

Package ‘decisionSupport’

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Type Package

Title Quantitative Support of Decision Making under Uncertainty

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Description Supporting the quantitative analysis of binary welfare based decision making processes using Monte Carlo simulations. Decision support is given on two levels: (i) The actual decision level is to choose between two alternatives under probabilistic uncertainty. This package calculates the optimal decision based on maximizing expected welfare. (ii) The meta decision level is to allocate resources to reduce the uncertainty in the underlying decision problem, i.e to increase the current information to improve the actual decision making process. This problem is dealt with using the Value of Information Analysis (VIA). The Expected Value of Information (EVI) for arbitrary prospective estimates can be calculated as well as Individual and Clustered Expected Value of Perfect Information (EVPI). The probabilistic calculations are done via Monte Carlo simulations. This Monte Carlo functionality can be used on its own.

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 'rdistq_fit.R'
 'rdist90ci_exact.R'
 'paramtnormci.R'
 'r0_1norm90ci_numeric.R'
 'random_estimate_1d.R'
 'random.R'
 'estimate.R'
 'mcSimulation.R'
 'welfareDecisionAnalysis.R'
 'eviSimulation.R'
 'individualEvpiSimulation.R'
 'estimate_read_csv_old.R'
 'uncertaintyAnalysis.R'
 'decisionSupport-package.R'
 'paramtnormci_fit.R'
 'paramtnormci_numeric.R'
 'rposnorm90ci_iter.R'

R topics documented:

as.data.frame.mcSimulation	3
corMat	4
corMat.estimate	4
decisionSupport	5
estimate	7
estimate_read_csv	7
estimate_read_csv_old	9
estimate_write_csv	10
eviSimulation	11
hist.mcSimulation	15
individualEvpiSimulation	16
mcSimulation	18
names.estimate	21
paramposnorm90ci	21
paramtnormci	22
paramtnormci_fit	23
paramtnormci_numeric	24
print.mcSimulation	25
print.summary.eviSimulation	26
print.summary.mcSimulation	26
print.summary.welfareDecisionAnalysis	27
r0_1norm90ci_numeric	27
random	28
random.default	28
random.estimate	29

random_estimate_ld	30
rdist90ci_exact	31
rdistq_fit	31
rmvnorm90ci_exact	33
row.names.estimate	33
rposnorm90ci_iter	34
rposnorm90ci_numeric	34
sort.summary.eviSimulation	35
summary.eviSimulation	36
summary.mcSimulation	36
summary.welfareDecisionAnalysis	37
uncertaintyAnalysis	38
welfareDecisionAnalysis	39

Index	42
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as.data.frame.mcSimulation
Coerce to a Data Frame.

Description

Functions to check if an object is a data frame, or coerce it if possible.

Usage

```
## S3 method for class 'mcSimulation'
as.data.frame(x, row.names = NULL, optional = FALSE,
  ..., stringsAsFactors = default.stringsAsFactors())
```

Arguments

x	An object of class mcSimulation.
row.names	NULL or a character vector giving the row names for the data frame. Missing values are not allowed.
optional	logical. If TRUE, setting row names and converting column names (to syntactic names: see make.names) is optional.
...	additional arguments to be passed to or from methods.
stringsAsFactors	logical: should the character vector be converted to a factor?

See Also

[as.data.frame](#)

corMat	<i>Return the Correlation Matrix of x.</i>
--------	--

Description

Return the correlation matrix of x.

Usage

```
corMat(rho)
```

Arguments

rho a distribution.

corMat.estimate	<i>Return the correlation matrix of an estimate object.</i>
-----------------	---

Description

This function returns the full correlation matrix of an [estimate](#) object.

Usage

```
## S3 method for class 'estimate'  
corMat(rho)
```

Arguments

rho an [estimate](#) object.

See Also

[estimate](#), [row.names.estimate](#), [names.estimate](#)

Description

The **decisionSupport** package supports the quantitative analysis of welfare based decision making processes using Monte Carlo simulations. This is an important part of the Applied Information Economics (AIE) approach developed in Hubbard (2014). These decision making processes can be categorized into two levels of decision making:

1. The actual problem of interest of a policy maker which we call the *underlying welfare based decision* on how to influence an ecological-economic system based on a particular information on the system available to the decision maker and
2. the *meta decision* on how to allocate resources to reduce the uncertainty in the underlying decision problem, i.e to increase the current information to improve the underlying decision making process.

The first problem, i.e. the underlying problem, is the problem of choosing the decision which maximizes expected welfare. The welfare function can be interpreted as a von Neumann-Morgenstern utility function. Whereas, the second problem, i.e. the meta decision problem, is dealt with using the *Value of Information Analysis (VIA)*. Value of Information Analysis seeks to assign a value to a certain reduction in uncertainty or, equivalently, increase in information. Uncertainty is dealt with in a probabilistic manner. Probabilities are transformed via Monte Carlo simulations.

Details

The functionality of this package is subdivided into three main parts: (i) the welfare based analysis of the underlying decision, (ii) the meta decision of reducing uncertainty and (iii) the Monte Carlo simulation for the transformation of probabilities and calculation of expectation values. Furthermore, there is a wrapper function around these three parts which aims at providing an easy-to-use interface.

Welfare based Analysis of the Underlying Decision Problem:

Welfare Decision Analysis: Implementation: [welfareDecisionAnalysis](#)

Utility Functions: Implementation: `ToDo`

The Meta Decision of Reducing Uncertainty: The meta decision of how to allocate resources for uncertainty reduction can be analyzed with this package in two different ways: via (i) Expected Value of Information Analysis or (ii) via Partial Least Squares (PLS) analysis and Variable Importance in Projection (VIP).

Expected Value of Information (EVI): Implementation: [eviSimulation](#), [individualEvpSimulation](#)

Partial Least Squares (PLS) analysis and Variable Importance in Projection (VIP): Implementation: `ToDo`

Solving the Practical Problem of Calculating Expectation Values by Monte Carlo Simulation:

Estimates: Implementation: [estimate](#)

Multivariate Random Number Generation: Implementation: [random.estimate](#)

Monte Carlo Simulation: Implementation: [mcSimulation](#)

Uncertainty Analysis: A wrapper function: Implementation: [uncertaintyAnalysis](#)

Package Options

ToDo

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References

Hubbard, Douglas W., *How to Measure Anything? - Finding the Value of "Intangibles" in Business*, John Wiley & Sons, Hoboken, New Jersey, 2014, 3rd Ed, <http://www.howtomeasureanything.com/>.

Hugh Gravelle and Ray Rees, *Microeconomics*, Pearson Education Limited, 3rd edition, 2004.

See Also

[welfareDecisionAnalysis](#), [eviSimulation](#), [mcSimulation](#)

estimate	Create an Estimate Object
----------	---------------------------

Description

This function creates an object of class estimate. #ToDo: detailed description #ToDo: Implement characterization of distribution by mean and sd. Eventually, also by other quantiles.

