaxIter = 6000,  
  MaxFunEvals = 6000,  
  TolFun = 1e-06,  
  NUM_RAND_STARTS = 50  
)
```

**Arguments**

<code>n</code>	The number of experimental runs.
<code>xmin</code>	The minimum value of the independent variable.
<code>xmax</code>	The maximum value of the independent variable.
<code>theta0</code>	The guess of the true value of the slope / intercept.
<code>f_hetero</code>	Specification of heteroskedasticity: the $h(x)$ which relates the value of the independent variable to the variance in the response around the line at that place or the proportional variance at that point. If NULL, homoskedasticity is assumed (this is the default behavior).
<code>MaxIter</code>	For the heteroskedastic design, a Nelder-Mead search is used (via the function <code>fminbnd</code> ). This is the <code>MaxIter</code> value for the search. Default is 6000. Lower if <code>n</code> is high.
<code>MaxFunEvals</code>	For the heteroskedastic design, a Nelder-Mead search is used (via the function <code>fminbnd</code> ). This is the <code>MaxFunEvals</code> value for the search. Default is 6000. Lower if <code>n</code> is high.
<code>TolFun</code>	For the heteroskedastic design, a Nelder-Mead search is used (via the function <code>fminbnd</code> ). This is the <code>TolFun</code> value for the search. Default is $1e-6$ . Increase for faster execution.
<code>NUM_RAND_STARTS</code>	For the heteroskedastic design, a Nelder-Mead search is used (via the function <code>fminbnd</code> ). The Nelder-Mead search must be given a starting location. Our implementation uses many starting locations. This parameter controls the number of additional random starting locations in the space $[xmin, xmax]$ . Default is 50.

**Value**

An `n`-vector of `x`-values which specifies the optimal design

**Author(s)**

Adam Kapelner

**Examples**

```
xmin = 5 / 15
xmax = 19 / 1
n = 10
theta0 = 0.053
opt_homo_design = oed_for_slope_over_intercept(n, xmin, xmax, theta0)
table(opt_homo_design)
```

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optDesignSlopeInt

*Optimal Designs for Estimating the Slope Divided by the Intercept*

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**Description**

Software which helps practitioners optimally design experiments that measure the slope divided by the intercept.

**Author(s)**

Adam Kapelner <kapelner@qc.cuny.edu>



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