

# Package: gmfd (via r-universe)

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**Type** Package

**Title** Inference and Clustering of Functional Data

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**Description** Some methods for the inference and clustering of univariate and multivariate functional data, using a generalization of Mahalanobis distance, along with some functions useful for the analysis of functional data. For further details, see Martino A., Ghiglietti, A., Ieva, F. and Paganoni A. M. (2017) <[arXiv:1708.00386](#)>.

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funData	<i>S3 Class for functional datasets. A class for univariate or multivariate functional dataset.</i>
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### Description

S3 Class for functional datasets. A class for univariate or multivariate functional dataset.

### Usage

```
funData(grid, data)
```

### Arguments

grid	the grid over which the functional dataset is defined.
data	a vector, a matrix or a list of vectors or matrices containing the functional data.

### Value

The function returns a S3 object of class funData, containing the grid over which the functional dataset is defined and a matrix or a list of vectors or matrices containing the functional data

### See Also

[gmfd\\_simulate](#)

### Examples

```
# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
```

```

m1 <- t^2 * ( 1 - t )
m2 <- t * ( 1 - t )^2

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}
# Simulate the functional data
x1 <- gmfd_simulate( n, m1, rho = rho, theta = theta )
x2 <- gmfd_simulate( n, m2, rho = rho, theta = theta )

FD <- funData( t, list( x1, x2 ) )

```

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funDist

*Distance function*


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## Description

This function allows you to compute the distance between two curves with the chosen metric.

## Usage

```

funDist(FD1, FD2, metric, p = NULL, lambda = NULL, phi = NULL,
        k_trunc = NULL)

```

## Arguments

FD1	a functional data object of type funData for the first curve
FD2	a functional data object of type funData for the second curve
metric	the chosen distance to be used: "L2" for the classical L2-distance, "trunc" for the truncated Mahalanobis semi-distance, "mahalanobis" for the generalized Mahalanobis distance.
p	a positive numeric value containing the parameter of the regularizing function for the generalized Mahalanobis distance.
lambda	a vector containing the eigenvalues in descending order of the functional data from which the curves are extracted.
phi	a matrix containing the eigenfunctions of the functional data in its columns from which the curves are extracted.
k_trunc	a positive numeric value representing the number of components at which the truncated mahalanobis distance must be truncated

**Value**

The function returns a numeric value indicating the distance between the two curves.

**References**

Ghiglietti A., Ieva F., Paganoni A. M. (2017). Statistical inference for stochastic processes: Two-sample hypothesis tests, *Journal of Statistical Planning and Inference*, 180:49-68.

Ghiglietti A., Paganoni A. M. (2017). Exact tests for the means of gaussian stochastic processes. *Statistics & Probability Letters*, 131:102–107.

**Examples**

```
# Define parameters:
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
m1 <- t^2 * ( 1 - t )

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}

# Simulate the functional data
z <- gmfd_simulate( n, m1, rho = rho, theta = theta )

# Extract two rows of the functional data
x <- funData( t, z[1, ] )
y <- funData( t, z[2, ] )

lambda <- eigen(cov(z))$values
phi <- eigen(cov(z))$vectors

d <- funDist( x, y, metric = "mahalanobis", p = 1, lambda = lambda, phi = phi )
```

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gmfd_diss	<i>Dissimilarity matrix function</i>
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### Description

This function computes the dissimilarity matrix containing the distances between the curves of the functional dataset

### Usage

```
gmfd_diss(FD, metric, p = NULL, k_trunc = NULL)
```

### Arguments

FD	a functional data object of type funData
metric	the chosen distance to be used. Choose "L2" for the classical L2-distance, "trunc" for the truncated Mahalanobis semi-distance, "mahalanobis" for the generalized Mahalanobis distance.
p	a positive numeric value containing the parameter of the regularizing function for the generalized Mahalanobis distance.
k_trunc	a positive numeric value representing the number of components at which the truncated mahalanobis distance must be truncated

### Value

The function returns a matrix of numeric values containing the distances between the curves.

### References

Ghiglietti A., Ieva F., Paganoni A. M. (2017). Statistical inference for stochastic processes: Two-sample hypothesis tests, *Journal of Statistical Planning and Inference*, 180:49-68.

Ghiglietti A., Paganoni A. M. (2017). Exact tests for the means of gaussian stochastic processes. *Statistics & Probability Letters*, 131:102–107.

### Examples

```
# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
m1 <- t^2 * ( 1 - t )
```

```

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}

# Simulate the functional data
x <- gmfd_simulate( n, m1, rho = rho, theta = theta )

FD <- funData( t, x )

D <- gmfd_diss( FD, metric = "L2" )

```

---

gmfd\_kmeans

*k-means clustering algorithm*


---

### Description

This function performs a k-means clustering algorithm on an univariate or multivariate functional data using a generalization of Mahalanobis distance.

### Usage

```
gmfd_kmeans(FD, n.cl = 2, metric, p = NULL, k_trunc = NULL)
```

### Arguments

FD	a functional data object of type funData.
n.cl	an integer representing the number of clusters.
metric	the chosen distance to be used: "L2" for the classical L2-distance, "trunc" for the truncated Mahalanobis semi-distance, "mahalanobis" for the generalized Mahalanobis distance.
p	a positive numeric value containing the parameter of the regularizing function for the generalized Mahalanobis distance.
k_trunc	a positive numeric value representing the number of components at which the truncated mahalanobis distance must be truncated

### Value

The function returns a list with the following components: `cluster`: a vector of integers (from 1 to `n.cl`) indicating the cluster to which each curve is allocated; `centers`: a list of `d` matrices (`k x T`) containing the centroids of the clusters

## References

- Martino A., Ghiglietti A., Ieva F., Paganoni A. M. (2017). A k-means procedure based on a Mahalanobis type distance for clustering multivariate functional data, *MOX report 44/2017*
- Ghiglietti A., Ieva F., Paganoni A. M. (2017). Statistical inference for stochastic processes: Two-sample hypothesis tests, *Journal of Statistical Planning and Inference*, 180:49-68.
- Ghiglietti A., Paganoni A. M. (2017). Exact tests for the means of gaussian stochastic processes. *Statistics & Probability Letters*, 131:102–107.

