

Package ‘pbo’

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Type Package

Title Probability of Backtest Overfitting

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Description Following the method of Bailey et al., computes for a collection of candidate models the probability of backtest overfitting, the performance degradation and probability of loss, and the stochastic dominance.

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URL <https://github.com/mrbcuda/pbo>

BugReports <https://github.com/mrbcuda/pbo/issues>

Depends utils, lattice

Suggests PerformanceAnalytics, foreach, grid, latticeExtra, testthat,
doParallel, knitr

VignetteBuilder knitr

NeedsCompilation no

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pbo-package*Probability of backtest overfitting.***Description**

Computes the probability of backtest overfitting

Details

Implements algorithms for computing the probability of backtest overfitting, performance degradation and probability of loss, and first- and second-order stochastic dominance, based on the approach specified in Bailey et al., September 2013. Provides a collection of pre-configured plots based on lattice graphics.

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References

See Bailey, David H. and Borwein, Jonathan M. and Lopez de Prado, Marcos and Zhu, Qiji Jim, The Probability of Back-Test Overfitting (September 1, 2013). Available at SSRN. See <http://ssrn.com/abstract=2326253> or <http://dx.doi.org/10.2139/ssrn.2326253>.

dotplot.pbo*PBO in-sample selection dot plot.***Description**

Draws an annotated dot plot of study selection sorted by in-sample selection frequency.

Usage

```
## S3 method for class 'pbo'
dotplot(x, data = NULL, main,
         xlab = "Sorted Study Number (N)", ylab = "IS Selection Frequency",
         show_config = TRUE, show_grid = TRUE, sel_threshold = 0, ...)
```

Arguments

<code>x</code>	a pbo object as returned by pbo .
<code>data</code>	should not be used
<code>main</code>	plot title, default computed internally, passed to dotplot .
<code>xlab</code>	x-axis label with default, passed to dotplot .
<code>ylab</code>	y-axis label with default, passed to dotplot .

show_config whether to show the study dimension annotations, default TRUE
 show_grid whether to show the grid panel, default TRUE
 sel_threshold the minimum in-sample frequency subsetting threshold, default 0; selection frequencies at or below this value will be omitted
 ... other parameters as passed to [dotplot](#).

See Also

[pbo](#), [histogram.pbo](#), [xyplot.pbo](#)

[histogram.pbo](#)

PBO rank logits histogram.

Description

Draws an annotated histogram of PBO rank logits.

Usage

```
## S3 method for class 'pbo'
histogram(x, data = NULL, show_pbo = TRUE,
           show_regions = TRUE, show_config = TRUE, col_bar = "#cc99cc",
           col_line = "#3366cc", ...)
```

Arguments

x an object of class pbo as returned by [pbo](#).
 data should not be used
 show_pbo whether to show the PBO value annotation, default TRUE
 show_regions whether to show the overfit region annotations, default TRUE
 show_config whether to show the study dimension annotations, default TRUE
 col_bar histogram bar fill color passed to histogram panel
 col_line density plot line color passed to density plot panel
 ... other parameters passed to [histogram](#), [densityplot](#), or [panel.abline](#).

Details

Uses **lattice** function [histogram](#), [densityplot](#), and [panel.abline](#) panels together with class-specific annotations.

See Also

[pbo](#), [dotplot.pbo](#), [xyplot.pbo](#)

pbo*Probability of backtest overfitting***Description**

Performs the probability of backtest overfitting computations.

Usage

```
pbo(m, s = 4, f = NA, threshold = 0, inf_sub = 6,
    allow_parallel = FALSE)
```

Arguments

- m** a $T \times N$ data frame of returns, where T is the samples per study and N is the number of studies.
- s** the number of subsets of **m** for CSCV combinations; must evenly divide **m**
- f** the function to evaluate a study's performance; required
- threshold** the performance metric threshold (e.g. 0 for Sharpe, 1 for Omega)
- inf_sub** infinity substitution value for reasonable plotting
- allow_parallel** whether to enable parallel processing, default FALSE

Details

This function performs the probability of backtest overfitting calculation using a combinatorially-symmetric cross validation (CSCV) approach.

