

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

Suggests ggplot2, reshape2

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DVDtest-package	<i>Difference between Varying Distributions Test (DVDtest)</i>
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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.

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See Also[DVDtest](#)

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, respectively. 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 estimating 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 (=gamlss()) 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 permuted 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 lengths 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)"`, respectively.
- Multicore calculation is dealing with the long list of data.frame rather than 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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