

Package: CEACT (via r-universe)

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Type Package

Title Cost-Effectiveness Analysis Toolkit for Clinical Trials

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Author Imad EL BADISY [aut, cre]

Maintainer Imad EL BADISY <elbadisyimad@gmail.com>

Description CEACT (Cost-Effectiveness Analysis Toolkit for Clinical Trials) is a comprehensive R package designed to facilitate the economic evaluation of healthcare interventions in clinical trial settings. It provides tools to compute and visualize Incremental Cost-Effectiveness Ratios (ICERs), Cost-Effectiveness Acceptability Curves (CEAC), Cost-Effectiveness Planes, and Net Monetary Benefit (NMB) tables. The package streamlines cost-utility analyses and supports both observed and bootstrap-based approaches.

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boot_icer	<i>Bootstrap Incremental Cost, Effect, ICER, and Net Benefit</i>
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Description

Performs non-parametric stratified bootstrap resampling for a two-arm trial-based economic evaluation. Resampling is stratified by trial group to preserve arm sizes.

Usage

```
boot_icer(formula, data, ref, R = 1000, ci.type = "bca", na.omit = TRUE)
```

Arguments

formula	A formula of the form <code>cost + effect ~ group</code> .
data	A data frame containing the variables in the formula.
ref	Reference group label.
R	Number of bootstrap replications.
ci.type	Confidence interval type passed to <code>boot::boot.ci()</code> .
na.omit	Logical; whether to remove rows with missing values.

Value

An object of class `boot_icer` containing a summary table, bootstrap distribution, observed estimates, formula, reference group, and matched call.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
res <- boot_icer(cost + effect ~ group, data = df, ref = "control", R = 200)
summary(res)
```

 cea

Cost-Effectiveness Analysis Summary for a Two-Arm Trial

Description

Computes observed cost-effectiveness summaries comparing one treatment group with one reference group. The incremental cost-effectiveness ratio is $ICER = (\text{mean}(\text{cost_treatment}) - \text{mean}(\text{cost_reference})) / (\text{mean}(\text{effect_treatment}) - \text{mean}(\text{effect_reference}))$.

Usage

```
cea(formula, data, ref, na.omit = TRUE)
```

Arguments

formula	A formula of the form <code>cost + effect ~ group</code> .
data	A data frame containing the variables used in the formula.
ref	Character string specifying the reference group.
na.omit	Logical; whether to remove rows with missing values.

Value

An object of class `cea`, a data frame with group means, standard deviations, differences, confidence intervals, and p-values. Attributes include `ICER`, `delta_cost`, `delta_effect`, `formula`, `ref`, and `call`.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
res <- cea(cost + effect ~ group, data = df, ref = "control")
summary(res)
```

 compute_nmb_ceac

Compute Net Monetary Benefit and CEAC Table

Description

Computes expected incremental net monetary benefit (INMB) and the probability that treatment is cost-effective across willingness-to-pay thresholds. For a two-arm trial, $INMB(k) = k * \text{DeltaEffect} - \text{DeltaCost}$.

Usage

```
compute_nmb_ceac(
  x,
  data = NULL,
  wtp_range = seq(0, 1e+05, 1000),
  ref = NULL,
  R = 1000,
  ...
)
```

Arguments

x	Either a <code>boot_icer</code> object or a formula of the form <code>cost + effect ~ group</code> .
data	Data frame required when x is a formula.
wtp_range	Numeric vector of willingness-to-pay thresholds.
ref	Reference group label required when x is a formula.
R	Number of bootstrap replications used when x is a formula.
...	Additional arguments passed to <code>boot_icer()</code> for formula input.

Value

An object of class "nmb_ceac" with columns WTP, ENMB, and Prob_CE.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
ceac_tbl <- compute_nmb_ceac(cost + effect ~ group, data = df,
                             ref = "control", R = 200,
                             wtp_range = seq(0, 50000, 5000))
head(ceac_tbl)
```

Description

Varies one treatment-arm parameter while holding observed reference-arm data fixed, then recomputes the ICER or incremental net monetary benefit.

Usage

```
dsa_icer(formula, data, param, range, ref, metric = "ICER", k = 1000)
```

Arguments

formula	A formula of the form <code>cost + effect ~ group</code> .
data	A data frame.
param	Name of the variable to vary, usually the cost or effect variable from formula.
range	Numeric vector of values assigned to the treatment arm.
ref	Reference group label.
metric	Either "ICER" or "INMB". "NMB" is accepted as an alias for "INMB".
k	Willingness-to-pay threshold used for INMB.

Value

A data frame with varied parameter values and resulting metric.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
dsa <- dsa_icer(cost + effect ~ group, data = df, param = "effect",
               range = seq(0.74, 0.82, 0.02), ref = "control",
               metric = "INMB", k = 20000)
head(dsa)
```

plot_ceac

Plot Cost-Effectiveness Acceptability Curve

Description

Plots the probability that treatment is cost-effective across willingness-to-pay thresholds.