Value

object of class **pbo** containing list of PBO calculation results and settings

References

Baily et al., "The Probability of Backtest Overfitting," http://papers.ssrn.com/sol3/papers.cfm?abstract_id=2326253

Examples

```
## Not run:
require(pbo)
require(PerformanceAnalytics)
n <- 100
t <- 1000
s <- 8
m <- data.frame(matrix(rnorm(n*t,mean=0,sd=1),
  nrow=t,ncol=n,byrow=TRUE,
  dimnames=list(1:t,1:n)),
```

```
check.names=FALSE)
p <- pbo(m,s,f=Omega,threshold=1)

## End(Not run)
```

pbo_show_config

Writes grid text to a default predetermined location.

Description

Writes grid text to a default predetermined location.

Usage

```
pbo_show_config(p)
```

Arguments

p an object of class **pbo** as returned by [pbo](#).

Note

Meant for internal use only.

xyplot.pbo

PBO xy-plots

Description

Draws an annotated plot of performance degradation and probability of loss.

Usage

```
## S3 method for class 'pbo'
xyplot(x, data = NULL, plotType = "cscv", show_eqn = TRUE,
       show_threshold = TRUE, show_config = TRUE, show_rug = TRUE,
       show_prob = TRUE, show_grid = TRUE, increment = 0.01,
       osr_threshold = 0, sel_threshold = 0, xlab, ylab, main, lwd = 1,
       ylab_left, ylab_right, col_bar, col_line, col_sd1 = "#3366cc",
       col_sd2 = "#339999", lty_sd = c(1, 2, 4), ...)
```

Arguments

<code>x</code>	a pbo object as returned by pbo .
<code>data</code>	should not be used
<code>plotType</code>	one of <code>cscv</code> , <code>degradation</code> , <code>dominance</code> , <code>pairs</code> , <code>ranks</code> or <code>selection</code> .
<code>col_bar</code>	histogram bar fill color
<code>col_line</code>	density plot line color
<code>col_sd1</code>	color of two first-order stochastic dominance lines
<code>col_sd2</code>	color of the single second-order stochastic dominance line
<code>xlab</code>	x-axis label, default computed if not provided
<code>ylab</code>	y-axis label, default computed if not provided
<code>main</code>	plot title, default computed if not provided
<code>lwd</code>	line width, default 1, passed to panels and legends
<code>lty_sd</code>	line type array for stochastic dominance plot, e.g. <code>c(2,3,5)</code>
<code>ylab_left</code>	dominance plot left-hand axis label
<code>ylab_right</code>	dominance plot right-hand axis label
<code>increment</code>	stochastic dominance distribution generator increment, e.g. 0.1 steps
<code>osr_threshold</code>	out-of-sample rank threshold for filtering, default 0
<code>sel_threshold</code>	selection frequency threshold for filtering, default 0
<code>show_eqn</code>	whether to show the line equation annotation, default TRUE
<code>show_threshold</code>	whether to show the probability of loss annotation, default TRUE
<code>show_config</code>	whether to show the study dimension annotations, default TRUE
<code>show_grid</code>	whether to show the panel grid, default TRUE
<code>show_prob</code>	whether to show the probability value in dominance plot, default TRUE
<code>show_rug</code>	whether to show scatter rugs near the axes, default TRUE
<code>...</code>	other parameters passed to xyplot or its panels

Details

Provides several variations of xy-plots suitable for presentation of PBO analysis results. Use the `plotType` argument to indicate which variation or result to plot:

- The `cscv` type shows in-sample and out-of-sample results by CSCV iteration case (default).
- The `degradation` type shows the performance degradation regression fit results and the probability of loss.
- The `dominance` type shows the results of the first-order and second-order stochastic dominance analysis using two axes.
- The `pairs` type shows the in-sample and out-of-sample case selections.
- The `ranks` type shows the sorted performance ranks results.
- The `selection` type shows the case selection frequencies.

See Also

[pbo](#), [histogram.pbo](#), [xyplot.pbo](#)

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