

# Package ‘DVDtest’

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**Type** Package

**Title** Difference between Varying Distributions Test (DVDtest)

**Version** 0.1

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**Author** Meng Xu, Philip Reiss

**Maintainer** Meng Xu <mxu@campus.haifa.ac.il>

**Description** See DVDtest.

**RoxygenNote** 6.1.1

**License** GPL (>= 2)

**Encoding** UTF-8

**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.

## Author(s)

Meng Xu, Philip Reiss

Maintainer: Meng Xu <mxu@campus.haifa.ac.il>

## See Also

[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.

**Details**

This is the Details section

**Value**

.index	a vector, evaluation grids
pval	a vector or matrix of (corrected) p values

**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 exclude are NULL (default settings), then defaults to  
`formula <- list(.value ~ s(.index) + s(.obs, bs = "re"), ~ s(.index))`  
and `exclude <- "s(.obs)"`, respectively.
- Now Normal distribution in `mgcv::gam` and BCCG, BCT and BCPE in `gamlss::gamlss` are supported by DVDtest for fitting a GAMLSS. Please contact the maintainer for 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(data1) <- tp
dg1 <- melt(data1)
colnames(dg1) <- c('.obs','.index','.value')
dg1$.obs <- as.factor(dg1$.obs)

colnames(data2) <- tp
rownames(data2) <- 1:nperson+2*nperson
dg2 <- melt(data2)
colnames(dg2) <- c('.obs','.index','.value')
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)
```

```
# 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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