Usage

```
estimate(..., correlation_matrix = NULL)
```

Arguments

- ... arguments that can be coerced to a data frame comprising the base of the estimate.
- correlation_matrix numeric matrix containing the correlations of the variables.

Details

The parameters in ... provide the base information of an estimate.

The structure of the estimate base information (mandatory): Mandatory columns:

Column name	R-type	Explanation
distribution	character	Distribution types
variable	character	Variable names

Value

An object of type estimate which is a list whith components base and correlation_matrix. base is a data.frame with mandatory column distribution. The row.names are the names of the variables. correlation_matrix is a symmetric matrix with row and column names being the subset of the variables supplied in base which are correlated. Its elements are the corresponding correlations.

See Also

row.names.estimate, names.estimate, corMat, estimate_read_csv, estimate_write_csv, random.estimate

estimate_read_csv	Read an Estimate from CSV - File.
-------------------	-----------------------------------

Description

This function reads an [estimate](#) from the specified csv files. In this context, an estimate of a variable is defined by its distribution type, its 90%-confidence interval [lower, upper] and its correlation to other variables. #ToDo: Implement characterization of distribution by mean and sd. Eventually, also by other quantiles.

Usage

```
estimate_read_csv(fileName, strip.white = TRUE, ...)
```

Arguments

fileName	Name of the file containing the base information of the estimate that should be read.
strip.white	logical. Allows the stripping of leading and trailing white space from unquoted character fields (numeric fields are always stripped). See scan for further details (including the exact meaning of 'white space'), remembering that the columns may include the row names.
...	Further parameters to be passed to read.csv .

Details

An estimate might consists of uncorrelated and correlated variables. This is reflected in the input file structure, which is described in the following.

Value

An object of type [estimate](#).

CSV input file structures

The estimate is read from one or two csv files: the basic csv file which is mandatory and the correlation csv file which is optional. The basic csv file contains the definition of the distribution of all variables ignoring potential correlations. The correlation csv file only defines correlations.

The structure of the basic input file (mandatory): File name structure: <basic-filename>.csv
Mandatory columns:

Column name	R-type	Explanation
lower	numeric	ToDo
upper	numeric	ToDo
distribution	character	ToDo
variable	character	ToDo

Optional columns:

Column name	R-type	Explanation
description	character	ToDo
median	numeric	ToDo

start	integer	ToDo
end	integer	ToDo
indicator	logical	ToDo

Columns without names are ignored. Rows where the variable field is empty are also dropped.

The structure of the correlation file (optional): File name structure: <basic-filename>_cor.csv
Columns and rows are named by the corresponding variables. Only those variables need to be present which are correlated with others. The element ["rowname", "columnname"] contains the correlation between the variables rowname and columnname. Uncorrelated elements can be left empty, i.e. as NA, or defined as 0. The element ["name", "name"] has to be set to 1. The matrix must be given in symmetric form.

See Also

[estimate_write_csv](#), [read.csv](#), [estimate](#)

estimate_read_csv_old	<i>Read an Estimate from CSV - File (depreciated standard).</i>
-----------------------	---

Description

This function reads an [estimate](#) from the specified csv files. In this context, an estimate of a variable is defined by its distribution type, its 90%-confidence interval [lower, upper] and its correlation to other variables. #ToDo: Implement characterization of distribution by mean and sd. Eventually, also by other quantiles.

Usage

```
estimate_read_csv_old(fileName, strip.white = TRUE, ...)
```

Arguments

fileName	Name of the file containing the base information of the estimate that should be read.
strip.white	logical. Allows the stripping of leading and trailing white space from unquoted character fields (numeric fields are always stripped). See scan for further details (including the exact meaning of 'white space'), remembering that the columns may include the row names.
...	Further parameters to be passed to read.csv .

Details

An estimate might consists of uncorrelated and correlated variables. This is reflected in the input file structure, which is described in the following.

Value

An object of type `estimate`.

CSV input file structures

The estimate is read from one or two csv files: the basic csv file which is mandatory and the correlation csv file which is optional. The basic csv file contains the definition of the distribution of all variables ignoring potential correlations. The correlation csv file only defines correlations.

The structure of the basic input file (mandatory): File name structure: <basic-filename>.csv
Mandatory columns:

Column name	R-type	Explanation
lower	numeric	ToDo
upper	numeric	ToDo
distribution	character	ToDo
variable	character	ToDo

Optional columns:

Column name	R-type	Explanation
description	character	ToDo
median	numeric	ToDo
start	integer	ToDo
end	integer	ToDo
indicator	logical	ToDo

Columns without names are ignored. Rows where the variable field is empty are also dropped.

The structure of the correlation file (optional): File name structure: <basic-filename>.csv_correlations.csv
#ToDo

See Also

`estimate_read_csv`, `read.csv`, `estimate`

<code>estimate_write_csv</code>	<i>Write an Estimate to CSV - File.</i>
---------------------------------	---

Description

This function writes an `estimate` to the specified csv file(s).

Usage

```
estimate_write_csv(estimate, fileName, varNamesAsColumn = TRUE,
  quote = FALSE, ...)
```

Arguments

estimate	Estimate object to write to file fileName.
fileName	character. Output file name which must end with .csv.
varNamesAsColumn	logical; If TRUE the variable names will be written as a separate column, otherwise as row names.
quote	a logical value (TRUE or FALSE) or a numeric vector. If TRUE, any character or factor columns will be surrounded by double quotes. If a numeric vector, its elements are taken as the indices of columns to quote. In both cases, row and column names are quoted if they are written. If FALSE, nothing is quoted. Parameter is passed on to write.csv .
...	Further parameters to be passed to write.csv .

Value

An object of type [estimate](#).

See Also

[estimate_read_csv](#), [estimate](#), [write.csv](#)

eviSimulation	<i>Expected Value of Information (EVI) Simulation.</i>
---------------	--

Description

The Expected Value of Information (EVI) is calculated based on a Monte Carlo simulation of the values of two different decision alternatives.

Usage

```
eviSimulation(model, currentEstimate, prospectiveEstimate, numberOfSimulations,
  functionSyntax = "data.frameNames")
```

Arguments

model	either a function or a list with two functions: <code>list(p1,p2)</code> . In the first case the function is the net benefit of project approval vs. the status quo. In the second case the element p1 is the function valuing the first project and the element p2 valuing the second project.
currentEstimate	estimate object describing the distribution of the input variables as currently estimated.

prospectiveEstimate [estimate](#) object describing the prospective distribution of the input variables which could hypothetically achieved by collecting more information, viz. improving the measurement.

numberOfSimulations integer; number of simulations to be used in the underlying Monte Carlo analysis

functionSyntax function character; function syntax used in the model function(s).

Details

This principle is along the line described in Hubbard (2014). The Expected Value of Information is the decrease in the EOL for an information improvement from the current estimate (I_{current}) to a better prospective (or hypothetical) information ($I_{\text{prospective}}$): $\text{EVI} := \text{EOL}(I_{\text{current}}) - \text{EOL}(I_{\text{prospective}})$. Thus, the EVI depends on the model for valueing a decision, the current information, i.e. the current estimate, and the specification of a hypothetical improvement in information, i.e. a prospective estimate.

