Package 'overlapping'

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Type Package

Title Estimation of Overlapping in Empirical Distributions

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Description Functions for estimating the overlapping area of two or more kernel density estimations from empirical data.

Depends R (>= 3.0.0), ggplot2, testthat

License GPL-2

Encoding UTF-8

NeedsCompilation no

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R topics documented:

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boot.overlap   *Nonparametric Bootstrap for estimated overlapping area*

Description

Bootstrap the estimated overlapping area of two or more kernel density estimations from empirical data.

Usage

```
boot.overlap( x, B = 1000, ... )
```

Arguments

x	list of numerical vectors to be compared; each vector is an element of the list
B	integer, number of bootstrap draws
...	options, see function overlap for details

Details

If the list x contains more than two elements (i.e. more than two distributions) it computes bootstrap overlapping between all q number of paired distributions. For example, if x contains three elements, then $q = 3$; if x contains four elements, then $q = 6$, and so on.

Value

It returns a list containing the following components:

OVboot_stats	Data frame $q \times 3$; each row containing the following statistics: estOV, estimated overlapping area, $\hat{\eta}$; bias, difference between the bootstrap's expected value and the observed value of the overlapping area: $E(\hat{\eta}^*) - \hat{\eta}$; se, bootstrap standard error $\sigma_{\hat{\eta}}$.
OVboot_dist	Matrix $B \times q$, B rows (bootstrap replicates) and q columns (depending on the number of elements of x); each column is a boostrap distribution of an overlapping index.

Note

Call function overlap.

Thanks to Jeremy Vollen for suggestions.

Author(s)

Massimiliano Pastore

References

Pastore, M. (2018). Overlapping: a R package for Estimating Overlapping in Empirical Distributions. The Journal of Open Source Software, 3 (32), 1023. doi: 10.21105/joss.01023

Pastore, M., Calcagnì, A. (2019). Measuring Distribution Similarities Between Samples: A Distribution-Free Overlapping Index. Frontiers in Psychology, 10:1089. doi: 10.3389/fpsyg.2019.01089

Examples

```
set.seed(20150605)
x <- list(X1=rnorm(100), X2=rt(50,8), X3=rchisq(80,2))

## bootstrapping
out <- boot.overlap( x, B = 10 )
out$OVboot_stats

# bootstrap quantile intervals
apply( out$OVboot_dist, 2, quantile, probs = c(.05, .9) )

# plot of bootstrap distributions
Y <- stack( data.frame( out$OVboot_dist ))
ggplot( Y, aes( values )) + facet_wrap( ~ind ) + geom_density()
```

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cutnumeric	*Numerical conversion*

Description

It divides a numerical variable x in classes, and returns for each class the central value.

Internal function, generally not to be called by the user.

Usage

```
cutnumeric( x, n = 1000 )
```

Arguments

x	numeric vector
n	number of classes

Details

It calls the cut function, and then converts factor classes in numeric classes, returning for each class its central value.

Value

It returns a numerical vector. The values are the central points of classes obtained by the function cut.

Author(s)

Massimiliano Pastore

See Also

[cut](cut)

Examples

```
x <- rnorm(50)
cutnumeric(x,5)
```

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final.plot	*Final plot*

Description

Graphical representation of estimated densities and overlapping area.

Usage

```
final.plot( x, OV = NULL )
```

Arguments

x	list of numerical vectors to be compared; each vector is an element of the list, see overlap
OV	Optional vector of overlapping areas obtained by overlap

Details

It requires the package ggplot2.

Author(s)

Massimiliano Pastore

Examples

```
set.seed(20150605)
x <- list(X1=rnorm(100),X2=rt(50,8),X3=rchisq(80,2))
out <- overlap(x)
final.plot(x,out$OV)
```

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overlap | *Overlapping estimation*

Description

It gives the overlapped estimated area of two or more kernel density estimations from empirical data.

Usage

```
overlap( x, nbins = 1024, plot = FALSE,
    partial.plot = FALSE, boundaries = NULL,
    return.complete.data = FALSE, ... )
```

Arguments

x	list of numerical vectors to be compared; each vector is an element of the list
nbins	number of equally spaced points at which the overlapping density is evaluated; see [density](#) for details
plot	logical, if TRUE, final plot of estimated densities and overlapped areas is produced
partial.plot	logical, if TRUE, partial paired distributions are plotted
boundaries	an optional list for bounded distributions, see Details
return.complete.data	
	logical, if TRUE, return a data frame with information used for computing overlapping (see Value).
...	optional arguments to be passed to function [density](#)

Details

If the list x contains more than two elements (i.e. more than two distributions) it computes overlapping between all paired distributions. Partial plots refer to these paired distributions.

If plot=TRUE, all overlapped areas are plotted. It requires ggplot2.

The optional list boundaries must contain two elements: from and to, indicating the empirical limits of input variables. Each element must be of length equal to the input data list x or, at least, length one when all boundaries are equal for all distributions. See examples below.

Value

It returns a list containing the following components:

DD	Data frame with information used for computing overlapping, containing the following variables (only if return.complete.data = TRUE): x, coordinates of the points where the density is estimated; y1 and y2, densities; ovy, density for estimating overlapping area (i.e. min(y1,y2)); ally, density for estimating whole area (i.e. max(y1,y2)); dominance, indicates which distribution has the highest density; k, label indicating which distributions are compared.

| OV | Estimates of overlapped areas relative to each pair of distributions. |
| xpoints | List of abscissas of intersection points among the density curves. |

Note

Call function `final.plot`.

Author(s)

Massimiliano Pastore

References

Pastore, M. (2018). Overlapping: a R package for Estimating Overlapping in Empirical Distributions. The Journal of Open Source Software, 3 (32), 1023. doi: 10.21105/joss.01023

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Examples

```
set.seed(20150605)
x <- list(X1=rnorm(100), X2=rt(50,8), X3=rchisq(80,2))
out <- overlap(x, plot=TRUE)
out$OV

# including boundaries
x <- list(X1=runif(100), X2=runif(100,.5,1))
boundaries <- list( from = c(0,.5), to = c(1,1) )
out <- overlap(x, plot=TRUE, boundaries=boundaries)
out$OV

# equal boundaries
x <- list(X1=runif(100), X2=runif(50), X3=runif(30))
boundaries <- list( from = 0, to = 1 )
out <- overlap(x, plot=TRUE, boundaries=boundaries)
out$OV

# changing kernel
out <- overlap(x, plot=TRUE, kernel="rectangular")
out$OV
```

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