Usage

```
plot_ceac(
  x,
  data = NULL,
  wtp_range = seq(0, 1e+05, 1000),
  ref = NULL,
  R = 1000,
  ...
)
```

Arguments

x	A <code>boot_icer</code> object, <code>nmb_ceac</code> object, or formula of the form <code>cost + effect ~ group</code> .
data	Data frame required when x is a formula.
wtp_range	Numeric vector of willingness-to-pay thresholds.

ref	Reference group label required when x is a formula.
R	Number of bootstrap replications used when x is a formula.
...	Additional arguments passed to <code>ggplot2::labs()</code> .

Value

A ggplot object.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
res <- boot_icer(cost + effect ~ group, data = df, ref = "control", R = 200)
plot_ceac(res, wtp_range = seq(0, 50000, 5000))
```

plot_ceplane *Plot Cost-Effectiveness Plane*

Description

Visualizes bootstrap replicates of incremental cost and incremental effect.

Usage

```
plot_ceplane(boot_icer_result, k = NULL, subtitle = NULL)
```

Arguments

boot_icer_result	A boot_icer object from <code>boot_icer()</code> .
k	Optional willingness-to-pay threshold shown as a straight line with slope k.
subtitle	Optional subtitle text.

Value

A ggplot object.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
res <- boot_icer(cost + effect ~ group, data = df, ref = "control", R = 200)
plot_ceplane(res, k = 20000)
```

plot_tornado	<i>Tornado Plot for Deterministic Sensitivity Analysis</i>
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Description

Creates a line-style sensitivity plot from a data frame generated by `dsa_icer()`.

Usage

```
plot_tornado(df, metric = "ICER")
```

Arguments

df	A data frame from <code>dsa_icer()</code> .
metric	Character string. Either "ICER" or "INMB".

Value

A ggplot object.

Examples

```
df <- simulate_ce_trial(n = 100, seed = 123)
dsa <- dsa_icer(cost + effect ~ group, data = df, param = "effect",
  range = seq(0.74, 0.82, 0.02), ref = "control")
plot_tornado(dsa, metric = "ICER")
```

simulate_ce_trial	<i>Simulate a Two-Arm Trial for Cost-Effectiveness Analysis</i>
-------------------	---

Description

Generates individual-level cost and effect outcomes for a simple two-arm randomized clinical trial. The function is intended for examples, tutorials, tests, and manuscript demonstrations.

Usage

```
simulate_ce_trial(
  n = 200,
  mean_cost = c(5000, 5600),
  sd_cost = c(900, 1000),
  mean_effect = c(0.72, 0.78),
  sd_effect = c(0.1, 0.11),
  rho = 0.15,
  seed = NULL,
  group_names = c("control", "treatment")
)
```

Arguments

n	Number of participants per arm.
mean_cost	Control and treatment mean costs.
sd_cost	Control and treatment cost standard deviations.
mean_effect	Control and treatment mean effects, for example QALYs.
sd_effect	Control and treatment effect standard deviations.
rho	Within-person cost-effect correlation used in a Gaussian construction.
seed	Optional random seed.
group_names	Character vector naming control and treatment arms.

Value

A data frame with cost, effect, and group.

Examples

```
trial <- simulate_ce_trial(n = 50, seed = 1)
head(trial)
```

trial_cea	<i>Example Trial-Based Cost-Effectiveness Dataset</i>
-----------	---

Description

A patient-level clinical-trial cost-effectiveness dataset with treatment assignment, total cost, QALYs, and baseline covariates. The dataset is used in teaching examples for trial-based economic evaluation and was described as being provided by the Health Services Research Unit for the textbook *Economic Evaluation in Clinical Trials*.

Usage

```
trial_cea
```

Format

A data frame with 500 rows and 10 variables:

- id** Patient identifier.
- treat** Treatment indicator, 1 for treatment and 0 for control.
- cost** Total cost in US dollars.
- qaly** Quality-adjusted life-years.
- dissev** Disease severity, ranging approximately from 0.025 to 0.729.
- race** Race indicator as supplied in the source dataset.
- bcost** Baseline cost in US dollars.
- blqaly** Baseline QALY.
- male** Sex indicator, 1 for male and 0 for female.
- group** Factor version of treat, with levels control and treatment.

Source

Health Services Research Unit example dataset used in Glick HA, Doshi JA, Sonnad SS, Polsky D. *Economic Evaluation in Clinical Trials*. Oxford University Press. Also distributed in the ceaR package as clintrial_cea.

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