Value

An object of class `eviSimulation` with the following elements:

current	welfareDecisionAnalysis object for currentEstimate
prospective	welfareDecisionAnalysis object for prospectiveEstimate
evi	Expected Value of Information (EVI) of gained by the prospective estimate w.r.t. the current estimate

See Also

[welfareDecisionAnalysis](#), [mcSimulation](#), [estimate](#)

Examples

```
#####
# Example 1 Only one prospective estimate:
#####
numberOfSimulations=10000
# Create the estimate object:
variable=c("revenue","costs")
distribution=c("posnorm","posnorm")
lower=c(10000, 5000)
upper=c(100000, 50000)
currentEstimate<-estimate(variable, distribution, lower, upper)
prospectiveEstimate<-currentEstimate
revenueConst<-mean(c(currentEstimate$base["revenue","lower"],
                     currentEstimate$base["revenue","upper"]))
prospectiveEstimate$base["revenue",]<-data.frame(distribution="const",
                                                  lower=revenueConst,
                                                  upper=revenueConst,
                                                  row.names="revenue",
                                                  stringsAsFactors=FALSE)

# (a) Define the model function without name for the return value:
profit<-function(x){
```

```

x$revenue-x$costs
}

# Calculate the Expected Value of Information:
eviSimulationResult<-eviSimulation(model=profit,
                                   currentEstimate=currentEstimate,
                                   prospectiveEstimate=prospectiveEstimate,
                                   numberOfSimulations=numberOfSimulations,
                                   functionSyntax="data.frameNames")

# Show the simulation results:
print(summary(eviSimulationResult))
#####
# (b) Define the model function with a name for the return value:
profit<-function(x){
  list(Profit=x$revenue-x$costs)
}
# Calculate the Expected Value of Information:
eviSimulationResult<-eviSimulation(model=profit,
                                   currentEstimate=currentEstimate,
                                   prospectiveEstimate=prospectiveEstimate,
                                   numberOfSimulations=numberOfSimulations,
                                   functionSyntax="data.frameNames")

# Show the simulation results:
print(summary((eviSimulationResult)))
#####
# (c) Two decision variables:
decisionModel<-function(x){
  list(Profit=x$revenue-x$costs,
       Costs=-x$costs)
}
# Calculate the Expected Value of Information:
eviSimulationResult<-eviSimulation(model=decisionModel,
                                   currentEstimate=currentEstimate,
                                   prospectiveEstimate=prospectiveEstimate,
                                   numberOfSimulations=numberOfSimulations,
                                   functionSyntax="data.frameNames")

# Show the simulation results:
print(summary((eviSimulationResult)))
#####
# Example 2 A list of prospective estimates:
#####
numberOfSimulations=10000
# Define the model function with a name for the return value:
profit<-function(x){
  list(Profit=x$revenue-x$costs)
}
# Create the estimate object:
variable=c("revenue","costs")
distribution=c("posnorm","posnorm")
lower=c(10000, 5000)
upper=c(100000, 50000)
currentEstimate<-estimate(variable, distribution, lower, upper)
perfectInformationRevenue<-currentEstimate

```

[illegible]

```

upper=revenueConst,
row.names="revenue",
stringsAsFactors=FALSE)

perfectInformationCosts<-currentEstimate
costsConst<-mean(c(currentEstimate$base["costs", "lower"], currentEstimate$base["costs", "upper"]))
perfectInformationCosts$base["costs", ]<-data.frame(distribution="const",
lower=costsConst,
upper=costsConst,
row.names="costs",
stringsAsFactors=FALSE)

prospectiveEstimate<-list(perfectInformationRevenue=perfectInformationRevenue,
perfectInformationCosts=perfectInformationCosts)

# Define the model function with two decision variables:
decisionModel<-function(x){
  list(Profit=x$revenue-x$costs,
Costs=-x$costs)
}

# Calculate the Expected Value of Information:
eviSimulationResult<-eviSimulation(model=decisionModel,
currentEstimate=currentEstimate,
prospectiveEstimate=prospectiveEstimate,
numberOfSimulations=numberOfSimulations,
functionSyntax="data.frameNames")

# Show the simulation results:
print(sort(summary(eviSimulationResult)),decreasing=TRUE,along="Profit")

```

hist.mcSimulation

*Plot Histogram of results of a Monte Carlo Simulation***Description**

This function plots the histograms of the results of [mcSimulation](#).

Usage

```

## S3 method for class 'mcSimulation'
hist(x, breaks = 100, col = NULL, xlab = NULL,
main = paste("Histogram of ", xlab), ..., colorQuantile = c("GREY",
"YELLOW", "ORANGE", "DARK GREEN", "ORANGE", "YELLOW", "GREY"),
colorProbability = c(1, 0.95, 0.75, 0.55, 0.45, 0.25, 0.05),
resultName = NULL)

```

Arguments

x	An object of class mcSimulation.
breaks	one of: <ul style="list-style-type: none"> a vector giving the breakpoints between histogram cells, a function to compute the vector of breakpoints,

- a single number giving the number of cells for the histogram,
- a character string naming an algorithm to compute the number of cells (see ‘Details’),
- a function to compute the number of cells.

In the last three cases the number is a suggestion only; the breakpoints will be set to [pretty](#) values. If breaks is a function, the x vector is supplied to it as the only argument.

col	a colour to be used to fill the bars. The default of NULL yields unfilled bars.
xlab	character; x label of the histogram. If it is not provided, i.e. equals NULL the name of the chosen variable by argument resultName is used.
main	character; main title of the histogram.
...	Further arguments to be passed to hist .
colorQuantile	character vector encoding the color of the quantiles defined in argument colorProbability.
colorProbability	numeric vector; defines the quantiles that shall be distinguished by the colors chosen in argument colorQuantile. Must be of the same length as colorQuantile.
resultName	character; indicating the name of the component of the simulation function (model_function) which results histogram shall be generated. If model_function is single valued, no name needs to be supplied. Otherwise, one valid name has to be specified. Defaults to NULL.

Value

an object of class "histogram". For details see [hist](#).

See Also

[mcSimulation](#), [hist](#). For a list of colors available in R see [colors](#).

individualEvpSimulation

Individual Expected Value of Perfect Information Simulation

Description

The Individual Expected Value of Perfect Information (Individual EVPI) is calculated based on a Monte Carlo simulation of the values of two different decision alternatives.

