Package'DVDtest'

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Type Package

Title Difference between Varying Distributions Test (DVDtest)

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Description See DVDtest.

RoxygenNote 6.1.1

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Imports gamlss, mgcv, parallel, gamlss.dist, ggplot2, reshape2

R topics documented:

DVDtest-package   *Difference between Varying Distributions Test (DVDtest)*

Description

This package contains a series of internal and external functions of Difference between Varying Distributions Test (DVDtest), which tests the pointwise group differences between two varying distributions.

Author(s)

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See Also

DVDtest

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DVDplot   *Plotting DVDtest*

Description

Plot a list of the DVDtest-related figures via ggplot2

Usage

```
DVDplot(tobj, kxlab = NULL, kylab = NULL, kname = NULL)
```

Arguments

tobj	a return test object of `DVDtest`
kxlab	a title for the x axis, `.index`
kylab	a title for the y axis, `.value`
kname	a name for k, e.g. ROI in the references

Details

Fig `pfig` illustrates a collection of the p value curves among all k. Fig `kfig` illustrates a collection list of the figures with varying distributions among all k, highlighted the zones of small p values in dark, where `ydata1` and `ydata2` mark as red and blue, repectively. The dashed and dotted lines denote the smooth mean function and +/- 2 sigma, respectively.

Value

a list of ggplot objects on p value curves and varying distributions

Note

Please contact the maintainer if need more details.

Author(s)

Meng Xu

References

reiss-EMR18.pdf

See Also

Examples in `DVDtest`

DVDtest   *Difference between Varying Distributions Test (DVDtest)*

Description

Testing the difference of two varying distributions.

Usage

```
DVDtest(ydata1, ydata2, nperm, grid, dist.method = "wass",
  mgcv.gam = TRUE, ..., exclude = NULL, permadj = FALSE,
  mc.cores = 1)
```

Arguments

ydata1
: a data.frame or a list of data.frame, containing at least 3 columns called '.obs', '.index' and '.value' which specify which curve the point belongs to (.obs) at which ('.index') it was observed and the observed value ('.value'). See details in the package refund. Other columns are available as well for modelling the varying distributions.

ydata2
: same as ydata1.

nperm
: a scalar, number of permutation

grid
: a vector, evaluation grids of .index

dist.method
: the distance measure to be used. This must be one of Wasserstein distance ('wass'), 'L2' distance, 'L1' distance and 'Hellinger'. Defaults to 'wass'.

mgcv.gam
: a logical variable, whether to apply mgcv::gam for eastimating distributions, whose parameters are a smooth function of a continuous variable. If FALSE, gamlss::gamlss is adopted.

...
: passed to arguments of gam or gamlss. If mgcv.gam = TRUE, ... should include formula, family (=gaulss()) and other optional arguments in mgcv::gam. Otherwise, ... passed to arguments inside of gamlss::gamlss.

exclude
: passed to exclude inside of predict.gam in case mgcv.gam = TRUE.

permadj
: a logical variable, whether to adjust the permutated data to cover the entire range, esp. in case of sparsity. Defaults to FALSE.

mc.cores
: passed to mc.cores inside of mclapply (not available on Windows unless mc.cores = 1). Defaults to 1. See extra info in Note.

Details

This is the Details section

Value

.index
: a vector, evaluation grids.

pval
: a vector or matrix of (corrected) p values.

vdparam
: a list of paramters of varying distributions.

Note

- If ydata1 and ydata2 are lists of data.frames, the lenghs of two lists must be the same.
- If mgcv.gam is TRUE, ... and exclue are NULL (default settings), then defaults to
  formula <- list(.value ~ s(.index) + s(.obs, bs = "re"), ~ s(.index))
  and exclude <- "s(.obs)", repectively.
- Multicore calculation is dealing with the permutation.
- Normal distribution in mgcv::gam and BCCG, BCT and BCPE in gamlss::gamlss are currently
  supported by DVDtest for fitting a GAMLSS-type varying distributions. Please contact the
  maintainer if need further supporting.

Author(s)

Meng Xu, Philip Reiss

References

reiss-EMR18.pdf

Examples

```
## Data Generation ##
 p <- 6
 mu1 <- function(t) 0.2*(p-1)*sin(pi*t)+t+1
 mu2 <- function(t) -0.2*(p-1)*sin(pi*t)+t+1
 sig1 <- function(t) t+1
 sig2 <- sig1
 nperson <- 10
 fun1 <- function(t) rnorm(nperson, mu1(t), sig1(t))
 fun2 <- function(t) rnorm(nperson, mu2(t), sig2(t))
 tp <- seq(0,1,,10)
 data1 <- sapply(tp,fun1)
 data2 <- sapply(tp,fun2)

 library(reshape2)
 colnames(data2) <- colnames(data1) <- tp
 rownames(data2) <- 1:nperson+2*nperson
 dg1 <- melt(data1)
 dg2 <- melt(data2)
 colnames(dg1) <- colnames(dg2) <- c('.obs','.index','.value')
 dg1$.obs <- as.factor(dg1$.obs)
 dg2$.obs <- as.factor(dg2$.obs)
 # library(ggplot2)
 # ggplot() + geom_line(data = dg1, aes(x = .index,y = .value, col = factor(.obs)))
 # + geom_line(data = dg2, aes(x = .index, y = .value, col = factor(.obs)))

 ngrid <- 50
 ev.grid <- seq(0, 1, , ngrid)
 nperm. <- 50

####
 simu.test <- DVDtest(dg1, dg2, nperm. ,ev.grid)
 simu.figs <- DVDplot(simu.test)
 simu.figs$pfig
 simu.figs$kfig[[1]]
```

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