# Package'DVDtest'

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Type Package  Title Difference between Varying Distributions Test (DVDtest)  Version 0.1  Date 2019-03-24  Author Meng Xu, Philip Reiss  Maintainer Meng Xu <mxu@campus.haifa.ac.il>  Description See DVDtest.  RoxygenNote 6.1.1  License GPL (&gt;= 2)  Encoding UTF-8  Imports gamlss, mgcv, ggplot2, reshape2, parallel, gamlss.dist  R topics documented:  DVDtest-package DVDtest  DVDtest  DVDtest  DVDtest  DVDtest  DVDtest  Description  This package contains a series of internal and external functions of Difference between Varying Distributions Test (DVDtest)  Author(s)  Meng Xu, Philip Reiss Maintainer: Meng Xu <mxu@campus.haifa.ac.il>  See Also  DVDtest</mxu@campus.haifa.ac.il></mxu@campus.haifa.ac.il>			
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DVDtest 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'). 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 are needed.
exclude	passed to exclude inside of predict.gam in the 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.

## **Details**

This is the Details section

#### Value

```
. index a vector, evaluation gridspval a vector or matrix of p values
```

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#### Note

• If the types of ydata1 and ydata2 are list of data. frames, the lengths of two lists must be the same.

• If mgcv.gam is TRUE, ... and exclue are NULL (default settings), then both defaults to formula <- list(.value~s(.index)+s(.obs, bs="re"), ~s(.index)) and exclude <- "s(.obs)"

### Author(s)

Meng Xu, Philip Reiss

#### References

reiss-EMR18.pdf

### **Examples**

```
## Data Generation ##
p <- 6
mu1 \leftarrow function(t) 0.2*(p-1)*sin(pi*t)+t+1
mu2 \leftarrow function(t) -0.2*(p-1)*sin(pi*t)+t+1
sig1 \leftarrow function(t) t+1
sig2 <- sig1
nperson <- 10
 fun1 <- function(t) rnorm(nperson, mu1(t), sig1(t))</pre>
 fun2 <- function(t) rnorm(nperson, mu2(t), sig2(t))</pre>
 tp <- seq(0,1,10)
 data1 <- sapply(tp,fun1)</pre>
data2 <- sapply(tp,fun2)</pre>
library(reshape2)
colnames(data1) <- tp</pre>
dg1 <- melt(data1)</pre>
colnames(dg1) <- c('.obs','.index','.value')</pre>
dg1$.obs <- as.factor(dg1$.obs)</pre>
colnames(data2) <- tp</pre>
 rownames(data2) <- 1:nperson+2*nperson</pre>
dg2 <- melt(data2)</pre>
 colnames(dg2) <- c('.obs','.index','.value')</pre>
 dg2$.obs <- as.factor(dg2$.obs)</pre>
 # 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)</pre>
 \#ggplot(data.frame(simu.test), aes(x = .index, y = pval)) + geom_line()
 # + geom_hline(yintercept = 0.05, linetype = 2, col = "red")
```

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