Usage

```
individualEvpSimulation(model, currentEstimate,
  perfectProspectiveNames = row.names(currentEstimate),
  perfectProspectiveValues = colMeans(random(rho = currentEstimate, n =
    numberOfSimulations)[, perfectProspectiveNames]), numberOfSimulations,
  functionSyntax = "data.frameNames")
```


Arguments

model	either a function or a list with two functions: <code>list(p1,p2)</code> . In the first case the function is the net benefit of project approval vs. the status quo. In the second case the element p1 is the function valuing the first project and the element p2 valuing the second project.
currentEstimate	estimate object describing the distribution of the input variables as currently estimated.
perfectProspectiveNames	character vector; input variable names that are assumed to be known perfectly with prospective information.
perfectProspectiveValues	numeric vector of the same length as <code>perfectProspectiveNames</code> with the corresponding values assumed to be known perfectly.
numberOfSimulations	integer; number of simulations to be used in the underlying Monte Carlo analysis
functionSyntax	function character; function syntax used in the model function(s).

Details

This principle is along the line described in Hubbard (2014). The Expected Value of Information is the decrease in the EOL for an information improvement from the current estimate (I_{current}) to a better prospective (or hypothetical) information ($I_{\text{prospective}}$): $\text{EVI} := \text{EOL}(I_{\text{current}}) - \text{EOL}(I_{\text{prospective}})$. If one variables under $I_{\text{prospective}}$ is assumed to be known with certainty the EVI is called the Individual Expected Value of Perfect Information (Individual EVPI). More precisely, if one assumes under $I_{\text{prospective}}$ to perfectly know (x_1, \dots, x_k) to equal (a_1, \dots, a_k) then one can specify the notation as Individual EVPI[$x_i = a_i$]. Summarizing, the Individual EVPI depends on the model for valuing a decision, the current information, i.e. the current estimate, and the specification of the variable that is assumed to be known with certainty, viz. the improvement in information, i.e. a prospective estimate.

Value

An object of class `eviSimulation` with the following elements:

current	welfareDecisionAnalysis object for currentEstimate
prospective	welfareDecisionAnalysis object for prospectiveEstimate
evi	Expected Value of Information (EVI) of gained by the prospective estimate w.r.t. the current estimate

See Also

[eviSimulation](#), [welfareDecisionAnalysis](#), [mcSimulation](#), [estimate](#)

Examples

```
# Number of simulations:
n=100000
# Create the current estimate from text:
```

```

estimateText<-"variable, distribution, lower, upper
      revenue1, posnorm,      100,   1000
      revenue2, posnorm,      50,   2000
      costs1,   posnorm,      50,   2000
      costs2,   posnorm,     100,   1000"
currentEstimate<-estimate(read.csv(header=TRUE, text=estimateText,
      strip.white=TRUE, stringsAsFactors=FALSE))

# The model function:
profitModel <- function(x){
  list(Profit=x$revenue1 + x$revenue2 - x$costs1 - x$costs2)
}
# Calculate the Individual EVPI:
individualEvpiResult<-individualEvpiSimulation(model=profitModel,
      currentEstimate=currentEstimate,
      numberOfSimulations=n,
      functionSyntax="data.frameNames")

# Show the simulation results:
print(sort(summary(individualEvpiResult)),decreasing=TRUE,along="Profit")

```

mcSimulation

Perform a Monte Carlo Simulation.

Description

This method solves the following problem. Given a multivariate random variable $x = (x_1, \dots, x_k)$ with joint probability distribution P , i.e.

$$x \sim P.$$

Then the continuous function

$$f : R^k \rightarrow R^l, y = f(x)$$

defines another random variable with distribution

$$y \sim f(P).$$

Given a probability density ρ of x that defines P the problem is the determination of the probability density ϕ that defines $f(P)$. This method samples the probability density ϕ of y by Monte Carlo simulation.

Usage

```

mcSimulation(estimate, model_function, ..., numberOfSimulations,
  randomMethod = "calculate", functionSyntax = "data.frameNames")

```

Arguments

estimate Filename or estimate object representing the joint probability distribution of the input variables.

model_function A numeric function; The function that describes the value of a certain project.


```

                                functionSyntax="data.frameNames")

# Show the simulation results:
print(summary(predictionProfit1))
hist(predictionProfit1,xlab="Profit")
#####
# (b) Define the model function with a name for the return value:
profit1<-function(x){
  list(Profit=x$revenue-x$costs)
}
# Perform the Monte Carlo simulation:
predictionProfit1<-mcSimulation( estimate=costBenefitEstimate,
                                model_function=profit1,
                                numberOfSimulations=100000,
                                functionSyntax="data.frameNames")

# Show the simulation results:
print(summary(predictionProfit1, classicView=TRUE))
hist(predictionProfit1)
#####
# (c) Using global names in the model function syntax
# (CAVE: currently slow!):
profit1<-function(){
  list(Profit=revenue-costs)
}
# Perform the Monte Carlo simulation:
predictionProfit1<-mcSimulation( estimate=costBenefitEstimate,
                                model_function=profit1,
                                numberOfSimulations=10000,
                                functionSyntax="globalNames")

# Show the simulation results:
print(summary(predictionProfit1, probs=c(0.05,0.50,0.95)))
hist(predictionProfit1)

#####
# Example 2(Reading the estimate from file):
#####
# Define the model function:
profit2<-function(x){
  Profit<-x[["sales"]]*(x[["productprice"]] - x[["costprice"]])
  list(Profit=Profit)
}
# Read the estimate of sales, productprice and costprice from file:
inputFileName=system.file("extdata","profit-4.csv",package="decisionSupport")
parameterEstimate<-estimate_read_csv(fileName=inputFileName)
print(parameterEstimate)
# Perform the Monte Carlo simulation:
predictionProfit2<-mcSimulation( estimate=parameterEstimate,
                                model_function=profit2,
                                numberOfSimulations=100000,
                                functionSyntax="data.frameNames")

# Show the simulation results:
print(summary(predictionProfit2))
hist(predictionProfit2)

```

names.estimate	<i>Return the column names of an estimate object.</i>
----------------	---

Description

This function returns the column names of an [estimate](#) object which is identical to `names(x$base)`.

Usage

```
## S3 method for class 'estimate'
names(x)
```

Arguments

`x` an [estimate](#) object.

See Also

[estimate](#), [row.names.estimate](#), [corMat.estimate](#)

paramposnorm90ci	<i>Return parameters of positive normal random distribution based on the 90%-confidence interval.</i>
------------------	---

Description

This function calculates the distribution parameters from the 90%-confidence interval.

Usage

```
paramposnorm90ci(lower, upper, relativeTolerance = 0.05, method = "numeric")
```

Arguments

<code>lower</code>	numeric; lower bound of the 90% confidence intervall.
<code>upper</code>	numeric; upper bound of the 90% confidence intervall.
<code>relativeTolerance</code>	numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval.
<code>method</code>	The method to calculate the parameters. Default is "numeric".

Details

#ToDo

paramtnormci	<i>Return parameters of truncated normal distribution based on a confidence interval.</i>
--------------	---

Description

This function calculates the distribution parameters, i.e. mean and sd, of a truncated normal distribution from an arbitrary confidence interval.

