

Package ‘ghcm’

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Type Package

Title Functional Conditional Independence Testing with the GHCM

Version 3.0.1

Description A statistical hypothesis test for conditional independence.

Given residuals from a sufficiently powerful regression, it tests whether the covariance of the residuals is vanishing. It can be applied to both discretely-observed functional data and multivariate data.

Details of the method can be found in Anton Rask Lundborg, Rajen D. Shah and Jonas Peters (2022) <[doi:10.1111/rssb.12544](https://doi.org/10.1111/rssb.12544)>.

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Encoding UTF-8

LazyData true

Imports CompQuadForm, Rcpp, splines

Depends R (>= 4.0.0)

RoxygenNote 7.2.3

Suggests graphics, stats, utils, refund, testthat, knitr, rmarkdown, bookdown, ggplot2, reshape2, dplyr, tidyr

URL <https://github.com/arlundborg/ghcm>

BugReports <https://github.com/arlundborg/ghcm/issues>

VignetteBuilder knitr

LinkingTo Rcpp

NeedsCompilation yes

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Contents

ghcm	2
ghcm_sim_data	2
ghcm_test	3
inner_product_matrix_splines	5

Index	7
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ghcm	<i>ghcm: A package for Functional Conditional Independence Testing</i>
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Description

To learn more about ghcm, start with the vignette: `'browseVignettes(package = "ghcm")'`

ghcm_sim_data	<i>GHCM simulated data</i>
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Description

A simulated dataset containing a combination of functional and scalar variables. Y_1 and Y_2 are scalar random variables and are both functions of Z . X , Z and W are functional, Z is a function of X and W is a function of Z .

Usage

`ghcm_sim_data`

`ghcm_sim_data_irregular`

Format

`ghcm_sim_data` is a data frame with 500 rows of 5 variables:

Y_1 Numeric vector.

Y_2 Numeric vector.

Z 500 x 101 matrix.

X 500 x 101 matrix.

W 500 x 101 matrix.

`ghcm_sim_data_irregular` is a list with 5 elements:

Y_1 Numeric vector.

Y_2 Numeric vector.

Z 500 x 101 matrix.

X A data frame with

.obs Integer between 1 and 500 indicating which curve the row corresponds to.

.index Function argument that the curve is evaluated at.

.value Value of the function.

W A data frame with

.obs Integer between 1 and 500 indicating which curve the row corresponds to.

.index Function argument that the curve is evaluated at.

.value Value of the function.

Details

In `ghcm_sim_data` the functional variables each consists of 101 observations on an equidistant grid on $[0, 1]$.

In `ghcm_sim_data_irregular` the functional variables `X` and `W` are instead only observed on a subsample of the original equidistant grid.

Source

The generation script can be found in the `data-raw` folder of the package.

ghcm_test

Conditional Independence Test using the GHCM

Description

Test whether X is independent of Y given Z using the Generalised Hilbertian Covariance Measure. The function is applied to residuals from regressing each of X and Y on Z respectively. Its validity is contingent on the performance of the regression methods. For a more in-depth explanation see the package vignette or the paper mentioned in the references.

Usage

```
ghcm_test(  
  resid_X_on_Z,  
  resid_Y_on_Z,  
  X_limits = NULL,  
  Y_limits = NULL,  
  alpha = 0.05  
)
```

Arguments

- `resid_X_on_Z, resid_Y_on_Z`
Residuals from regressing X (Y) on Z with a suitable regression method. If X (Y) is uni- or multivariate or functional on a constant, fixed grid, the residuals should be supplied as a vector or matrix with no missing values. If instead X (Y) is functional and observed on varying grids or with missing values, the residuals should be supplied as a "melted" data frame with
- .obs** Integer indicating which curve the row corresponds to.
 - .index** Function argument that the curve is evaluated at.
 - .value** Value of the function.
- Note that in the irregular case, a minimum of 4 observations per curve is required.
- `X_limits, Y_limits`
The minimum and maximum values of the function argument of the X (Y) curves. Ignored if X (Y) is not functional.
- `alpha`
Numeric in the unit interval. Significance level of the test.

Value

- An object of class `ghcm` containing:
- `test_statistic` Numeric, test statistic of the test.
 - `p` Numeric in the unit interval, estimated p-value of the test.
 - `alpha` Numeric in the unit interval, significance level of the test.
 - `reject` TRUE if $p < \alpha$, FALSE otherwise.

References

Please cite the following paper: Anton Rask Lundborg, Rajen D. Shah and Jonas Peters: "Conditional Independence Testing in Hilbert Spaces with Applications to Functional Data Analysis" *Journal of the Royal Statistical Society: Series B (Statistical Methodology)* 2022 [doi:10.1111/rssb.12544](https://doi.org/10.1111/rssb.12544).

Examples

```
if (require(refund)) {
  set.seed(1)
  data(ghcm_sim_data)
  grid <- seq(0, 1, length.out = 101)

  # Test independence of two scalars given a functional variable

  m_1 <- pfr(Y_1 ~ lf(Z), data=ghcm_sim_data)
  m_2 <- pfr(Y_2 ~ lf(Z), data=ghcm_sim_data)
  ghcm_test(resid(m_1), resid(m_2))

  # Test independence of a regularly observed functional variable and a
  # scalar variable given a functional variable
```

```

    m_X <- pffr(X ~ ff(Z), data=ghcm_sim_data, chunk.size=31000)
    ghcm_test(resid(m_X), resid(m_1))

# Test independence of two regularly observed functional variables given
# a functional variable

    m_W <- pffr(W ~ ff(Z), data=ghcm_sim_data, chunk.size=31000)
    ghcm_test(resid(m_X), resid(m_W))

data(ghcm_sim_data_irregular)
n <- length(ghcm_sim_data_irregular$Y_1)
Z_df <- data.frame(.obs=1:n)
Z_df$Z <- ghcm_sim_data_irregular$Z
# Test independence of an irregularly observed functional variable and a
# scalar variable given a functional variable

    m_1 <- pfr(Y_1 ~ lf(Z), data=ghcm_sim_data_irregular)
    m_X <- pffr(X ~ ff(Z), ydata = ghcm_sim_data_irregular$X,
    data=Z_df, chunk.size=31000)
    ghcm_test(resid(m_X), resid(m_1), X_limits=c(0, 1))

# Test independence of two irregularly observed functional variables given
# a functional variable

    m_W <- pffr(W ~ ff(Z), ydata = ghcm_sim_data_irregular$W,
    data=Z_df, chunk.size=31000)
    ghcm_test(resid(m_X), resid(m_W), X_limits=c(0, 1), Y_limits=c(0, 1))

}

```

```
inner_product_matrix_splines
```

Computes the matrix of L2 inner products of the splines given in list_of_splines as produced by splines::interpSpline. The splines are assumed to be functions on the interval [from, to].

Description

Computes the matrix of L2 inner products of the splines given in list_of_splines as produced by splines::interpSpline. The splines are assumed to be functions on the interval [from, to].

Usage

```
inner_product_matrix_splines(list_of_splines, from, to)
```

Arguments

`list_of_splines` list of `interpSpline` objects.
`from, to` limits of integration.

Value

matrix of inner products.

Index

* datasets

ghcm_sim_data, 2

ghcm, 2

ghcm_sim_data, 2

ghcm_sim_data_irregular
(ghcm_sim_data), 2

ghcm_test, 3

inner_product_matrix_splines, 5