## See Also

[funDist](#)

## Examples

```
# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
m1 <- t^2 * ( 1 - t )

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}

s <- 0
for (k in 4:K) {
  s <- s + sqrt( rho[k] ) * theta[k, ]
}

m2 <- m1 + s

# Simulate the functional data
x1 <- gmfd_simulate( n, m1, rho = rho, theta = theta )
x2 <- gmfd_simulate( n, m2, rho = rho, theta = theta )

# Create a single functional dataset containing the simulated datasets:
FD <- funData(t, rbind( x1, x2 ) )
```

```
output <- gmfd_kmeans( FD, n.cl = 2, metric = "mahalanobis", p = 10^6 )
```

---

gmfd\_simulate                      *Simulation of a functional sample*

---

### Description

Simulate a univariate functional sample using a Karhunen Loeve expansion.

### Usage

```
gmfd_simulate(size, mean, covariance = NULL, rho = NULL, theta = NULL)
```

### Arguments

size	a positive integer indicating the size of the functional sample to simulate.
mean	a vector representing the mean of the sample.
covariance	a matrix from which the eigenvalues and eigenfunctions must be extracted.
rho	a vector of the eigenvalues in descending order to be used for the simulation.
theta	a matrix containing the eigenfunctions in its columns to be used for the simulation.

### Value

The function returns a functional data object of type funData.

### Examples

```
# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
# with the Karhunen - Loève expansion
m1 <- t^2 * ( 1 - t )

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
}
```

```

else
  theta[k, ] <- rep( 1, P )
}

# Simulate the functional data
x <- gmfd_simulate( n, m1, rho = rho, theta = theta )

```

gmfd\_test

*Two-sample hypothesis tests***Description**

Performs a two sample hypothesis tests on two samples of functional data.

**Usage**

```
gmfd_test(FD1, FD2, conf.level = 0.95, stat_test, p = NULL,
          k_trunc = NULL)
```

**Arguments**

FD1	a functional data object of type <code>funData</code> of the first sample.
FD2	a functional data object of type <code>funData</code> of the second sample.
conf.level	confidence level of the test.
stat_test	the chosen test statistic to be used: "L2" for the classical L2-distance, "L2_trunc" for the truncated L2-distance, "trunc" for the truncated Mahalanobis semi-distance, "mahalanobis" for the generalized Mahalanobis distance
p	a vector of positive numeric value containing the parameters of the regularizing function for the generalized Mahalanobis distance.
k_trunc	a positive numeric value representing the number of components at which the truncated mahalanobis distance must be truncated

**Value**

The function returns a list with the following components:

`statistic` the value of the test statistic.

`quantile` the value of the quantile.

`p.value` the p-value for the test.

**References**

Ghiglietti A., Ieva F., Paganoni A. M. (2017). Statistical inference for stochastic processes: Two-sample hypothesis tests, *Journal of Statistical Planning and Inference*, 180:49-68.

Ghiglietti A., Paganoni A. M. (2017). Exact tests for the means of gaussian stochastic processes. *Statics & Probability Letters*, 131:102–107.

**See Also**[funDist](#)**Examples**

```

# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
m1 <- t^2 * ( 1 - t )

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}

s <- 0
for ( k in 4:K ) {
  s <- s + sqrt( rho[k] ) * theta[k,]
}

m2 <- m1 + 0.1 * s

# Simulate the functional data
x1 <- gmfd_simulate( n, m1, rho = rho, theta = theta )
x2 <- gmfd_simulate( n, m2, rho = rho, theta = theta )
FD1 <- funData( t, x1 )
FD2 <- funData( t, x2 )
output <- gmfd_test( FD1, FD2, 0.95, "mahalanobis", p = 10^5 )

```

---

`plot.funData`*A method to plot funData objects*

---

**Description**

This function performs the plot of a functional dataset stored in an object of class `funData`.

**Usage**

```
## S3 method for class 'funData'
plot(x, ...)
```

**Arguments**

x                    the univariate functional dataset in form of funData object.  
 ...                  additional graphical parameters to be used in plotting functions

**See Also**

[funData](#)

**Examples**

```
# Define parameters
n <- 50
P <- 100
K <- 150

# Grid of the functional dataset
t <- seq( 0, 1, length.out = P )

# Define the means and the parameters to use in the simulation
m1 <- t^2 * ( 1 - t )
m2 <- t * ( 1 - t )^2

rho <- rep( 0, K )
theta <- matrix( 0, K, P )
for ( k in 1:K ) {
  rho[k] <- 1 / ( k + 1 )^2
  if ( k%%2 == 0 )
    theta[k, ] <- sqrt( 2 ) * sin( k * pi * t )
  else if ( k%%2 != 0 && k != 1 )
    theta[k, ] <- sqrt( 2 ) * cos( ( k - 1 ) * pi * t )
  else
    theta[k, ] <- rep( 1, P )
}
# Simulate the functional data
x1 <- gmfd_simulate( n, m1, rho = rho, theta = theta )
x2 <- gmfd_simulate( n, m2, rho = rho, theta = theta )

FD <- funData( t, list( x1, x2 ) )

plot(FD)
```

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