Usage

```
paramtnormci(p, ci, lowerTrunc = -Inf, upperTrunc = Inf,  
  relativeTolerance = 0.05, method = "numeric")
```

Arguments

p	numeric 2-dimensional vector; probabilities of upper and lower bound of the corresponding confidence interval.
ci	numeric 2-dimensional vector; lower and upper bound of the confidence interval.
lowerTrunc	numeric; lower truncation point of the distribution ($\geq -\text{Inf}$).
upperTrunc	numeric; upper truncation point of the distribution ($\leq \text{Inf}$).
relativeTolerance	numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval. If this deviation is greater than <code>relativeTolerance</code> a warning is given.
method	The method to calculate the parameters. Default is "numeric".

Details

For details of the truncated normal distribution see [tnorm](#).

Value

A list with elements mean and sd.

Warning

This method has not been tested systematically!

See Also

[tnorm](#)

paramtnormci_fit	<i>Fit parameters of truncated normal distribution based on a confidence interval.</i>
------------------	--

Description

This function fits the distribution parameters, i.e. mean and sd, of a truncated normal distribution from an arbitrary confidence interval and, facultatively, the median.

Usage

```
paramtnormci_fit(p, ci, median = mean(ci), lowerTrunc = -Inf,
  upperTrunc = Inf, relativeTolerance = 0.05, fitMethod = "Nelder-Mead",
  ...)
```

Arguments

p	numeric 2-dimensional vector; probabilities of upper and lower bound of the corresponding confidence interval.
ci	numeric 2-dimensional vector; lower, i.e ci[[1]], and upper bound, i.e ci[[2]], of the confidence interval.
median	if NULL: truncated normal is fitted only to lower and upper value of the confidence interval; if numeric: truncated normal is fitted on the confidence interval and the median simultaneously. For details cf. below.
lowerTrunc	numeric; lower truncation point of the distribution ($\geq -\text{Inf}$).
upperTrunc	numeric; upper truncation point of the distribution ($\leq \text{Inf}$).
relativeTolerance	numeric; the relative tolerance level of deviation of the generated probability levels from the specified confidence interval. If the relative deviation is greater than relativeTolerance a warning is given.
fitMethod	optimization method used in constrOptim .
...	further parameters to be passed to constrOptim .

Details

For details of the truncated normal distribution see [tnorm](#).

The cummulative distribution of a truncated normal $F_{\mu,\sigma}(x)$ gives the probability that a sampled value is less than x . This is equivalent to saying that for the vector of quantiles $q = (q_{p_1}, \dots, q_{p_k})$ at the corresponding probabilities $p = (p_1, \dots, p_k)$ it holds that

$$p_i = F_{\mu,\sigma}(q_{p_i}), \quad i = 1, \dots, k$$

In the case of arbitrary postulated quantiles this system of equations might not have a solution in μ and σ . A least squares fit leads to an approximate solution:

$$\sum_{i=1}^k (p_i - F_{\mu,\sigma}(q_{p_i}))^2 = \min$$

defines the parameters μ and σ of the underlying normal distribution. This method solves this minimization problem for two cases:

1. $ci[[1]] < median < ci[[2]]$: The parameters are fitted on the lower and upper value of the confidence interval and the median, formally:
 $p_1 = p[[1]]$, $p_2 = 0.5$ and $p_3 = p[[2]]$;
 $q_{p_1} = ci[[1]]$, $q_{0.5} = median$ and $q_{p_3} = ci[[2]]$
2. $median = NULL$: The parameters are fitted on the lower and upper value of the confidence interval only, formally:
 $p_1 = p[[1]]$, $p_2 = p[[2]]$;
 $q_{p_1} = ci[[1]]$, $q_{p_2} = ci[[2]]$

The $(p[[2]] - p[[1]])$ - confidence interval must be symmetric in the sense that $p[[1]] + p[[2]] = 1$.

Value

A list with elements mean and sd.

Warning

This method has not been tested systematically!

See Also

[tnorm](#), [constrOptim](#)

`paramtnormci_numeric` *Return parameters of truncated normal distribution based on a confidence interval.*

Description

This function calculates the distribution parameters, i.e. mean and sd, of a truncated normal distribution from an arbitrary confidence interval.

Usage

```
paramtnormci_numeric(p, ci, lowerTrunc = -Inf, upperTrunc = Inf,
  relativeTolerance = 0.05)
```

Arguments

<code>p</code>	numeric 2-dimensional vector; probabilities of upper and lower bound of the corresponding confidence interval.
<code>ci</code>	numeric 2-dimensional vector; lower and upper bound of the confidence interval.
<code>lowerTrunc</code>	numeric; lower truncation point of the distribution ($\geq -\text{Inf}$).

upperTrunc numeric; upper truncation point of the distribution ($\leq \text{Inf}$).
relativeTolerance numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval. If this deviation is greater than relativeTolerance a warning is given.

Details

For details of the truncated normal distribution see [tnorm](#).

Value

A list with elements mean and sd.

Warning

This method has not been tested systematically!

See Also

[tnorm](#)

print.mcSimulation	<i>Print Basic Results from Monte Carlo Simulation.</i>
--------------------	---

Description

This function prints basic results from Monte Carlo simulation and returns it invisible.

Usage

```
## S3 method for class 'mcSimulation'
print(x, ...)
```

Arguments

x An object of class mcSimulation.
... Further arguments #ToDo

See Also

[mcSimulation](#)

```
print.summary.eviSimulation
```

Print the Summarized EVI Simulation Results.

Description

This function prints the summary of of eviSimulation obtained by [summary.eviSimulation](#).

Usage

```
## S3 method for class 'summary.eviSimulation'  
print(x, ...)
```

Arguments

x	An object of class summary.eviSimulation.
...	Further arguments #ToDo

See Also

[eviSimulation](#)

```
print.summary.mcSimulation
```

Print the Summary of a Monte Carlo Simulation.

Description

This function prints the summary of of mcSimulation obtained by [summary.mcSimulation](#).

Usage

```
## S3 method for class 'summary.mcSimulation'  
print(x, ...)
```

Arguments

x	An object of class mcSimulation.
...	Further arguments #ToDo

See Also

[mcSimulation](#)

```
print.summary.welfareDecisionAnalysis
      Print the Summarized Decsion Analysis Results..
```

Description

This function prints the summary of of welfareDecisionAnalysis obtained by [summary.welfareDecisionAnalysis](#).

Usage

```
## S3 method for class 'summary.welfareDecisionAnalysis'
print(x, ...)
```

Arguments

x	An object of class summary.welfareDecisionAnalysis.
...	Further arguments #ToDo

See Also

[welfareDecisionAnalysis](#)

```
r0_1norm90ci_numeric  Generate normal random numbers truncated to [0,1] based on the
                        90%-confidence interval.
```

Description

This function generates normal random numbers truncated to $[0, 1]$ based on the 90% confidence interval calculating the distribution parameter numerically from the 90%-confidence interval.

Usage

```
r0_1norm90ci_numeric(n, lower, upper, relativeTolerance = 0.05)
```

Arguments

n	Number of generated observations.
lower	numeric; lower bound of the 90% confidence intervall.
upper	numeric; upper bound of the 90% confidence intervall.
relativeTolerance	numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval.

Details

#ToDo

random	<i>Generate random numbers for a certain probability distribution.</i>
--------	--

Description

This function generates multivariate random numbers for general multivariate distributions.

Usage

```
random(rho, n, method, ...)
```

Arguments

rho	Distribution to be randomly sampled.
n	Number of generated observations.
method	Particular method to be used for random number generation.
...	Optional arguments to be passed to the particular random number generating function.

random.default	<i>Generate random numbers based on the first two moments of a certain probability distribution.</i>
----------------	--

Description

This function generates random numbers for general multivariate distributions that can be characterized by the joint first two moments, viz. the mean and covariance.

Usage

```
## Default S3 method:
random(rho, n, method, ...)
```

Arguments

rho	list; Distribution to be randomly sampled.
n	Number of generated observations
method	Particular method to be used for random number generation.
...	Optional arguments to be passed to the particular random number generating function.

random.estimate	<i>Generate Random Numbers for an Estimate.</i>
-----------------	---

Description

This function generates random numbers for general multivariate distributions that are defined as an [estimate](#).

Usage

```
## S3 method for class 'estimate'
random(rho, n, method = "calculate", ...)
```

Arguments

rho	estimate object; Multivariate distribution to be randomly sampled.
n	Number of generated observations
method	Particular method to be used for random number generation.
...	Optional arguments to be passed to the particular random number generating function.

Details

Generation of uncorrelated components: Implementation: [random_estimate_1d](#)

Generation of correlated components: Implementation: [rmvnorm90ci_exact](#)

See Also

[estimate](#)

Examples

```
variable=c("revenue","costs")
distribution=c("norm","norm")
lower=c(10000, 5000)
upper=c(100000, 50000)
estimateObject<-estimate(variable, distribution, lower, upper)
x<-random(rho=estimateObject, n=10000)
apply(X=x, MARGIN=2, FUN=quantile, probs=c(0.05, 0.95))
cor(x)
colnames(x)
summary(x)
hist(x[, "revenue"])
hist(x[, "costs"])
```

random_estimate_1d	<i>Generate univariate random numbers based on an estimate.</i>
--------------------	---

Description

This function generates random numbers for general univariate distributions.

Usage

```
random_estimate_1d(rho, n, method = "calculate", ...)
```

Arguments

rho	estimate object; Univariate distribution to be randomly sampled.
n	Number of generated observations
method	Particular method to be used for random number generation.
...	Optional arguments to be passed to the particular random number generating function.

Details

method can be either "calculate" (the default) or "fit".

The following table shows the available distributions and the implemented generation method:

Identification	Distribution	method
const	ToDo	calculate
norm	Normal distribution	calculate, fit
posnorm	ToDo	calculate
Ø_1norm	ToDo	calculate
beta	Beta distribution	fit
cauchy	ToDo	fit
logis	ToDo	fit
t	ToDo	fit
chisq	ToDo	fit
chisqnc	ToDo: implement?	fit
exp	ToDo	fit
f	ToDo	fit
gamma	ToDo	fit
lnorm	ToDo	fit
unif	ToDo	calculate, fit
weibull	ToDo	fit
triang	ToDo	fit
gompertz	ToDo	fit
pert	ToDo	fit
tnorm	Truncated normal distribution	fit

See Also

For method="calculate": [rdist90ci_exact](#), [rposnorm90ci_numeric](#) and [r0_1norm90ci_numeric](#);
for method="fit": [rdistq_fit](#)

rdist90ci_exact	<i>Generate univariate random numbers based on the 90%-confidence interval.</i>
-----------------	---

Description

This function generates random numbers for general univariate distributions based on the 90% confidence interval.

Usage

```
rdist90ci_exact(distribution, n, lower, upper)
```

Arguments

- distribution character; A character string that defines the univariate distribution to be randomly sampled.
- n Number of generated observations.
- lower numeric; lower bound of the 90% confidence intervall.
- upper numeric; upper bound of the 90% confidence intervall.

Details

The follwing table shows the available distributions and their identification as a character string:

Distribution encoding	Distribution
const	ToDo
norm	ToDo
pos_norm	ToDo
norm_0_1	ToDo
pois	ToDo
binom	ToDo
unif	ToDo
lnorm	ToDo
lnorm_lim2	ToDo

rdistq_fit	<i>Generate univariate random numbers based on quantiles.</i>
------------	---

Description

This function generates random numbers for a set of univariate distributions based on the distribution quantiles. Internally, this is achieved by fitting the distribution function to the given quantiles using `rriskFitdist.perc`.

Usage

```
rdistq_fit(distribution, n, percentiles = c(0.05, 0.5, 0.95), quantiles,
  tolConv = 0.001, fit.weights = rep(1, length(percentiles)))
```

Arguments

distribution	A character string that defines the univariate distribution to be randomly sampled.
n	Number of generated observations.
percentiles	Numeric vector giving the percentiles.
quantiles	Numeric vector giving the quantiles.
tolConv	positive numerical value, the absolute convergence tolerance for reaching zero by fitting distributions <code>get.norm.par</code> will be shown.
fit.weights	numerical vector of the same length as a probabilities vector <code>p</code> containing positive values for weighting quantiles. By default all quantiles will be weighted by 1.

Details

The following table shows the available distributions and their identification as a character string:

Identification	Distribution	Number of quantiles
<code>norm</code>	Normal distribution	≥ 2
<code>beta</code>	Beta distribution	ToDo
<code>cauchy</code>	ToDo	ToDo
<code>logis</code>	ToDo	ToDo
<code>t</code>	ToDo	ToDo
<code>chisq</code>	ToDo	ToDo
<code>chisqnc</code>	ToDo: implement?	ToDo
<code>exp</code>	ToDo	ToDo
<code>f</code>	ToDo	ToDo
<code>gamma</code>	ToDo	ToDo
<code>lnorm</code>	ToDo	ToDo
<code>unif</code>	ToDo	ToDo
<code>weibull</code>	ToDo	ToDo
<code>triang</code>	Triangular Distribution, Note: package <code>mc2d</code> needed.	ToDo
<code>gompertz</code>	Gompertz distribution, Note: package <code>eha</code> needed.	ToDo
<code>pert</code>	The (modified) PERT distribution, Note: package <code>mc2d</code> needed.	ToDo
<code>tnorm</code>	Truncated normal distribution	ToDo

The default for `percentiles` is 0.05, 0.5 and 0.95, so for the default, the `quantiles` argument should

be a vector with 3 elements. If this is to be longer, the percentiles argument has to be adjusted to match the length of quantiles.

Value

ToDo

rmvnorm90ci_exact	<i>Generate normal distributed multivariate random numbers based on the 90%-confidence interval.</i>
-------------------	--

Description

This function generates normal distributed multivariate random numbers based on the 90%-confidence interval.

Usage

```
rmvnorm90ci_exact(n, lower, upper, correlationMatrix)
```

Arguments

n	Number of generated observations.
lower	numeric vector; lower bound of the 90% confidence intervall.
upper	numeric vector; upper bound of the 90% confidence intervall.
correlationMatrix	numeric symmetric matrix; correlation matrix; In particular, all diagonal elements must be equal to 1.

row.names.estimate	<i>Return the variable names of an estimate object.</i>
--------------------	---

Description

This function returns the variable names of an [estimate](#) object which is identical to `row.names(x$base)`.

Usage

```
## S3 method for class 'estimate'
row.names(x)
```

Arguments

x	an estimate object.
---	-------------------------------------

See Also

[estimate](#), [names.estimate](#), [corMat.estimate](#)

rposnorm90ci_iter	<i>Generate positive normal random numbers based on the 90%-confidence interval.</i>
-------------------	--

Description

This function generates positive normal random numbers based on the 90% confidence interval using an iteration algorithm.

Usage

```
rposnorm90ci_iter(n, lower, upper, relativeTolerance = 0.05, maxIter = 40)
```

Arguments

n	Number of generated observations.
lower	numeric; lower bound of the 90% confidence intervall.
upper	numeric; upper bound of the 90% confidence intervall.
relativeTolerance	numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval.
maxIter	numeric; maximum number of iterations.

Details

The generation of random numbers is repeated until the generated 90% - confidence interval is close enough to the desired value.

rposnorm90ci_numeric	<i>Generate positive normal random numbers based on the 90%-confidence interval.</i>
----------------------	--

Description

This function generates positive normal random numbers based on the 90% confidence interval calculating the distribution parameter numerically from the 90%-confidence interval.

Usage

```
rposnorm90ci_numeric(n, lower, upper, relativeTolerance = 0.05)
```

Arguments

n	Number of generated observations.
lower	numeric; lower bound of the 90% confidence intervall.
upper	numeric; upper bound of the 90% confidence intervall.
relativeTolerance	numeric; the relative tolerance level of deviation of the generated confidence interval from the specified interval.

Details

#ToDo

sort.summary.eviSimulation

Sort Summarized EVI Simulation Results..

Description

Sort summarized EVI simulation results according to their EVI.

Usage

```
## S3 method for class 'summary.eviSimulation'
sort(x, decreasing = TRUE, ...,
     along = row.names(x$summary$evi)[[1]])
```

Arguments

x	An object of class summary.eviSimulation.
decreasing	logical; if the evi should be sorted in decreasing order.
...	Further arguments #ToDo
along	character; the name of the valuation variable along which evi should be sorted.

Value

An object of class summary.eviSimulation.

See Also

[eviSimulation](#), [summary.eviSimulation](#), [sort](#)

summary.eviSimulation *Summarize EVI Simulation Results*

Description

summary.eviSimulation produces result summaries of the results of Expected Value of Information (EVI) simulation obtained by the function [eviSimulation](#).

Usage

```
## S3 method for class 'eviSimulation'
summary(object, ..., digits = max(3,
  getOption("digits") - 3))
```

Arguments

object	An object of class eviSimulation.
...	Further arguments passed to summary.welfareDecisionAnalysis .
digits	how many significant digits are to be used for numeric and complex x. The default, NULL, uses getOption("digits") . This is a suggestion: enough decimal places will be used so that the smallest (in magnitude) number has this many significant digits, and also to satisfy nsmall. (For the interpretation for complex numbers see signif .)

Value

An object of class summary.eviSimulation.

See Also

[eviSimulation](#), [print.summary.eviSimulation](#), [summary.welfareDecisionAnalysis](#)

summary.mcSimulation *Summarize Results from Monte Carlo Simulation.*

Description

A summary of the results of a Monte Carlo simulation obtained by the function [mcSimulation](#) is produced.

Usage

```
## S3 method for class 'mcSimulation'
summary(object, ..., digits = max(3,
  getOption("digits") - 3), variables.y = names(object$y), variables.x = if
  (classicView) names(object$x), classicView = FALSE, probs = c(0, 0.1,
  0.25, 0.5, 0.75, 0.9, 1))
```

Arguments

object	An object of class mcSimulation.
...	Further arguments passed to <code>summary.data.frame</code> (classicView=TRUE) or <code>format</code> (classicView=FALSE).
digits	how many significant digits are to be used for numeric and complex x. The default, NULL, uses <code>getOption("digits")</code> . This is a suggestion: enough decimal places will be used so that the smallest (in magnitude) number has this many significant digits, and also to satisfy nsmall. (For the interpretation for complex numbers see <code>signif</code> .)
variables.y	character or character vector; Names of the components of the simulation function (model_function) which results shall be displayed. Defaults to all components.
variables.x	character or character vector; Names of the components of the input variables to the simulation function, i.e. the names of the variables in the input estimate which random sampling results shall be displayed. Defaults to all components.
classicView	logical; if TRUE the results are summarized using <code>summary.data.frame</code> , if FALSE further output is produced and the quantile information can be chosen. Cf. section Value and argument probs. Default is FALSE.
probs	numeric vector of quantiles that shall be displayed if classicView=FALSE.

Value

An object of class `summary.mcSimulation`.

chance_loss
chance_zero
chance_gain

See Also

`mcSimulation`, `print.summary.mcSimulation`, `summary.data.frame`

`summary.welfareDecisionAnalysis`

Summarize Decsion Analysis Results.

Description

`summary.welfareDecisionAnalysis` produces result summaries of the results of decision analysis simulation obtained by the function `welfareDecisionAnalysis`.

Usage

```
## S3 method for class 'welfareDecisionAnalysis'
summary(object, ..., digits = max(3,
  getOption("digits") - 3))
```

Arguments

object	An object of class welfareDecisionAnalysis.
...	Further arguments passed to format .
digits	how many significant digits are to be used for numeric and complex x. The default, NULL, uses getOption("digits") . This is a suggestion: enough decimal places will be used so that the smallest (in magnitude) number has this many significant digits, and also to satisfy nsmall. (For the interpretation for complex numbers see signif .)

Value

An object of class `summary.welfareDecisionAnalysis`.

See Also

[welfareDecisionAnalysis](#), [print.summary.welfareDecisionAnalysis](#), [format](#)

uncertaintyAnalysis	<i>Uncertainty Analysis Wrapper Function.</i>
---------------------	---

Description

This function performs a Monte Carlo simulation from input files and analyses the results via Partial Least Squares Regression (PLSR) and calculates the Variable Importance on Projection (VIP). Results are saved as plots.

Usage

```
uncertaintyAnalysis(inputFilePath, outputDirectory, modelFunction,
  NumberOfSimulations, randomMethod = "calculate",
  functionSyntax = "globalNames", write_table = TRUE, indicators = FALSE,
  log_scales = FALSE, oldInputStandard = FALSE)
```

Arguments

inputFilePath	Path to input csv file, which gives the input estimate .
outputDirectory	Path where the result plots and tables are saved.
modelFunction	The model function.

NumberOfSimulations	The number of Monte Carlo simulations to be performed.
randomMethod	ToDo
functionSyntax	ToDo
write_table	logical; If the full Monte Carlo simulation results and PLSR results should be written to file.
indicators	logical; If indicator variables should be respected specially.
log_scales	logical; If the scales in the pls plots should be logarithmic.
oldInputStandard	logical; If the old input standard should be used (estimate_read_csv_old).

See Also

[mcSimulation](#), [estimate](#), [estimate_read_csv](#)

welfareDecisionAnalysis

Analysis of the Underlying Welfare Based Decision Problem

Description

The optimal choice between two different opportunities is calculated. This decision is based on minimizing the Expected Net Loss (ENL).

Usage

```
welfareDecisionAnalysis(estimate, model, numberOfSimulations,
  functionSyntax = "data.frameNames")
```

Arguments

estimate	estimate object describing the distribution of the input variables.
model	either a function or a list with two functions: <code>list(p1,p2)</code> . In the first case the function is the net benefit of project approval vs. the status quo. In the second case the element p1 is the function valuing the first project and the element p2 valuing the second project.
numberOfSimulations	integer; number of simulations to be used in the underlying Monte Carlo analysis
functionSyntax	function character; function syntax used in the model function(s).

Details

This principle is along the line described in Hubbard (2014). The Expected Opportunity Loss (EOL) is defined as the Expected Net Loss (ENL) for the best decision. The best decision minimises the ENL. The EOL is always conditional on the available information (I): $EOL=EOL(I)$. Here, the available information is the supplied estimate. One can show that in the case of two alternatives, minimization of EOL is equivalent to maximization of the Expected Net Benefit.

Value

An object of class `welfareDecisionAnalysis` with the following elements:

<code>enbPa</code>	Expected Net Loss (ENL) in case of project approval (PA)
<code>enbSq</code>	Expected Net Loss (ENL) in case of status quo (SQ)
<code>eol</code>	Expected Opportunity Loss (EOL)
<code>optimalChoice</code>	The optimal choice, i.e. either project approval (PA) or the status quo (SQ)

See Also

[mcSimulation](#), [estimate](#), [summary.welfareDecisionAnalysis](#)

Examples

```
#####
# Example 1 (Creating the estimate from the command line):
#####
# Create the estimate object:
variable=c("revenue","costs")
distribution=c("posnorm","posnorm")
lower=c(10000, 5000)
upper=c(100000, 50000)
costBenefitEstimate<-estimate(variable, distribution, lower, upper)
# (a) Define the model function without name for the return value:
profit<-function(x){
  x$revenue-x$costs
}
# Perform the decision analysis:
myAnalysis<-welfareDecisionAnalysis(estimate=costBenefitEstimate,
                                     model=profit,
                                     numberOfSimulations=100000,
                                     functionSyntax="data.frameNames")

# Show the analysis results:
print(summary((myAnalysis)))
#####
# (b) Define the model function with a name for the return value:
profit<-function(x){
  list(Profit=x$revenue-x$costs)
}
# Perform the decision analysis:
myAnalysis<-welfareDecisionAnalysis(estimate=costBenefitEstimate,
                                     model=profit,
                                     numberOfSimulations=100000,
                                     functionSyntax="data.frameNames")

# Show the analysis results:
print(summary((myAnalysis)))
#####
# (c) Two decision variables:
decisionModel<-function(x){
  list(Profit=x$revenue-x$costs,
       Costs=-x$costs)
}
```



```
# Perform the decision analysis:
myAnalysis<-welfareDecisionAnalysis(estimate=costBenefitEstimate,
                                     model=decisionModel,
                                     numberOfSimulations=100000,
                                     functionSyntax="data.frameNames")

# Show the analysis results:
print(summary((myAnalysis)))
```

Index

as.data.frame, [3](#)
as.data.frame.mcSimulation, [3](#)

beta, [30](#), [32](#)

colors, [16](#)
constrOptim, [23](#), [24](#)
corMat, [4](#), [7](#)
corMat.estimate, [4](#), [21](#), [33](#)

data.frame, [7](#)
decisionSupport, [5](#)
decisionSupport-package
 (decisionSupport), [5](#)

estimate, [4](#), [6](#), [7](#), [8–12](#), [17](#), [19](#), [21](#), [29](#), [33](#),
 [38–40](#)
estimate_read_csv, [7](#), [7](#), [10](#), [11](#), [39](#)
estimate_read_csv_old, [9](#), [39](#)
estimate_write_csv, [7](#), [9](#), [10](#)
eviSimulation, [5](#), [6](#), [11](#), [17](#), [26](#), [35](#), [36](#)

fit, [30](#)
format, [37](#), [38](#)

getOption, [36–38](#)
gompertz, [32](#)

hist, [16](#)
hist.mcSimulation, [15](#), [19](#)

individualEvpSimulation, [5](#), [16](#)

make.names, [3](#)
mcSimulation, [6](#), [12](#), [15–17](#), [18](#), [25](#), [26](#), [36](#),
 [37](#), [39](#), [40](#)

names.estimate, [4](#), [7](#), [21](#), [33](#)
norm, [30](#), [32](#)

paramposnorm90ci, [21](#)
paramtnormci, [22](#)

paramtnormci_fit, [23](#)
paramtnormci_numeric, [24](#)
pert, [32](#)
pretty, [16](#)
print.mcSimulation, [19](#), [25](#)
print.summary.eviSimulation, [26](#), [36](#)
print.summary.mcSimulation, [26](#), [37](#)
print.summary.welfareDecisionAnalysis,
 [27](#), [38](#)

r0_1norm90ci_numeric, [27](#), [31](#)
random, [28](#)
random.default, [28](#)
random.estimate, [6](#), [7](#), [19](#), [29](#)
random_estimate_1d, [29](#), [30](#)
rdist90ci_exact, [31](#), [31](#)
rdistq_fit, [31](#), [31](#)
read.csv, [8–10](#)
rmvnorm90ci_exact, [29](#), [33](#)
row.names, [7](#)
row.names.estimate, [4](#), [7](#), [21](#), [33](#)
rposnorm90ci_iter, [34](#)
rposnorm90ci_numeric, [31](#), [34](#)
rriskFitdist.perc, [32](#)

scan, [8](#), [9](#)
signif, [36–38](#)
sort, [35](#)
sort.summary.eviSimulation, [35](#)
summary.data.frame, [37](#)
summary.eviSimulation, [26](#), [35](#), [36](#)
summary.mcSimulation, [19](#), [26](#), [36](#)
summary.welfareDecisionAnalysis, [27](#), [36](#),
 [37](#), [40](#)

tnorm, [22–25](#), [30](#), [32](#)
triang, [32](#)

uncertaintyAnalysis, [6](#), [38](#)

welfareDecisionAnalysis, [5](#), [6](#), [12](#), [17](#), [27](#),
[37](#), [38](#), [39](#)
write.csv, [11](#)